COVID-19
Scientific literature review

A digest of peer-reviewed articles from top-ranked journals on selected research topics related to SARS-CoV-2 and COVID-19

UPDATE OF
03 JUNE 2021

L’ANRS|Emerging Infectious Diseases shares a selection of the most relevant articles published on COVID-19 on a weekly basis. This literature review not only presents a selection of references, but also highlights the key points and messages from each article. It does not include pre-print articles.

Our objective is to help the scientific community, health-workers and public health decision makers, being up to date with the latest scientific research.

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Additional links:
Haute Autorité de Santé: https://www.scoop.it/topic/coronavirus-covid-19-has-veille?nosug=1
MODCOV19: https://modcov19.math.cnrs.fr/veille_public/
<table>
<thead>
<tr>
<th>Journal and date</th>
<th>Title</th>
<th>Authors and link</th>
<th>Field of expertise</th>
<th>Key facts</th>
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<td>Lancet Microbe 02JUN2021</td>
<td>Co-infections, secondary infections, and antimicrobial use in patients hospitalised with COVID-19 during the first pandemic wave from the ISARIC WHO CCP-UK study: a multicentre, prospective cohort study</td>
<td>Russel C.D., et al. UK gotopaper</td>
<td>Clinic</td>
<td>Aim: to describe microbiologically confirmed co-infections and secondary infections, and antimicrobial use, in patients admitted to hospital with COVID-19 (ISARIC, CCP-UK study). Results &gt; Analysis of data from 48 902 patients admitted to hospital (Feb 6 and June 8, 2020). Median age 74 years (IQR 59–84), 20 786 (42.6%) of 48 765 women. Microbiological investigations were recorded for 8649 (17.7%) of 48 902 patients, with clinically significant COVID-19-related respiratory or bloodstream culture results recorded for 1107 patients. &gt; 762 (70.6%) of 1080 infections were secondary, occurring more than 2 days after hospital admission. &gt; Staphylococcus aureus and Haemophilus influenzae were the most common pathogens causing respiratory co-infections (diagnosed ≥2 days after admission), with Enterobacteriaceae and S aureus most common in secondary respiratory infections. &gt; Bloodstream infections were most frequently caused by Escherichia coli and S aureus. &gt; Among patients with available data, 13 390 (37.0%) of 36 145 had received antimicrobials in the community for this illness episode before hospital admission. &gt; 39 258 (85.2%) of 46 061 patients with inpatient antimicrobial data received one or more antimicrobials at some point during their admission (highest for patients in critical care). &gt; Frequent use of broad-spectrum agents and use of carbapenem rather than carbapenem-sparing alternatives was identified. In patients admitted to hospital with COVID-19, bacterial infections are rare, and more likely to be secondary infections. The frequency and nature of antimicrobial use are concerning.</td>
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<td>JAMA Intern Med. 28MAY2021</td>
<td>Assessment of SARS-CoV-2 Reinfection 1 Year After Primary Infection in a Population in Lombardy, Italy</td>
<td>Vitale J., et al. Italy gotopaper</td>
<td>Public Health / Epidemiology</td>
<td>Aim: to investigate the one-year incidence (Feb 2021) of SARS-CoV-2 primary infection and reinfection among individuals who underwent diagnostic RT-PCR between Feb-July 2020 in Lombardy, Italy. Reinfections were defined by a second RT-PCR positivity beyond 90 days after complete resolution of the first infection and with at least 2 consecutive negative test results between episodes. Results &gt; Demographic characteristics: Median age 59 (IQR 40-78) years, positive cases were older and geographically distributed more in the industrial area of Legnano. &gt; During follow-up (mean [SD], 280 [41] days) 5 reinfections (0.31%; 95% CI, 0.03%-0.58%) were confirmed among 1579 positive patients. &gt; Most of reinfeected patients were evaluated, treated, and followed in hospitals or dedicated COVID-19 ambulatories. One was hospitalized, and 4 had a close relationship with health facilities. Mean (SD) interval between primary infection and reinfection was &gt;230 (90) days. &gt; Of 13 496 persons initially not infected with SARS-CoV-2, 528 (3.9%; 95% CI, 3.5%-4.2%) subsequently developed a primary infection. &gt; The incidence density per 100 000 person days was 1.0 (95% CI, 0.5-1.5) for reinfections compared with 15.1 (95% CI, 14.5-15.7) for new infections, while the incidence rate ratio adjusted for age, sex, ethnicity, and the sanitary area was 0.07 (95% CI, 0.06-0.08). The study results suggest that reinfections are rare events and patients who have recovered from COVID-19 have a lower risk of reinfection 1 year later. Observation ended before SARS-CoV-2 variant spread.</td>
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| **JAMA Oncol. 28MAY2021** | Evaluation of Seropositivity Following BNT162b2 Messenger RNA Vaccination for SARS-CoV-2 in Patients Undergoing Treatment for Cancer | Massarweh A., et al. Israel gotopaper | Vaccines - Immunisation | **Aim:** To evaluate rates of antispike (anti-S) antibody response to a BNT162b2 vaccine in patients with cancer who are undergoing systemic treatment vs healthy controls.  

**Methods:**  
Prospective cohort study included 102 adult patients with solid tumors undergoing active intravenous anticancer treatment and 78 controls who received the second dose of the BNT162b2 vaccine at least 12 days before enrollment. Serum samples were analyzed and the titers of the IgG antibodies against SARS-CoV-2 spike receptor-binding domain  

**Results:**  
The analysis included 180 participants, which comprised 102 patients with cancer (median [interquartile range (IQR)] age, 66 years; 58 men [57%]) and 78 healthy controls (median [IQR] age, 62 [49-70] years; 25 men [32%]).  
The most common tumor type was gastrointestinal (29 [28%]). In the patient group, 92 (90%) were seropositive for SARS-CoV-2 antispike IgG antibodies after the second vaccine dose, whereas in the control group, all were seropositive.  
The median IgG titer in the patients with cancer was significantly lower than that in the controls (1931 [IQR, 509-4386] AU/ml vs 7160 [IQR, 3129-11 241] AU/m).  
In a multivariable analysis, the only variable that was significantly associated with lower IgG titers was treatment with chemotherapy plus immunotherapy (β, −3.5; 95% CI, −5.6 to −1.5). |
| **JAMA Oncol. 28MAY2021** | Difference in SARS-CoV-2 Antibody Status Between Patients With Cancer and Health Care Workers During the COVID-19 Pandemic in Japan | Yazaki S., et al. Japan gotopaper | Vaccines - Immunisation | **Aim:** To evaluate serum SARS-CoV-2 antibody status in patients with cancer and health care workers (HCWs) during the COVID-19 pandemic in Japan.  

**Methods:**  
Participants were enrolled for this prospective cross-sectional study between August 3 and October 30, 2020, from 2 comprehensive cancer centers in the epidemic area around Tokyo, Japan.  

**Results:**  
A total of 500 patients with cancer (median age, 62.5 years [range, 21-88 years]; 265 men [55.4%]) and 1190 HCWs (median age, 40 years [range, 20-70 years]; 382 men [25.4%]) were enrolled.  
The seroprevalence was 1.0% (95% CI, 0.33%-2.32%) in patients and 0.67% (95% CI, 0.29%-1.32%) in HCWs (P = .48).  
The N-IgG and S-IgG antibody levels were significantly lower in patients than in HCWs (N-IgG: β, −0.38; 95% CI, −0.55 to −0.21; P < .001; and S-IgG: β, −0.39; 95% CI, −0.54 to −0.23; P < .001).  
Additionally, among patients, N-IgG levels were significantly lower in those who received chemotherapy than in those who did not (median N-IgG levels, 0.1 [IQR, 0.1-0.5] vs 0.1 [IQR, 0.1-0.3], P = .04).  
In contrast, N-IgG and S-IgG levels were significantly higher in patients who received immune checkpoint inhibitors than in those who did not (median N-IgG levels: 0.2 [IQR, 0.1-0.5] vs 0.1 [IQR, 0.0-0.3], P = .02; S-IgG levels: 0.15 [IQR, 0.0-0.3] vs 0.1 [IQR, 0.0-0.2], P = .02).  
The seroprevalence of SARS-CoV-2 antibodies did not differ between the 2 groups; however, findings suggest that comorbid cancer and treatment with systemic therapy, including chemotherapy and immune checkpoint inhibitors, may influence the immune response to SARS-CoV-2. |
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| PNAS 28MAY2021   | Intracounty modeling of COVID-19 infection with human mobility: Assessing spatial heterogeneity with business traffic, age, and race | Hou X., et al. USA gotopaper | Public Health / Epidemiology | **Aim:** To develop models incorporating spatial heterogeneity and mobility flows to make predictions on the effect of nonpharmaceutical interventions  
**Methods:**  
> SEIR epidemic model capturing social and geographic heterogeneity within a small geographic region (a single county), with stochastic transmission rate and heterogeneous local mobility  
> Unsupervised machine learning to partition a county into local clusters using human mobility flow data from SafeGraph, giving smartphone coordinates over time  
> Other data: test results (Public Health Offices of City of Madison & Dane County and Milwaukee County) between March 11-August 14, 2020; demographic and socioeconomic attributes (US Census Bureau)  
> Online data assimilation and the ensemble Kalman filter method are used to update the model parameters along time  
> Study of several scenarios of reopening policies  
**Findings:**  
> The spatial clustering constructed from mobility data is strongly correlated with demographic heterogeneity (e.g. in terms of race and ethnicity composition, age structure)  
> There is a strong heterogeneity of the estimated effective reproduction number across clusters  
> In a college town (Dane County), the most important heterogeneity is age structure  
> In a large city area (Milwaukee County), racial and ethnic heterogeneity becomes more apparent  
> Scenario studies indicate a strong response of the spread rate to various reopening policies  
**Conclusion:** Policy makers may need to take social and geographic heterogeneities into account very carefully when designing policies for mitigating the ongoing spread of Covid-19 and reopening |
| NEJM 27MAY2021   | Safety, Immunogenicity, and Efficacy of the BNT162b2 Covid-19 Vaccine in Adolescents | Frenck R.W., et al. International gotopaper | Vaccines | Safe, effective vaccines are needed to protect adolescents, facilitate in-person learning and socialization, and contribute to herd immunity  
**Methods**  
> Ongoing multinational, placebo-controlled, observer-blinded trial  
> Participants randomly assigned in a 1:1 ratio to receive two injections, 21 days apart, of 30 μg of BNT162b2 or placebo.  
> Immunogenicity objective: noninferiority of the immune response to BNT162b2 in 12- to 15-year-old participants as compared with that in 16- to 25-year-old participants  
> Safety (reactogenicity and adverse events) and efficacy against confirmed Covid-19 (onset ≥7 days after dose 2) in 12-15- yo cohort.  
**Findings**  
> 1131 adolescents 12 to 15 years of age received BNT162b2; 1129 received placebo.  
> BNT162b2 had a favorable safety and side-effect profile, with mainly transient mild to moderate reactogenicity (mainly injection-site pain [79-86% of participants], fatigue [60-66%], headache [55-65%]).  
> No vaccine-related serious adverse events and few overall severe adverse events.  
> The GMT of SARS-CoV-2 50% neutralizing titers after dose 2 in 12- to 15-year-old participants relative to 16- to 25-year-old participants was 1.76 (95% confidence interval [CI], 1.47 to 2.10), which met the noninferiority criterion of a lower boundary of the two-sided 95% confidence interval greater than 0.67 and indicated a greater response in the 12- to 15-year-old cohort.  
> Among participants without evidence of previous SARS-CoV-2 infection, no Covid-19 cases with an onset of 7 or more days after dose 2 were noted among BNT162b2 recipients, and 16 cases occurred among placebo recipients.  
> The observed VE was 100% (95% CI, 75.3 to 100).  
**Conclusions** The BNT162b2 vaccine in 12-to-15-year-old recipients had a favorable safety profile, produced a greater immune response than in young adults, and was highly effective against Covid-19. |
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<th>Field of expertise</th>
<th>Key facts</th>
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- Phase 3, 1:1 randomised, double-blind, adaptive, placebo-controlled, international multicentre trial.  
- Patients with COVID-19 diagnosed by PCR testing or clinical criteria who were not being treated in hospital, ≥40-year-old and who had at least one high-risk characteristic.  
- Treatment: orally administered colchicine (0.5 mg twice/day for 3 days, then once/day for 27 days thereafter) or matching placebo.  
**Primary efficacy endpoint:** composite of death or hospital admission for COVID-19.  
**Results:**  
> Trial enrolment: March 23, 2020 - Dec 22, 2020. 4488 patients included (53.9% women; median age 54.0 years, IQR 47.0–61.0), 2235 were randomly assigned to colchicine and 2253 to placebo.  
> Overall, the primary endpoint occurred in 104 (4.7%) of 2235 patients in the colchicine group and 131 (5.8%) of 2253 patients in the placebo group (odds ratio [OR] 0.79, 95% CI 0.61–1.03; p=0.081).  
> Among the 4159 patients with PCR-confirmed COVID-19, the primary endpoint occurred in 96 (4.6%) of 2075 patients in the colchicine group and 126 (5.6%) of 2084 patients in the placebo group (OR 0.75, 0.57–0.99; p=0.042).  
> Serious adverse events were reported in 108 (4.9%) of 2195 patients in the colchicine group and 139 (6.3%) of 2217 patients in the placebo group (p=0.051); pneumonia occurred in 63 (2.9%) of 2195 patients in the colchicine group and 92 (4.1%) of 2217 patients in the placebo group (p=0.021). Diarrhoea was reported in 300 (13.7%) of 2195 patients in the colchicine group and 161 (7.3%) of 2217 patients in the placebo group (p<0.0001).  
**In community-treated patients** including those without diagnostic test, the **effect of colchicine on COVID-19-related clinical events was not statistically significant.** Among patients with PCR-confirmed COVID-19, colchicine led to a lower rate of the composite of death or hospital admission than placebo. |
| Lancet Infect Dis. 27MAY2021 | Same-day SARS-CoV-2 antigen test screening in an indoor mass-gathering live music event: a randomised controlled trial | Revollo B., et al. Spain gotopaper | Public Health / Epidemiology | **Primary outcome:** difference in incidence of RT-PCR-confirmed SARS-CoV-2 infection at 8 days, control vs. intervention groups.  
**Findings:**  
> 1047 people randomized. 465 were finally included in the final analysis for the experimental group and 491 in the control group.  
> At baseline, 15 (3%) of 495 individuals in the control group and 13 (3%) of 465 in the experimental group tested positive on TMA despite a negative Ag-RDT result.  
> The RT-PCR test was positive in one case in each group and cell viral culture was negative in all cases.  
> 8 days after the event, two (<1%) individuals in the control arm had a positive Ag-RDT and PCR result, whereas no Ag-RDT nor RT-PCR positive results were found in the intervention arm.  
> The Bayesian estimate for the incidence between the experimental and control groups was −0.15% (95% CI −0.72 to 0.44).  
**Conclusions**  
Preliminary evidence on indoor mass-gathering event safety during a COVID-19 outbreak under a comprehensive preventive intervention. |
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<td><strong>Clin Infect Dis.</strong>&lt;br&gt;27MAY2021</td>
<td>Risk Factors for Death Among the First 80 543 COVID-19 Cases in China: Relationships Between Age, Underlying Disease, Case Severity, and Region</td>
<td>Zhang Y., et al.&lt;br&gt;China [gotopaper]</td>
<td>Public Health / Epidemiology</td>
<td>Examining risk factors for COVID-19 death&lt;br&gt;&lt;br&gt;<strong>Methods</strong>&lt;br&gt; &gt; A total of 80 543 COVID-19 cases reported in China, nationwide, through April 8, 2020 were included&lt;br&gt; &lt;br&gt;<strong>Findings</strong>&lt;br&gt; &gt; Overall national case fatality ratio (CFR) was 5.64%&lt;br&gt; &gt; Risk factors for death were older age, presence of underlying disease, worse case severity, and near-epicenter region&lt;br&gt; &gt; CFR increased from 0.35% (30-39 years) to 18.21% (≥70 years) without underlying disease&lt;br&gt; &gt; Regardless of age, CFR increased from 2.50% for no underlying disease to 7.72% for 1, 13.99% for 2, and 21.99% for ≥3&lt;br&gt; &gt; CFR increased with worse case severity from 2.80% (mild), to 12.51% (severe) and 48.60% (critical) regardless of region&lt;br&gt; &gt; Compared to other regions, CFR was much higher in Wuhan regardless of case severity (mild: 3.83% versus 0.14% in Hubei and 0.03% elsewhere; moderate: 4.60% versus 0.21% and 0.06%; severe: 15.92% versus 5.84% and 1.86%; and critical: 58.57% versus 49.80% and 18.39%).&lt;br&gt; &gt; Older patients regardless of underlying disease and patients with underlying disease regardless of age were at elevated risk of death.&lt;br&gt; &gt; Higher death rates near the outbreak epicenter and during the surge of cases reflect the deleterious effects of allowing health systems to become overwhelmed.</td>
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<td><strong>JAMA</strong>&lt;br&gt;26MAY2021</td>
<td>Effect of 2 Inactivated SARS-CoV-2 Vaccines on Symptomatic COVID-19 Infection in Adults</td>
<td>Kaabi N.A., et al.&lt;br&gt;International [gotopaper]</td>
<td>Vaccines</td>
<td>Evaluation of the efficacy and adverse events of 2 inactivated COVID-19 vaccines.&lt;br&gt;<strong>Methods</strong>&lt;br&gt; Randomized, double-blind, phase 3 trial. United Arab Emirates and Bahrain. &gt; 18 years of age without history of COVID-19. Participants randomized to receive 1 of 2 inactivated vaccines developed from SARS-CoV-2 WIV04 (5 µg/dose; n = 13 459) and HB02 (4 µg/dose; n = 13 465) strains or an aluminum hydroxide (alum)–only control (n = 13 458). 2 IM injections 21 days apart.&lt;br&gt; Primary outcome: efficacy against laboratory-confirmed symptomatic COVID-19 14 days following a second vaccine dose among participants who had no virologic evidence of SARS-CoV-2 infection at randomization.&lt;br&gt; Secondary outcome: efficacy against severe COVID-19.&lt;br&gt; <strong>Findings</strong>&lt;br&gt; &gt; 40 382 participants randomized (mean age, 36.1 years; 32 261 [84.4%] men), &gt; 38 206 (94.6%) who received 2 doses were included in the primary efficacy analysis.&lt;br&gt; &gt; Symptomatic COVID-19 was identified in 26 participants in the WIV04 group (12.1 [95% CI, 8.3-17.8] per 1000 person-years), 21 in the HB02 group (9.8 [95% CI, 6.4-15.0] per 1000 person-years), and 95 in the alum-only group (44.7 [95% CI, 36.6-54.6] per 1000 person-years). VE: 72.8% (95% CI, 58.1%-82.4%) for WIV04 and 78.1% (95% CI, 64.8%-86.3%) for HB02 (P &lt; .001 for both).&lt;br&gt; &gt; Two severe cases of COVID-19 occurred in the alum-only group and none occurred in the vaccine groups.&lt;br&gt; &gt; Adverse reactions 7 days after each injection occurred in 41.7% to 46.5% of participants in the 3 groups; serious adverse events were rare and similar in the 3 groups (WIV04: 64 [0.5%]; HB02: 59 [0.4%]; alum-only: 78 [0.6%]).&lt;br&gt; <strong>Conclusions</strong>&lt;br&gt; Treatment of adults with either of 2 inactivated SARS-CoV-2 vaccines significantly reduced the risk of symptomatic COVID-19, and serious adverse events were rare.</td>
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| **Nature** 27MAY2021 | BNT162b2 vaccine induces neutralizing antibodies and poly-specific T cells in humans | Sahin U., et al. Germany [gotopaper](#) | Vaccines | **Background**
BNT162b2, a lipid nanoparticle (LNP) formulated nucleoside-modified messenger RNA (mRNA) that encodes the SARS-CoV-2 spike glycoprotein (S) stabilized in the prefusion conformation, has demonstrated 95% efficacy in preventing COVID-19.

**Aim:** to extend the previous phase 1/2 trial report and present BNT162b2 prime/boost induced immune response data from a second phase 1/2 trial in healthy adults (18-55 years of age).

**Results**
- BNT162b2 elicited strong antibody responses, with SARS-CoV-2 serum 50% neutralizing geometric mean titers up to 3.3-fold above those observed in COVID-19 human convalescent samples (HCS) one-week post-boost.
- BNT162b2-elicited sera neutralized 22 pseudoviruses bearing SARS-CoV-2 S variants.
- Most participants had a strong IFNγ- or IL-2-positive CD8+ and CD4+ T helper type 1 (TH1) T cell response, detectable throughout the full observation period of nine weeks following the boost.
- pMHC multimer technology identified several BNT162b2-induced epitopes that were presented by frequent MHC alleles and conserved in mutant strains.
- One-week post-boost, epitope-specific CD8+ T cells of the early differentiated effector-memory phenotype comprised 0.02-2.92% of total circulating CD8+ T cells and were detectable (0.01-0.28%) eight weeks later.
- BNT162b2 elicits an adaptive humoral and poly-specific cellular immune response against epitopes conserved in a broad range of variants at well tolerated doses.


**Methods**
- Prespecified interim analysis of an ongoing randomized, double-blind, phase 3 trial in the United Arab Emirates and Bahrain among adults 18 years and older without known history of COVID-19. ClinicalTrials.gov identifier: NCT04510207;
- Participants were randomized to receive 1 of 2 inactivated vaccines developed from SARS-CoV-2 WIV04 (5 µg/dose; n = 13,459) and HB02 (4 µg/dose; n = 13,465) strains or an aluminum hydroxide (alum)-only control (n = 13,458) (2 intramuscular injections 21 days apart)

**Outcomes:** efficacy against laboratory-confirmed symptomatic COVID-19 14 days following a second vaccine dose among participants who had no virologic evidence of SARS-CoV-2 infection at randomization.

**Efficacy against severe COVID-19:** Incidence of adverse events and reactions was collected in participants who received at least 1 dose.

**Findings:**
- 40,382 participants (mean age 36.1 yrs; 32,261 [84.4%] men)
- 38,206 (94.6%) received 2 doses
- During a median (range) follow-up duration of 77 (1-121) days, symptomatic COVID-19 was identified in 26 participants in the WIV04 group (12.1 [95% CI, 8.3-17.8] per 1000 person-years), 21 in the HB02 group (9.8 [95% CI, 6.4-15.0] per 1000 person-years), and 95 in the alum-only group (44.7 [95% CI, 36.6-54.6] per 1000 person-years), resulting in a vaccine efficacy, compared with alum-only, of 72.8% (95% CI, 58.1%-82.4%) for WIV04 and 78.1% (95% CI, 64.8%-86.3%) for HB02 (P < .001 for both).
- Two severe cases of COVID-19 occurred in the alum-only group and none occurred in the vaccine groups.
- Adverse reactions 7 days after each injection occurred in 41.7% to 46.5% of participants in the 3 groups; serious adverse events were rare and similar in the 3 groups (WIV04: 64 [0.5%]; HB02: 59 [0.4%]; alum-only: 78 [0.6%]).

**Conclusion**
Treatment of adults with either of 2 inactivated SARS-CoV-2 vaccines significantly reduced the risk of symptomatic COVID-19, and serious adverse events were rare.
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| PNAS 25MAY2021   | Just 2% of SARS-CoV-2-positive individuals carry 90% of the virus circulating in communities | Yang Q., et al. USA | Public Health / Epidemiology | **Aim:** Describing the estimated distribution of viral loads among SARS-CoV-2-positive asymptomatic individuals detected through random surveillance  
**Methods:**  
> Surveillance through repeated testing of asymptomatic individuals on the University of Colorado Boulder campus (72500 saliva samples; 1405 positive, mostly from unique individuals).  
> Detection by qRT-PCR of saliva samples  
> Estimation of viral loads from Ct values  
> Comparison to datasets of symptomatic individuals  

**Findings:**  
> Distributions of viral loads are similar between asymptomatic and symptomatic populations  
> Large heterogeneity of Ct values (hence viral loads): at a given time point 2% of positive individuals host 90% of the circulated virions  

**Limitations:**  
The authors interpret the difference in viral loads as exclusively due to individual variation in peak viral load. However, the authors do control for the time since infection, and therefore should not dismiss the fact that viral loads changes over the course of infection, which mechanically leads to a whole range of Ct values in a population sampled at a given point in time.  

**Conclusions:**  
At a given point in time, the vast majority of circulating virions in communities are found within the bodies of a small number of individuals. |
| JAMA Netw Open 25MAY2021 | Association of Circulating Sex Hormones With Inflammation and Disease Severity in Patients With COVID-19 | Dhindsa S., et al. USA | Clinic | **Contribution of sex hormones to severe COVID-19 illness in men:** association of concentrations of serum testosterone, estradiol, and insulinlike growth factor 1 (IGF-1, concentrations of which are regulated by sex hormone signaling) with COVID-19 severity.  

**Methods**  
> Prospective cohort study  
> Collection of serum samples from patients with COVID-19 (diagnosed using nasopharyngeal swabs).  
> Testosterone, estradiol, and IGF-1 concentrations measured at the time of presentation (day 0) and at day 3, 7, 14, 28 after admission  

**Main Outcomes** Baseline hormone concentrations compared among patients who had severe COVID-19 vs those with mild COVID-19.  

**Findings**  
> Among 152 patients (90 [59.2%] men; 62 [40.8%] women; mean [SD] age, 63 [16] years), 143 patients (94.1%) were hospitalized.  
> Among 66 men with severe COVID-19, median testosterone concentrations were lower at day 0 (53 [18 to 114] ng/dL vs 151 [95 to 217] ng/dL; P = .01) and day 3 (19 [6 to 68] ng/dL vs 111 [49 to 274] ng/dL; P = .006) compared with 24 men with milder disease.  
> Testosterone concentrations were inversely associated with concentrations of interleukin 6 (β = −0.43; 95% CI, −0.52 to −0.17; P < .001), C-reactive protein (β = −0.38; 95% CI, −0.78 to −0.16; P = .004), interleukin 1 receptor antagonist (β = −0.29; 95% CI, −0.64 to −0.06; P = .02), hepatocyte growth factor (β = −0.46; 95% CI, −0.69 to −0.25; P < .001), and interferon γ–inducible protein 10 (β = −0.32; 95% CI, −0.62 to −0.10; P = .007).  
> Estradiol and IGF-1 concentrations were not associated with COVID-19 severity in men.  
> Testosterone, estradiol, and IGF-1 concentrations were similar in women with and without severe COVID-19.  
> Gene set enrichment analysis revealed upregulated hormone signaling pathways in CD14+CD16- (ie, classical) monocytes and CD14–CD16+ (ie, nonclassical) monocytes in male patients with COVID-19 who needed intensive care unit treatment vs those who did not.  

**Conclusion**  
Lower testosterone concentrations during hospitalization were associated with increased disease severity and inflammation in men. |
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- Phase 3 multi-centre open-label, randomized 1:1:1:1:1, adaptive, controlled trial (DisCoVeRy).  
- Results for the lopinavir/ritonavir-containing arms and for the hydroxychloroquine arm, stopped prematurely, are reported.  
Primary outcome: clinical status at day 15, measured by the WHO 7-point ordinal scale.  
Secondary outcomes included SARS-CoV-2 quantification in respiratory specimens, pharmacokinetic and safety analyses.  
Results  
> 583 participants (lopinavir/ritonavir, n=145; lopinavir/ritonavir-IFN-β-1a, n=145; hydroxychloroquine, n=145; control, n=148), 418 (71.7%) males, median age 63yrs (IQR, 54-71), 211 (36.2%) had severe disease.  
> The day-15 clinical status was not improved with investigational treatments: lopinavir/ritonavir vs. control, adjusted odds ratio (aOR) 0.83, (95%CI 0.55-1.26, P=0.39); lopinavir/ritonavir-IFN-ß-1a vs. control, aOR 0.69 (95%CI 0.45-1.04, P=0.08); hydroxychloroquine vs. control, aOR 0.93 (95%CI 0.62-1.41, P=0.75).  
> No significant effect of investigational treatment was observed on SARS-CoV-2 clearance.  
> Trough plasma concentrations of lopinavir and ritonavir were higher than those expected, while those of hydroxychloroquine were those expected with the dosing regimen.  
> The occurrence of Serious Adverse Events was significantly higher in participants allocated to the lopinavir/ritonavir-containing arms.  
Conclusions  
In adults hospitalized for COVID-19, lopinavir/ritonavir, lopinavir/ritonavir-IFN-ß-1a and hydroxychloroquine did not improve the clinical status at day 15, nor SARS-CoV-2 clearance in respiratory tract specimens. |
- Vaccines with heart, kidney, liver, or pancreas transplants.  
- Spike protein antibodies monitored before and after vaccination.  
Findings  
> 950 patients of the 2666 within receiving at least 1 dose of an mRNA vaccine (BNT162b2 vaccine [Pfizer-BioNTech], n = 942; mRNA-1273 vaccine [Moderna], n = 8) and had anti–SARS-CoV-2 antibodies monitored. Fifty patients had vaccination without monitoring of antibodies, 80 patients were planned to be vaccinated within the month , and 257 patients declined the vaccine. No feedback from the remaining 1329 patients.  
> 895 of the 950 patients had an available serologic screening just before the first injection. Prevalence of anti–SARS-CoV-2 antibodies: 2.1% (95% CI, 1.3% to 3.3%; n = 19 of 895). Only 5 of the 19 patients who were seropositive previously had symptomatic COVID-19.  
> A total of 576 patients benefited from a second injection at day 28. The prevalence of anti–SARS-CoV-2 antibodies before the second injection was 6.4% (CI, 4.6% to 8.8%; n = 37 of 576).  
> In 367 patients who had a 4-week follow-up after the second dose, the prevalence of anti–SARS-CoV-2 antibodies increased from 1.4% (CI, 0.4% to 3.2%; n = 5 of 367) at baseline to 6.3% (CI, 4.0% to 9.3%; n = 23 of 367) at day 28 and 33.8% (CI, 29.0% to 38.9%; n = 124 of 367) 1 month after the second dose  
> The tolerance of mRNA vaccines was excellent, with no serious adverse events reported, except in 1 patient with a liver transplant who developed paresthesia of the lower limb.  
Conclusion  
> In immunocompromised patients, such as recipients of SOT, a weak humoral response to mRNA vaccines is reported  
> Recipients of liver transplant showed a better humoral response than recipients of other organs. |
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| Science Immunol. 25MAY2021 | SARS-CoV-2 variants of concern partially escape humoral but not T-cell responses in COVID-19 convalescent donors and vaccinees | Geers D., et al. Netherlands gotopaper | Immunology | **Aim:** to study humoral and cellular immune responses to wild type SARS-CoV-2 and the B.1.1.7 and B.1.351 variants of concern in a cohort of 121 BNT162b2 mRNA-vaccinated health care workers (HCW) who did not have evidence of infection before vaccination.  
> Twenty-three HCW recovered from mild COVID-19 disease and exhibited a recall response with high levels of SARS-CoV-2-specific functional antibodies and virus-specific T cells after a single vaccination.  
> Specific immune responses were also detected in seronegative HCW after one vaccination, but a second dose was required to reach high levels of functional antibodies and cellular immune responses in all individuals.  
> Vaccination-induced antibodies cross-neutralized the variants B.1.1.7 and B.1.351, but the neutralizing capacity and Fc-mediated functionality against B.1.351 was consistently 2- to 4-fold lower than to the homologous virus.  
> Peripheral blood mononuclear cells were stimulated with peptide pools spanning the mutated S regions of B.1.1.7 and B.1.351 to detect cross-reactivity of SARS-CoV-2-specific T cells with variants. **No differences in CD4+ T-cell activation in response to variant antigens was observed**, indicating that the B.1.1.7 and B.1.351 S proteins do not escape T-cell-mediated immunity elicited by wild type S protein.  
Some variants can partially escape humoral immunity induced by SARS-CoV-2 infection or BNT162b2 vaccination, but S-specific CD4+ T-cell activation is not affected by the mutations in the B.1.1.7 and B.1.351 variants. |

| Science 25MAY2021 | Estimating infectiousness throughout SARS-CoV-2 infection course | Jones T.C., et al. Germany / USA gotopaper | Virology | **Aim:** to analyse viral load and whether samples yield a replicating virus isolate in cell culture (parameters for quantifying viral infection and shedding).  
Sample: 25,381 German SARS-CoV-2 cases, including 6110 from test centres attended by pre-symptomatic, asymptomatic, and mildly-symptomatic (PAMS) subjects, 9519 who were hospitalised, and 1533 B.1.1.7 lineage infections.  
> The youngest had mean log10 viral load 0.5 (or less) lower than older subjects and an estimated ~78% of the peak cell culture replication probability, due to smaller swab sizes, unlikely to be clinically relevant.  
> Viral loads above 109 copies per swab were found in 8% of subjects, one-third of whom were PAMS, with mean age 37.6.  
> PAMS subjects in apparently-healthy groups can be expected to be as infectious as hospitalised patients at the time of detection.  
> Estimate of 4.3 days from onset of shedding to peak viral load (8.1) and cell culture isolation probability (0.75).  
> B.1.1.7 subjects had mean log10 viral load 1.05 higher than non-B.1.1.7, with estimated cell culture replication probability 2.6 times higher.  
**Accurate estimations can be directly obtained from two easily-measured virological parameters, viral load and sample cell culture infectivity.** |

| Cell 24MAY2021 | An infectivity-enhancing site on the SARS-CoV-2 spike protein targeted by antibodies | Liu Y., et al. Japan gotopaper | Virology | The effects of antibodies against spike protein domains other than the RBD are largely unknown.  
> Screening of a series of anti-spike monoclonal antibodies from COVID-19 patients showed that some of antibodies against the N-terminal-domain (NTD) induced the open conformation of receptor binding domain (RBD) and thus enhanced the binding capacity of the spike protein to ACE2 and infectivity of SARS-CoV-2.  
> Mutational analysis revealed that all the infectivity-enhancing antibodies recognized a specific site on the NTD.  
> Structural analysis demonstrated that all the infectivity-enhancing antibodies bound to NTD in a similar manner.  
> Divalent bridging of spikes is required to induce RBD-up sate.  
> The antibodies against this infectivity-enhancing site were detected at high levels in severe patients.  
> Antibodies against the infectivity-enhancing site were identified in uninfected donors, albeit at a lower frequency.  
**These findings demonstrate that not only neutralizing antibodies but also enhancing antibodies are produced during SARS-CoV-2 infection.** |
**Nature 24MAY2021**

**Title:** SARS-CoV-2 infection induces long-lived bone marrow plasma cells in humans

**Authors and link:** Turner J.S., et al. USA

**Field of expertise:** Immunology

**Key facts:**

**Aim:** to determine whether SARS-CoV-2 infection induces antigen-specific long-lived bone marrow plasma cells (BMPCs) in humans.

**Methods:**
Blood samples were collected approximately 1 month after onset of symptoms from 77 SARS-CoV-2 convalescent volunteers (49% female, 51% male, median age 49), the majority of whom had experienced mild illness (7.8% hospitalized).
Follow-up blood samples were collected three times at approximately 3-month intervals.
Additionally, bone marrow aspirates were collected from 18 of the participants 7 to 8 months after infection and from 11 healthy volunteers with no history of SARS-CoV-2 infection or vaccination.

**Findings:**
> In patients who experienced mild infections (n=77), serum anti-SARS-CoV-2 spike (S) antibodies decline rapidly in the first 4 months after infection and then more gradually over the following 7 months, remaining detectable at least 11 months after infection.
> Anti-S antibody titers correlated with the frequency of S-specific BMPCs obtained from bone marrow aspirates of 18 SARS-CoV-2 convalescent patients 7 to 8 months after infection. S-specific BMPCs were not detected in aspirates from 11 healthy subjects with no history of SARS-CoV-2 infection.
> S-binding BMPCs are quiescent, indicating that they are part of a long-lived compartment. Consistently, circulating resting memory B cells directed against the S protein were detected in the convalescent individuals.

SARS-CoV-2 infection induces a robust antigen-specific, long-lived humoral immune response in humans.

**Clin Microbiol Infect. 23MAY2021**

**Title:** Clinical outcomes in COVID-19 patients infected with different SARS-cov-2 variants in marseille, France

**Authors and link:** Dao T.L., et al. France

**Field of expertise:** Virology

**Key facts:**

**Clinical and epidemiological aspects associated with different predominant lineages circulating in Marseille from March 2020 to January 2021.**

**Methods**
> Single-center retrospective cohort study
> Characteristics of patients infected with four different SARS-CoV-2 variants were documented from medical files.

**Outcome:** occurrence of clinical failure, defined as hospitalization (for outpatients), transfer to the intensive-care unit (inpatients), death (all)

**Findings**
> 254 patients were infected with clade 20A (20AS), 85 with Marseille-1 (M1V), 190 with Marseille-4 (M4V) and 211 with N501Y (N501YV) variants.
(i) 20AS presented a bell-shaped epidemiological curve and nearly disappeared around May 2020.
(ii) M1V reached a very weak peak, then disappeared after a month-and-a-half.
(iii) M4V appeared in July presented an atypical wave form during seven months.
(iv) N501YV was only recently appeared.
> As compared to 20AS, patients infected with M1V were less likely to report dyspnoea (aOR=0.50, p=0.04), rhinitis (aOR=0.57, p=0.04) and to be hospitalised (aOR=0.22, p=0.002).
> Patients infected with M4V were more likely to report fever than those with 20AS and M1V (aOR=2.49, p<0.0001 and aOR=2.30, p=0.007, respectively) and to be hospitalised than those with M1V (aOR=4.81, p=0.003).
> Patients infected with N501YV reported lower rate of rhinitis (aOR=0.50, p=0.001) and anosmia (aOR=0.57, p=0.02), as compared to those infected with 20AS.
> A lower rate of hospitalisation associated with N501YV infection as compared to 20AS and M4V (aOR=0.33, p=0.0001 and aOR=0.27, p=0.0001, respectively).

**Conclusions**
The four lineages have presentations which differ from one other, epidemiologically and clinically. This supports SARS-CoV-2 genomic surveillance through next-generation sequencing.
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<td>Lancet 22MAY2021</td>
<td>Patient care and clinical outcomes for patients with COVID-19 infection admitted to African high-care or intensive care units (ACCOS): a multicentre, prospective, observational cohort study</td>
<td>ACCCOS Investigators Group International gotopaper</td>
<td>Public Health / Epidemiology</td>
<td>Resources, comorbidities, and critical care interventions are associated with mortality in critically ill African patients due to COVID-19. <strong>Methods</strong> &gt; Multicentre, prospective, observational cohort study in adults (aged 18 years or older) with suspected or confirmed COVID-19 infection referred to intensive care or high-care units in 64 hospitals in ten African countries (ie, Egypt, Ethiopia, Ghana, Kenya, Libya, Malawi, Mozambique, Niger, Nigeria, and South Africa). ClinicalTrials.gov, NCT04367207. Primary outcome: in-hospital mortality censored at 30 days. <strong>Findings</strong> &gt; 6779 patients referred to critical care. &gt; Of these, 3752 (55·3%) patients were admitted and 3140 (83·7%) patients from 64 hospitals in ten countries participated (mean age 55·6 years; 1890 [60·6%] of 3118 participants were male). &gt; In-hospital mortality within 30 days of admission was 48·2% (95% CI 46·4–50·0; 1483 of 3077 patients). &gt; Factors that were independently associated with mortality were (i) increasing age per year (odds ratio 1·03; 1·02–1·04); (ii) HIV/AIDS (1·91; 1·31–2·79); (iii) diabetes (1·25; 1·01–1·56); (iv) chronic liver disease (3·48; 1·48–8·18); (v) chronic kidney disease (1·89; 1·28–2·78); (vi) delay in admission due to a shortage of resources (2·14; 1·42–3·22); (vii) quick sequential organ failure assessment score at admission (for one factor [1·44; 1·01–2·04], for two factors [2·0; 1·33–2·99], and for three factors [3·66, 2·12–6·33]); (viii) respiratory support (high flow oxygenation [2·72; 1·46–5·08]; continuous positive airway pressure [3·93; 2·13–7·26]; invasive mechanical ventilation [15·27; 8·51–27·37]); (ix) cardiorespiratory arrest within 24 h of admission (4·43; 2·25–8·73); (x) vasopressor requirements (3·67; 2·77–4·86). &gt; Steroid therapy was associated with survival (0·55; 0·37–0·81). <strong>Conclusion</strong> Increased mortality was associated with insufficient critical care resources, as well as the comorbidities of HIV/AIDS, diabetes, chronic liver disease, and kidney disease, and severity of organ dysfunction at admission.</td>
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<td>Clin Infect Dis. 20MAY2021</td>
<td>Factors Associated with Readmission in the US Following Hospitalization with COVID-19</td>
<td>Verna E.C., et al. USA gotopaper</td>
<td>Clinic</td>
<td>Aim: to estimate the rate and risk factors associated with COVID-19-related readmission and inpatient mortality. <strong>Methods:</strong> Retrospective cohort study utilizing deidentified chargemaster data from 297 hospitals across 40 US states on patients hospitalized with COVID-19 February 15–June 09, 2020. Multivariable logistic regression was used to measure risk factor associations with 30-day readmission and in-hospital mortality. <strong>Findings:</strong> &gt; Among 29,659 patients, 1,070 (3.6%) were readmitted. &gt; Readmitted patients were more likely to have diabetes, hypertension, cardiovascular disease (CVD), chronic kidney disease (CKD) vs those not readmitted and to present on first admission with acute kidney injury (15.6% vs. 9.2%), congestive heart failure (6.4% vs. 2.4%), and cardiomyopathy (2.1% vs. 0.8%). &gt; Higher odds of readmission were observed in patients age &gt;60 vs. 18 to 40 (odds ratio [OR]=1.92), and admitted in the Northeast vs. West (OR=1.43) or South (OR=1.28). &gt; Comorbidities including diabetes (OR=1.34), CVD (OR=1.46), CKD stage 1-5 (OR=1.51) and stage 5 (OR=2.27) were associated with higher odds of readmission. &gt; 12.3% of readmitted patients died during second hospitalization. Readmission was associated with certain comorbidities and acute conditions during first hospitalization.</td>
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| Science 20MAY2021 | Face masks effectively limit the probability of SARS-CoV-2 transmission | Cheng Y., et al. USA [gotopaper](#) | Public Health / Epidemiology | **Aim:** to develop a quantitative model of airborne virus exposure that can explain these contrasting results and provide a basis for quantifying the efficacy of face masks.  
**Methods:**  
The analysis was focused on respiratory particles and droplets with diameters smaller than 100 μm (traditional physical definition of aerosols).  
**Findings:**  
> Mask efficacy strongly depends on airborne virus abundance. Based on direct measurements of SARS-CoV-2 in air samples and population-level infection probabilities, the authors find that the virus abundance in most environments is sufficiently low for masks to be effective in reducing airborne transmission.  
> A person typically emits a total number of about $3 \times 10^6$ particles during a 30 min period. This very large number implies that indoor environments are usually in a respiratory particle-rich regime. Surgical masks with particle collection efficiencies around ~50% cannot prevent the release of millions of particles per person and their inhalation by others.  
> For SARS-CoV-2, the viral load of infectious individuals can vary by orders of magnitude. The authors find that most environments and contacts are under conditions of low virus abundance (virus-limited) where surgical masks are effective at preventing virus spread.  
> More advanced masks and other protective equipment are required in potentially virus-rich indoor environments including medical centers and hospitals.  
> Masks are particularly effective in combination with other preventive measures like ventilation and distancing. |
Exoproteome-targeting autoantibodies can exert a wide range of functional effects such as perturbation of cell signaling (as with the case of anti-IFN-I autoantibodies11,12) and targeted killing of specific cell populations via Fc receptors (FcR) and/or complement.  
**Methods:**  
A high-throughput autoantibody (AAb) discovery technique called Rapid Extracellular Antigen Profiling (REAP)7 was used to screen the cohort.  
**Findings:**  
> COVID-19 patients exhibit dramatic increases in autoantibody reactivities compared to uninfected controls, with a high prevalence of autoantibodies against immunomodulatory proteins including cytokines, chemokines, complement components, and cell surface proteins.  
> These autoantibodies perturb immune function and impair virological control by inhibiting immunoreceptor signaling and by altering peripheral immune cell composition.  
> Murine surrogates of these autoantibodies exacerbate disease severity in a mouse model of SARS-CoV-2 infection.  
> Analysis of autoantibodies against tissue-associated antigens revealed associations with specific clinical characteristics and disease severity.  
These findings implicate a pathological role for exoproteome-directed autoantibodies in COVID-19 with diverse impacts on immune functionality and associations with clinical outcomes. |
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| BMJ 19MAY2021    | Risk of clinical sequelae after the acute phase of SARS-CoV-2 infection: retrospective cohort study | Daugherty S.E., et al. USA [gotopaper](#) | Clinic | **Aim:** to evaluate the excess risk and relative hazards for developing incident clinical sequelae after the acute phase of SARS-CoV-2 infection in adults aged 18-65  
**Methods:**  
Retrospective cohort study from individuals aged 18-65 with continuous enrollment in the health plan from January 2019 to the date of a diagnosis of SARS-CoV-2 infection.  
Three merged data sources from a large United States health plan: a large national administrative claims database, an outpatient laboratory testing database, and an inpatient hospital admissions database.  
**Findings:**  
>14% of adults aged ≤65 who were infected with SARS-CoV-2 (27,074 of 193,113) had at least one new type of clinical sequelae that required medical care after the acute phase of the illness, which was 4.95% higher than in the 2020 comparator group.  
The risk for specific new sequelae attributable to SARS-CoV-2 infection after the acute phase, including chronic respiratory failure, cardiac arrhythmia, hypercoagulability, encephalopathy, peripheral neuropathy, amnesia (memory difficulty), diabetes, liver test abnormalities, myocarditis, anxiety, and fatigue, was significantly greater than in the three comparator groups (2020, 2019, and viral lower respiratory tract illness groups).  
Significant risk differences because of SARS-CoV-2 infection ranged from 0.02 to 2.26 per 100 people, and hazard ratios ranged from 1.24 to 25.65 compared with the 2020 comparator group.  
Individuals who were older, had pre-existing conditions, and were admitted to hospital because of covid-19 were at greatest excess risk.  
Younger adults (aged ≤50), those with no pre-existing conditions, or those not admitted to hospital for covid-19 also had an increased risk of developing new clinical sequelae. |
| JAMA Netw Open 19MAY2021 | Assessment of the Association of Vitamin D Level With SARS-CoV-2 Seropositivity Among Working-Age Adults | Li Y., et al. USA [gotopaper](#) | Public Health / Epidemiology | **Aim:** to examine whether low levels of vitamin D (<20 ng/mL or < 30 ng/mL) are associated with SARS-CoV-2 seropositivity, an indicator of previous infection.  
**Results:**  
The 18,148 individuals included in this study had test results for SARS-CoV-2 IgG in 2020 and vitamin D levels from the prepandemic and pandemic periods. Median (interquartile range) age was 47 (37-56) years, 12,170 (67.1%) were women, 900 (5.0%) were seropositive, 4,498 (24.8%) had a vitamin D level <20 ng/mL, and 10,876 (59.9%) had a vitamin D level <30 ng/mL before the pandemic.  
In multivariable models adjusting for age, sex, race/ethnicity, education, body mass index, blood pressure, smoking status, and geographical location, SARS-CoV-2 seropositivity was not associated with having a vitamin D level <20 ng/mL before (odds ratio [OR], 1.04; 95% CI, 0.88-1.22) or during (OR, 0.93; 95% CI, 0.79-1.09) the pandemic; it was also not associated with having a vitamin D level <30 ng/mL before (OR, 1.09; 95% CI, 0.93-1.27) or during (OR, 1.05; 95% CI, 0.91-1.23) the pandemic. Similar results were observed in propensity score analyses.  
SARS-CoV-2 seropositivity was associated with obesity (OR, 1.26; 95% CI, 1.08-1.46), not having a college degree (OR, 1.40; 95% CI, 1.21-1.62), and Asian (OR, 1.46; 95% CI, 1.13-1.87), Black (OR, 2.74; 95% CI, 2.25-3.34), Hispanic (OR, 2.65; 95% CI, 2.15-3.27), American Indian or Alaska Native, and Native Hawaiian or other Pacific Islander (OR, 2.01; OR, 1.54-2.62) race/ethnicity, and was inversely associated with high blood pressure (OR, 0.82; 95% CI, 0.70-0.96), smoking (OR, 0.60; 95% CI, 0.47-0.78), and residing in the US Northeast (OR, 0.75; 95% CI, 0.62-0.92) and West (OR, 0.54; 95% CI, 0.44-0.67).  
**Conclusions**  
In this cohort study, SARS-CoV-2 seropositivity was not associated with low levels of vitamin D independently of other risk factors. |
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<td>Nature Med. 18MAY2021</td>
<td>Phase 1 randomized trial of a plant-derived virus-like particle vaccine for COVID-19</td>
<td>Ward B.J., et al. Canada gotopaper</td>
<td>Vaccines</td>
<td>Safety and immunogenicity data from the virus-like particle vaccine candidate produced by MEDICAGO. (CoVLP: NCT04450004). <strong>Methods</strong>  &gt; Phase 1 blinded, dose escalation, randomized controlled study of a virus-like particle vaccine candidate produced in plants that displays the SARS-CoV-2 spike glycoprotein  &gt; Adults (18–55 years, n = 180) receiving two intramuscular doses of CoVLP (3.75 μg, 7.5 μg, and 15 μg) 21 d apart, alone or adjuvanted with AS03 or CpG1018 or placebo. <strong>Primary outcomes:</strong> short-term tolerability/safety and immunogenicity of CoVLP formulations assessed by neutralizing antibody (NAb) and cellular responses. <strong>Findings</strong>  &gt; All formulations were well tolerated, and adverse events after vaccination were generally mild to moderate, transient and highest in the adjuvanted groups.  &gt; No CoVLP dose effect on serum NAbs, but titers increased significantly with both adjuvants.  &gt; After the second dose, NAbs in the CoVLP + AS03 groups were more than tenfold higher than titers in Coronavirus 2019 convalescent sera.  &gt; Spike protein-specific interferon-γ and interleukin-4 cellular responses were also induced.</td>
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<td>Nature Med. 18MAY2021</td>
<td>Neutralizing antibody levels are highly predictive of immune protection from symptomatic SARS-CoV-2 infection</td>
<td>Khoury D.S., et al. Australia gotopaper</td>
<td>Immunology</td>
<td>Analysis of the relationship between in vitro neutralization levels and observed protection from SARS-CoV-2 infection (data from seven current vaccines and from convalescent cohorts) <strong>Findings</strong>  &gt; The neutralization level for 50% protection against detectable SARS-CoV-2 infection to be 20.2% of the mean convalescent level (95% confidence interval (CI) = 14.4–28.4%).  &gt; Estimated neutralization level required for 50% protection from severe infection was significantly lower (3% of the mean convalescent level; 95% CI = 0.7–13%, P = 0.0004).  &gt; Modeling of the decay of the neutralization titer over the first 250 d after immunization predicts that a significant loss in protection from SARS-CoV-2 infection will occur, although protection from severe disease should be largely retained.  &gt; Neutralization titers against some SARS-CoV-2 variants of concern are reduced compared with the vaccine strain. <strong>Conclusion</strong> Neutralization level is highly predictive of immune protection, and provide an evidence-based model of SARS-CoV-2 immune protection that will assist in developing vaccine strategies to control the future trajectory of the pandemic.</td>
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<td>Science Immunol. 18MAY2021</td>
<td>Pharmacological activation of STING blocks SARS-CoV-2 infection</td>
<td>Minghua L., et al. USA gotopaper</td>
<td>Therapeutics</td>
<td>Since pretreatment with IFNs can block viral infection, we reasoned that pharmacological activation of innate immune pathways could control SARS-CoV-2 infection <strong>Methods</strong>  &gt; To identify potent antiviral innate immune agonists, we screened a panel of 75 microbial ligands that activate diverse signaling pathways and identified cyclic dinucleotides (CDNs), canonical STING agonists, as antiviral. <strong>Findings</strong>  &gt; SARS-CoV-2 evades interferon (IFN) activation in respiratory epithelial cells, resulting in a delayed response in bystander cells.  &gt; Since CDNs have poor bioavailability, we tested the small molecule STING agonist diABZI, and found that it potently inhibits SARS-CoV-2 infection of diverse strains including variants of concern (B.1.351) by transiently stimulating IFN signaling.  &gt; Importantly, diABZI restricts viral replication in primary human bronchial epithelial cells and in mice in vivo. Our study provides evidence that activation of STING may represent a promising therapeutic strategy to control SARS-CoV-2.</td>
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| Science Immunol. 18MAY2021 | A diamidobenzimidazole STING agonist protects against SARS-CoV-2 infection | Humphries F., et al. USA gotopaper | Therapeutics | Describe a diamidobenzimidazole compound: diABZI-4

**Findings**

> diABZ-4 activates STING and is highly effective in limiting SARS-CoV-2 replication in cells and animals.
> diABZI-4 inhibited SARS-CoV-2 replication in lung epithelial cells.
> Administration of diABZI-4 intranasally before or even after virus infection conferred complete protection from severe respiratory disease in K18-ACE2-transgenic mice infected with SARS-CoV-2.
> Intranasal delivery of diABZI-4 induced a rapid short-lived activation of STING, leading to transient proinflammatory cytokine production and lymphocyte activation in the lung associated with inhibition of viral replication.

**Our study supports the use of diABZI-4 as a host-directed therapy which mobilizes antiviral defenses for the treatment and prevention of COVID-19.** |

**Primary outcome:** incidence rate of a SARS-CoV-2 infection confirmed with rt-PCR, between 7 to 27 days after second dose (protection-period), as compared to days 1 to 7 after the first dose, where no protection by the vaccine is assumed (reference-period).

**Results**

> Data of 1,178,597 individuals vaccinated with BNT162b2 were analyzed (mean age 47.7 years [SD=18.1], 48.4% males) of whom 872,454 (74.0%) reached the protection period.
> Overall, 4514 infections occurred during the reference period compared to 728 during the protection period, yielding a weighted mean daily incidence of 5.48 per 100,000 (95% CI: 26.1-115.0 per 100,000) and 5.4 per 100,000 (95% CI: 3.5-8.4 per 100,000), respectively.
> The vaccine effectiveness in preventing infection was 90% (95%CI: 79%-95%) and 94% (95%CI: 88%-97%) against COVID-19.
> Among immunosuppressed patients, vaccine effectiveness against infection was 71% (95%CI: 37%-87%).
> The adjusted hazard ratios for hospitalization in those infected were 0.82 (95%CI:0.36-1.88), 0.45 (95%CI:0.23-0.90), and 0.56 (95%CI:0.36-0.89) in the age groups 16-44, 45-64 and ≥75, respectively.

**The effectiveness of the BNT162b2 vaccine is comparable to the one reported in the phase III clinical trial.** |
| Blood 14MAY2021 | Frequency of positive anti-PF4/polyanion antibody tests after COVID-19 vaccination with ChAdOx1 nCoV-19 and BNT162b2 | Thiele T., et al. Germany gotopaper | Vaccines | - Vaccination with COVID-19 vaccine ChAdOx1 nCoV-19 (AstraZeneca) has been associated with rare vaccine-induced immune thrombotic thrombocytopenia (VITT).
- Affected patients test strongly positive in PF4/polyanion enzyme immunoassays (EIAs) and serum-induced platelet activation is maximal in the presence of PF4.

**Aim:** to determine the frequency of anti-PF4/polyanion antibodies in healthy vaccines and to assess if PF4/polyanion EIA-positive sera exhibit platelet-activating properties after vaccination with ChAdOx1 nCoV-19 (n=138) or BNT162b2 (BioNTech/Pfizer; n=143).

> 19 of 281 participants tested positive for anti-PF4/polyanion antibodies post-vaccination (All: 6.8% [95%CI, 4.4-10.3]; BNT162b2: 5.6% [95%CI, 2.9-10.7]; ChAdOx1 nCoV-19: 8.0% [95%CI, 4.5-13.7%]).
> Optical densities were mostly low (between 0.5-1.0 units; reference range, <0.50) and none of the PF4/polyanion EIA-positive samples induced platelet activation in the presence of PF4.

**Positive PF4/polyanion EIAs can occur after SARS-CoV-2 vaccination with both mRNA- and adenoviral vector-based vaccines, but the majority of these antibodies likely have minor (if any) clinical relevance. Pathogenic platelet-activating antibodies that cause VITT do not occur commonly following vaccination.** |
### Changes in in-hospital mortality in the first wave of COVID-19: a multicentre prospective observational cohort study using the WHO Clinical Characterisation Protocol

**Aim:** To quantify potential drivers of mortality rates in in-hospital patients and to identify groups of patients who remain at high risk of dying in hospital.

**Methods:**
- A three-way decomposition mediation analysis using natural effects models to explore associations between week of admission and in-hospital mortality was performed.

**Findings:**
- 80,713 patients were recruited, of whom 63,972 were eligible and included in the study.
- Unadjusted weekly in-hospital mortality declined from 32.3% in March 9 to April 26, 2020, to 16.4% in June 15 to Aug 2, 2020.
- Reductions in mortality were observed in all age groups, in all ethnic groups, for both sexes, and in patients with and without comorbidities.
- After adjustment, there was a 32% reduction in the risk of mortality per 7-week period (odds ratio [OR] 0.68).
- The higher proportions of patients with severe disease and comorbidities earlier in the first wave (March and April) than in June and July accounted for 10.2% of this reduction.
- Changes in respiratory support and use of steroids accounted for 22.2%, OR 0.95 (0.94–0.95) of the reduction in in-hospital mortality.

A significant reduction in in-hospital mortality was associated with differences in respiratory support and critical care use, which could partly reflect accrual of clinical knowledge.

### Convalescent plasma in patients admitted to hospital with COVID-19 (RECOVERY): a randomised controlled, open-label, platform trial

**Aim:** To evaluate the safety and efficacy of convalescent plasma therapy in patients admitted to hospital with COVID-19.

**Primary outcome:** 28-day mortality, analysed on an intention-to-treat basis.

**Findings:**
- 11558 (71%) of 16287 patients enrolled in RECOVERY were assigned to either the convalescent plasma group or the usual care group (May 28, 2020, and Jan 15, 2021).
- There was no significant difference in 28-day mortality between the two groups: 1399 (24%) of 5795 patients in the convalescent plasma group and 1408 (24%) of 5763 patients in the usual care group died within 28 days (rate ratio 1.00, 95% CI 0.93–1.07; p=0.95).
- The 28-day mortality rate ratio was similar in all prespecified subgroups of patients, including in those patients without detectable SARS-CoV-2 antibodies at randomisation.
- Allocation to convalescent plasma had no significant effect on the proportion of patients discharged from hospital within 28 days (3832 [66%] patients in the convalescent plasma group vs 3822 [66%] patients in the usual care group; rate ratio 0.99, 95% CI 0.94–1.03; p=0.57).
- Among those not on invasive mechanical ventilation at randomisation, there was no significant difference in the proportion of patients meeting the composite endpoint of progression to invasive mechanical ventilation or death (1568 [29%] of 5493 patients in the convalescent plasma group vs 1568 [29%] of 5448 patients in the usual care group; rate ratio 0.99, 95% CI 0.93–1.05; p=0.79).

In patients hospitalised with COVID-19, high-titre convalescent plasma did not improve survival or other prespecified clinical outcomes.
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**Methods**
- Case-control study.
- 156,930 adults aged 70 years and older who reported symptoms of covid-19 between 8 December 2020 and 19 February 2021 and were successfully linked to vaccination data in the National Immunisation Management System.
- Interventions: Vaccination with BNT162b2 or ChAdOx1-S.

**Main outcome** PCR confirmed symptomatic SARS-CoV-2 infections, admissions to hospital for covid-19, and deaths with covid-19.

**Findings**
- Participants aged 80 years and older vaccinated with BNT162b2 before 4 January 2021 had a higher odds of testing positive for covid-19 in the first nine days after vaccination (odds ratio up to 1.48, 95% confidence interval 1.23 to 1.77). Vaccine effects were noted 10 to 13 days after vaccination, reaching a vaccine effectiveness of 70% (95% confidence interval 59% to 78%). From 14 days after the second dose the vaccination effectiveness was estimated to 89% (85% to 93%). Vaccine effectiveness reached 61% (51% to 69%) from 28 to 34 days after vaccination, then plateaued.
- With ChAdOx1-S, effects were seen from 14 to 20 days after vaccination, reaching an effectiveness of 60% (41% to 73%) from 28 to 34 days, increasing to 73% (27% to 90%) from day 35 onwards.
- Further 43% (33% to 52%) reduced risk of emergency hospital admission and 51% (37% to 62%) reduced risk of death was observed in those who had received one dose of BNT162b2.
- Participants who had received one dose of ChAdOx1-S had a further 37% (3% to 59%) reduced risk of emergency hospital admission. Follow-up was insufficient to assess the effect of ChAdOx1-S on mortality.
- Combined with the effect against symptomatic disease, a single dose of either vaccine was about 80% effective at preventing admission to hospital with covid-19 and a single dose of BNT162b2 was 85% effective at preventing death with covid-19.

**Conclusion**
Vaccination with either one dose of BNT162b2 or ChAdOx1-S was associated with a significant reduction in symptomatic covid-19 in older adults, and with further protection against severe disease. Both vaccines showed similar effects. Protection was maintained for the duration of follow-up (>6 weeks). A second dose of BNT162b2 was associated with further protection against symptomatic disease. A clear effect of the vaccines against the B.1.1.7 variant was found.
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| Nature 12MAY2021 | The epidemiological impacts of the NHS COVID-19 App | Wymant C., et al. UK gotopaper | Public Health / Epidemiology | **Aim:** Investigate the impact of the NHS COVID-19 app for England and Wales, from its launch on 24 September 2020 through to the end of December 2020, using data aggregated at the level of local authorities.  

**Methods:**  
> Estimation of the secondary attack rate (SAR) in app-notified individuals based on a probabilistic model for how many positive test results would be expected among those recently notified.  
> Evaluation of the number of cases averted by the app based on notifications and SAR, using a mechanistic probabilistic modelling approach.  
> Evaluation of the number of cases averted by the app using a stratified statistical approach, allowing to address confounding factors.  

**Findings:** 16.5 million users (28% of the total population) sent approximately 1.7 million exposure notifications, 4.4 per index case consenting to contact tracing.  
> SAR was estimated at 6.0%, comparable to the SAR for manually traced close contacts.  
> Modelling based on the notifications and SAR gave 284,000 (108,000-450,000) cases averted by the app.  
> Statistical comparison of matched neighbouring local authorities gave 594,000 (317,000-914,000) cases averted by the app.  
> For every percentage point increase in app users, the number of cases can be reduced by 0.8% (modelling) or 2.3% (statistical analysis).  

**Limitation:**  
> It is an observational study: no randomized or systematic experiment resulted in different app uptake in different places.  

**Conclusion:**  
> These findings provide evidence for continued development and deployment of privacy-preserving contact tracing apps in populations that are awaiting full protection from vaccines.  
> Digital tracing is best understood as part of a system of non-pharmaceutical interventions, not in isolation. Also, it is not a substitute for manual tracing, both being valuable. |
| BMJ 11MAY2021 | Use of repurposed and adjuvant drugs in hospital patients with covid-19: multinational network cohort study | Prats-Uribe A., et al. UK / USA gotopaper | Therapeutics | **Aim:** to investigate the use of repurposed and adjuvant drugs in patients admitted to hospital with covid-19 across three continents.  

**Methods:**  
> Multinational network cohort study.  
> Data collected from Hospital electronic health records from the United States, Spain, and China, and nationwide claims data from South Korea.  
> 303 264 patients admitted to hospital with covid-19 from January 2020 to December 2020  

**Findings:**  
> Of the 303 264 patients included, 290 131 were from the US, 7599 from South Korea, 5230 from Spain, and 304 from China.  
> 3455 drugs were identified.  
> Common repurposed drugs were hydroxychloroquine (used in from ≤5 (<2%) patients in China to 2165 (85.1%) in Spain), azithromycin (from 15 (4.9%) in China to 1473 (57.9%) in Spain), combined lopinavir and ritonavir (from 156 (<2%) in the VA-OMOP US to 2,652 (34.9%) in South Korea and 1285 (50.5%) in Spain), and umifenovir (0% in the US, South Korea, and Spain and 238 (78.3%) in China).  
> Use of adjunctive drugs varied greatly, with the five most used treatments being enoxaparin, fluoroquinolones, ceftriaxone, vitamin D, and corticosteroids.  
> Hydroxychloroquine use increased rapidly from March to April 2020 but declined steeply in May to June and remained low for the rest of the year.  
> The use of dexamethasone and corticosteroids increased steadily during 2020. |
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**3 cohorts:**  
- GCS-NeuroCOVID all COVID-19 – hospitalized patients with COVID-19 with and without neurological manifestations (n = 3055; 57% men, mean age 59.9 years [95% CI, 59.3-60.6]).  
- GCS-NeuroCOVID COVID-19 neurological cohort – patients hospitalized with COVID-19 who had confirmed neurological manifestations (n = 475; 55% men, and the mean age 62.6 [61.1-64.1]).  
- ENERGY cohort – patients with COVID-19 who received formal neurological consultation. (n = 214; 62% men, mean age 67 years [52-78]).  
**Results**  
> A total of 3083 of 3743 patients (82%) across cohorts had any neurological manifestation (self-reported neurological symptoms and/or clinically captured neurological sign and/or syndrome).  
> The most common self-reported symptoms included headache (1385 of 3732 patients [37%]) and anosmia or ageusia (977 of 3700 patients [26%]).  
> The most prevalent neurological signs and/or syndromes were acute encephalopathy (1845 of 3740 patients [49%]), coma (649 of 3737 patients [17%]), and stroke (222 of 3737 patients [6%]), while meningitis and/or encephalitis were rare (19 of 3741 patients [0.5%]).  
> Presence of clinically captured neurologic signs and/or syndromes was associated with increased risk of in-hospital death (adjusted odds ratio [aOR], 5.99; 95% CI, 4.33-8.28) after adjusting for study site, age, sex, race, and ethnicity.  
> Presence of preexisting neurological disorders (aOR, 2.23; 95% CI, 1.80-2.75) was associated with increased risk of developing neurological signs and/or syndromes with COVID-19.  
**Conclusions**  
In this multicohort study, neurological manifestations were prevalent among patients hospitalized with COVID-19 and were associated with higher in-hospital mortality. |
| Nature Med. 11MAY2021 | Neutralizing antibody responses to SARS-CoV-2 in symptomatic COVID-19 is persistent and critical for survival | Dispinseri S., et al. Italy gotopaper | Immunology | **Antibody responses of 162 COVID-19 symptomatic patients in the COVID-BioB cohort followed longitudinally for up to eight months from symptom onset to find SARS-CoV-2 neutralization, as well as antibodies either recognizing SARS-CoV-2 spike antigens and nucleoprotein, or specific for S2 antigen of seasonal beta-coronaviruses and hemagglutinin of the H1N1 flu virus.**  
**Findings**  
> The presence of neutralizing antibodies within the first weeks from symptoms onset correlates with time to a negative swab result (p = 0.002). The lack of neutralizing capacity correlates with an increased risk of a fatal outcome (p = 0.008).  
> Neutralizing antibody titers progressively drop after 5–8 weeks but are still detectable up to 8 months in the majority of recovered patients regardless of age or co-morbidities, with IgG to spike antigens providing the best correlate of neutralization.  
> Antibody responses to seasonal coronaviruses are temporarily boosted, and parallel those to SARS-CoV-2 without dampening the specific response or worsening disease progression.  
**Conclusions**  
Compromised immune responses to the SARS-CoV-2 spike to be a major trait of COVID-19 patients with critical conditions, and thereby inform on the planning of COVID-19 patient care and therapy prioritization. |
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<tr>
<td>Clin Infect Dis.</td>
<td>Trends over time in the risk of adverse outcomes among patients with SARS-CoV-2 infection</td>
<td>Loannou G.N., et al. USA</td>
<td>Public Health / Epidemiology</td>
<td>We aimed to describe trends in adverse outcomes among patients who tested positive for SARS-CoV-2 between February and September 2020 within a national healthcare system.</td>
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<td>11MAY2021</td>
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<td>gotopaper</td>
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<td><strong>Methods</strong>  &gt; Identified enrollees in the national U.S. Veterans Affairs healthcare system who tested positive for SARS-CoV-2 between 2/28/2020 and 9/30/2020 (n=55,952), with follow-up extending to 11/19/2020  &gt; Determined trends over time in incidence of the following outcomes that occurred within 30 days of testing positive: hospitalization, intensive care unit (ICU) admission, mechanical ventilation and death.</td>
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<td><strong>Findings</strong>  &gt; Between February and July 2020, there were marked <em>downward trends</em> in the 30-day incidence of hospitalization (44.2% to 15.8%), ICU admission (20.3% to 5.3%), mechanical ventilation (12.7% to 2.2%), and death (12.5% to 4.4%), which subsequently plateaued between July and September 2020.  &gt; These trends persisted after adjustment for sociodemographic characteristics, comorbid conditions, documented symptoms and laboratory tests, including among subgroups of patients hospitalized, admitted to the ICU or treated with mechanical ventilation.  &gt; From February to September, there were decreases in the use of hydroxychloroquine (56.5% to 0%), azithromycin (48.3% to 16.6%) vasopressors (20.6% to 8.7%), and dialysis (11.6% to 3.8%) and increases in the use of dexamethasone (3.4% to 53.1%), other corticosteroids (4.9% to 29.0%) and remdesivir (1.7% to 45.4%) among hospitalized patients.  &gt; The risk of adverse outcomes in SARS-CoV-2-positive patients decreased markedly between February and July, with subsequent stabilization from July to September. These trends were not explained by changes in measured baseline patient characteristics and may reflect changing treatment practices or viral pathogenicity.</td>
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<td>Clin Microbiol Infect.</td>
<td>Persistent COVID-19 symptoms are highly prevalent 6 months after hospitalization: results from a large prospective cohort</td>
<td>Ghosn J., et al. France</td>
<td>Public Health / Epidemiology</td>
<td>We have assessed, in the longitudinal prospective French COVID-19 cohort, symptoms that persisted 6 months after admission for COVID-19.  &gt; M6 data were available for 1137 participants (Hospitalized patients with virologically confirmed COVID-19). Median age was 61 years (IQR 51–71) and 288 (29%, 95% CI 26–32%) were admitted to intensive care unit (ICU) during the acute phase.  &gt; 650 (68%, 95% CI 65–71%) and 639 (60%, 95% CI 57–63%) participants had at least one symptom at M3 and M6 visit, respectively, mostly fatigue, dyspnoea, joint pain and myalgia.  &gt; At M6, 255 (24%, 95% CI 21–27%) of participants had three or more persistent symptoms.  &gt; The presence of three or more symptoms at M6 was independently associated with female gender (adjusted odds ratio (aOR) 2.40, 95% CI 1.75–3.30), having three or more symptoms at admission (aOR 2.04, 95% CI 1.45–2.89) and ICU admission/transfer during acute phase (aOR 1.55, 95% CI 1.09–2.18), but not significantly with age or having two or more comorbidities.  &gt; 125 (29%, 95% CI 25–34%) of those who initially had a professional occupation were not back to work at M6.  &gt; A fourth of individuals admitted to hospital for COVID-19 still had three or more persistent symptoms at M6. Longitudinal follow-up of individuals with severe COVID-19 is warranted to better understand the pathophysiology underlying this long-term persistence.</td>
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**Methods:** Population-based cohort study using the Danish prescription, patient, and health insurance registries. All individuals with a positive or negative RT-PCR test for SARS-CoV-2 in Denmark between Feb 27 and May 31, 2020. The outcomes of interest were delayed acute complications, chronic disease, hospital visits due to persisting symptoms, and prescription drug use.  
**Findings:** > 10 498 eligible individuals tested positive for SARS-CoV-2 in Denmark from Feb 27 to May 31, 2020, of whom 8983 (85·6%) were alive and not admitted to hospital 2 weeks after their positive test. The matched SARS-CoV-2-negative reference population not admitted to hospital consisted of 80 894 individuals.  
> Compared with SARS-CoV-2-negative individuals, SARS-CoV-2-positive individuals were not at an increased risk of initiating new drugs (RD <0·1%) except bronchodilating agents, specifically short-acting β2-agonists (117 [1·7%] of 6935 positive individuals vs 743 [1·3%] of 57 206 negative individuals. And triptans (33 [0·4%] of 8292 vs 198 [0·3%] of 72 828.  
> There was an increased risk of receiving hospital diagnoses of dyspnoea (103 [1·2%] of 8676 vs 499 [0·7%] of 76 728; RD +0·6%; RR 2·00) and venous thromboembolism (20 [0·2%] of 8785 vs 110 [0·1%] of 78 872; RD +0·1%; RR 1·77) for SARS-CoV-2-positive individuals compared with negative individuals.  
> Prior event rate ratio-adjusted rate ratios of overall general practitioner visits (1·18) and outpatient hospital visits (1·10), but not hospital admission, showed increases among SARS-CoV-2-positive individuals compared with SARS-CoV-2-negative individuals.  
The absolute risk of severe post-acute complications after SARS-CoV-2 infection not requiring hospital admission is low. |
| *Clin Microbiol Infect.* 09MAY2021 | Outbreak investigation of symptomatic SARS-CoV-2 VOC 202012/01-lineage B.1.1.7 infection in healthcare workers, Italy | Loconsole D., et al. Italy [gotopaper](#) | Public Health / Epidemiology - Variants | **Aim:** to describe an outbreak of SARS-CoV-2 lineage B.1.1.7 infection in three HCWs in a hospital setting; two of the HCWs were fully vaccinated (i.e., had received two doses).  
**Methods:** Two physicians and one nurse working on the same shift on February 20, 2021, were involved in the outbreak. Real-time PCR, antigen tests, and serological tests for the IgG anti-spike protein of SARS-CoV-2 were performed, along with whole-genome sequencing (WGS).  
**Findings:** >SARS-CoV-2 infection was confirmed in all three HCWs; all presented with mild symptoms of COVID-19.  
The two physicians were fully vaccinated with BNT162b2 vaccine, with the second dose administered 1 month before symptom onset. Both had high titres of IgG anti-spike antibodies at the time of diagnosis.  
>WGS confirmed that all virus strains were VOC 202012/01-lineage B.1.1.7, suggesting a common source of exposure.  
>Epidemiological investigation revealed that the suspected source was a SARS-CoV-2-positive patient who required endotracheal intubation due to severe COVID-19. All procedures were carried out using a full suite of personal protective equipment (PPE).  
This mini-outbreak highlights some important issues about the efficacy of vaccines against transmission of SARS-CoV-2 variants, the high risk of exposure among HCWs, and the need for optimized implementation of PPE in hospitals. |
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Results  
> Antibody response was not detected in 26 of 392 (6.6%) COVID-19 convalescent subjects.  
> Over 9 months, the level of antibodies decreased by 50% but stabilized at 6 months and prevailed a protective level up to 9 months.  
> No differences were found regarding IgG SARS-CoV-2 antibody levels for age, gender, and major blood types, over-time.  
> COVID-19 asymptomatic subjects did not differ in antibody level overtime from subjects with mild to severe disease.  
> Repeated paired IgG SARS-COV-2 antibody level analyses disclosed that over 6 and 9 months, 15.3% (9 of 59) and 15.8% (3 of 19) of subjects became SARS-COV-2 IgG seronegative, respectively, all with low antibody level at 3 months. Rate of antibody decline was not affected by age, gender, or clinical symptomatology.  
> In a subgroup of recovering subjects, memory B-cell response up to 9-months post infection was undetectable in 31.8% (14/44) of subjects with no correlation to age, SARS-COV-2 antibody level, or time post-infection.  
Conclusions  
Majority of COVID-19 convalescent subjects develop IgG SARS-CoV-2 antibody response that prevails a protective level over a period of up to 9-months. |
| Lancet Rheumatol. 07MAY2021 | Non-steroidal anti-inflammatory drug use and outcomes of COVID-19 in the ISARIC Clinical Characterisation Protocol UK cohort: a matched, prospective cohort study | Drake T.M., et al. UK [gotopaper](https://doi.org/10.1016/S2468-2657(20)30086-6) | Therapeutics | We aimed to characterise the safety of NSAIDs and identify whether pre-existing NSAID use was associated with increased severity of COVID-19 disease.  
Methods  
> Prospective, multicentre cohort study included patients of any age admitted to hospital with a confirmed or highly suspected SARS-CoV-2 infection leading to COVID-19  
> We used propensity score matching to further estimate effects of NSAIDS while accounting for covariate differences in populations.  
Findings  
> Jan 17 and Aug 10, 2020, we enrolled 78 674 patients across 255 health-care facilities in England, Scotland, and Wales. 72 179 patients had death outcomes available for matching. 40 406 (56.2%) of 71 915 were men, 31 509 (43.8%) were women.  
> In this cohort, 4211 (5.8%) patients were recorded as taking systemic NSAIDs before admission to hospital.  
> At hospital admission, we observed no significant differences in severity between exposure groups.  
> After adjusting for explanatory variables, NSAID use was not associated with worse in-hospital mortality (matched OR 0.95, 95% CI 0.84–1.07; p=0.35), critical care admission (1.01, 0.87–1.17; p=0.89), requirement for invasive ventilation (0.96, 0.80–1.17; p=0.069), requirement for non-invasive ventilation (1.12, 0.96–1.32; p=0.14), requirement for oxygen (1.00, 0.89–1.12; p=0.97), or occurrence of acute kidney injury (1.08, 0.92–1.26; p=0.33).  
NSAID use is not associated with higher mortality or increased severity of COVID-19. Policy makers should consider reviewing issued advice around NSAID prescribing and COVID-19 severity. |
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<td>Sci Rep. 07MAY2021</td>
<td>Incorporating false negative tests in epidemiological models for SARS-CoV-2 transmission and reconciling with seroprevalence estimates</td>
<td>Bhattacharyya R., et al. India gotopaper</td>
<td>Public Health / Epidemiology</td>
<td><strong>Aims:</strong> Estimate the number of unreported COVID-19 cases and deaths by taking into account the false negative rate of RT-PCR tests in the Delhi area.</td>
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<td><strong>Methods:</strong></td>
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<td>&gt; Modelisation of the epidemic dynamic by an age-structured SEIR model taking into account the tests and false negatives.</td>
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<td>&gt; Estimation of the model parameters by a well-known stochastic algorithm.</td>
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<td>&gt; Model-based results are compared with data form serological surveys.</td>
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<td><strong>Findings:</strong> The number of Covid-19 cases and deaths in the Delhi area is dramatically underestimated.</td>
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<td>&gt; In July 2020, the underreporting factor was (34-53) for the number of cases, and (8-13) for the number of deaths.</td>
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<td>&gt; In January 2021, the underreporting factor remains (13-22) for the number of cases, and (3-7) for the number of deaths.</td>
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<td><strong>Limits:</strong></td>
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<td>&gt; The model does not take into account false positive of PCR tests.</td>
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<td>&gt; Estimation of the false negative rate of PCR tests has a large impact on the estimation of the underreporting factors, limiting their accuracy.</td>
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<td><strong>Conclusion:</strong></td>
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<td>&gt; Epidemic modeling can provide a less expensive method, with similar accuracy to large serological surveys, for estimating underreporting of Covid-19 cases.</td>
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<td>&gt; This model predicts a very large number of unreported deaths due to Covid-19, making the cost of achieving herd immunity by letting the epidemics run its course even higher than previously expected.</td>
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<td>Clin Infect Dis. 06MAY2021</td>
<td>Interacting Epidemics in Amazonian Brazil: Prior Dengue Infection Associated with Increased COVID-19 Risk in a Population-Based Cohort Study</td>
<td>Nicolete V.C., et al. International gotopaper</td>
<td>Public Health / Epidemiology</td>
<td><strong>Immunology after dengue virus (DENV) infection has been suggested to cross-protect from severe SARS-CoV-2 infection and mortality.</strong></td>
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<td><strong>Methods</strong></td>
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<td>&gt; Serological surveys in proven prior DENV infection diagnosed subjects before the coronavirus 2019 (COVID-19) pandemic</td>
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<td>&gt; Outcome: reduced the risk of SARS-CoV-2 infection and clinically apparent COVID-19 over the next 13 months</td>
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<td><strong>Findings</strong></td>
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<td>&gt; Anti-DENV IgG was found in 37.0% of 1,285 cohort participants in 2019</td>
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<td>&gt; In 2020, 35.2% of the participants had anti-SARS-CoV-2 IgG and 57.1% of the 448 SARS-CoV-2 seropositives reported clinical manifestations at the time of infection.</td>
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<td>&gt; Participants aged &gt;60 were twice more likely to have symptomatic COVID-19 than under-five children.</td>
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<td>&gt; Locally circulating SARS-CoV-2 isolates were assigned to the B.1.1.33 lineage.</td>
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<td><strong>Conclusion</strong></td>
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<td>Contrary to the cross-protection hypothesis, prior DENV infection was associated with twice the risk of clinically apparent COVID-19 upon SARS-CoV-2 infection</td>
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<td>JAMA 06MAY2021</td>
<td>Association Between Vaccination With BNT162b2 and Incidence of Symptomatic and Asymptomatic SARS-CoV-2 Infections Among Health Care Workers</td>
<td>Angel Y., et al. Israel gotopaper</td>
<td>Vaccines</td>
<td>Importance Randomized clinical trials have provided estimates of the effectiveness of the BNT162b2 vaccine against symptomatic SARS-CoV-2 infection, but its effect on asymptomatic infections remains unclear. Association of vaccination with the Pfizer-BioNTech BNT162b2 vaccine with symptomatic and asymptomatic SARS-CoV-2 infections among health care workers. Methods &gt; Single-center, retrospective cohort study (Tel Aviv, Israel). &gt; Data collected on symptomatic and asymptomatic SARS-CoV-2 infections confirmed via PCR tests in HCW undergoing regular screening with nasopharyngeal swabs &gt; Comparison of the incidence of infection between fully vaccinated and unvaccinated participants Primary outcome IRR for symptomatic and asymptomatic SARS-CoV-2 infection of fully vaccinated vs unvaccinated HCW Findings &gt; 6710 health care workers followed up for a median period of 63 days &gt; 5953 health care workers (88.7%) received at least 1 dose of the BNT162b2 vaccine, 5517 (82.2%) received 2 doses, and 757 (11.3%) were not vaccinated &gt; Symptomatic SARS-CoV-2 infection occurred in 8 fully vaccinated HCW and 38 unvaccinated HCW (incidence rate, 4.7 vs 149.8 per 100 000 person-days) &gt; Asymptomatic SARS-CoV-2 infection occurred in 19 fully vaccinated HCW and 17 unvaccinated HCW (incidence rate, 11.3 vs 67.0 per 100 000 person-days, respectively) Conclusions Receipt of the BNT162b2 vaccine compared with no vaccine was associated with a significantly lower incidence of symptomatic and asymptomatic SARS-CoV-2 infection more than 7 days after the second dose.</td>
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<td>Clin Infect Dis. 05MAY2021</td>
<td>SARS-CoV-2 RNAemia predicts clinical deterioration and extrapulmonary complications from COVID-19</td>
<td>Ram-Mohan N., et al. USA gotopaper</td>
<td>Clinic</td>
<td>Aim: to characterise relationships between SARS-CoV-2 RNAemia and disease severity, clinical deterioration, and specific extrapulmonary complications (EPCs). RNAemia was quantified by quantitative (qPCR) and digital (dPCR) PCR. &gt; 23.0% (44/191) of SARS-CoV-2 positive patients had viral RNA detected in plasma by dPCR, compared to 1.4% (2/147) by qPCR. &gt; Most patients with serial measurements had undetectable RNAemia within 10 days of symptom onset, reached maximum clinical severity within 16 days, and symptom resolution within 33 days. &gt; Initially RNAemic patients were more likely to manifest severe disease (OR 6.72 [95% CI, 2.45 – 19.79]), worsening of disease severity (OR 2.43 [95% CI, 1.07 – 5.38]), and EPCs (OR 2.81 [95% CI, 1.26 – 6.36]). RNA load correlated with maximum severity ($r = 0.47$ [95% CI, 0.20 – 0.67]). dPCR is more sensitive than qPCR for the detection of SARS-CoV-2 RNAemia, which is a robust predictor of eventual COVID-19 severity and oxygen requirements, as well as EPCs.</td>
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| JAMA 05MAY2021   | Antibody Response to 2-Dose SARS-CoV-2 mRNA Vaccine Series in Solid Organ Transplant Recipients | Boyarsky B.J., et al. USA | Vaccines | In this study, we assessed antibody response after the second dose. **Methods**  
> Transplant recipients without prior polymerase chain reaction–confirmed COVID-19 were recruited from across the US to participate in this prospective cohort  
> Those who completed the 2-dose SARS-CoV-2 mRNA vaccine series between December 16, 2020, and March 13, 2021, were included and followed up through April 13, 2021  
**Findings**  
> Overall, of the 658 participants, 98 (15%) had measurable antibody response after dose 1 and dose 2; 301 (46%) had no antibody response after dose 1 or dose 2; and 259 (39%) had no antibody response after dose 1 but subsequent antibody response after dose 2  
> Among all 658 participants, median (IQR) antibody levels after dose 2 were 2.14 U/mL (<0.4–245.8) (Roche) and 1.23 arbitrary units (0.13–6.38) (EUROIMMUN)  
> Among the 357 with detectable antibody after dose 2, median (IQR) antibody levels were 142.1 U/mL (9.44–250) (Roche) and 6.48 arbitrary units (3.75–8.72) (EUROIMMUN) overall;  
> Among the 473 receiving antimetabolites, 38 participants (8%) had antibody response after dose 1 and dose 2; 268 (57%) had no antibody response after dose 1 but subsequent antibody after dose 2. |

| Lancet Resp Med. 05MAY2021 | 3-month, 6-month, 9-month, and 12-month respiratory outcomes in patients following COVID-19-related hospitalisation: a prospective study | Wu X., et al. China/UK | Clinic | Aim: to describe the temporal trends in respiratory outcomes over 12 months in patients hospitalised for severe COVID-19 and to investigate the associated risk factors. **Methods:**  
Prospective, longitudinal, cohort study, patients admitted to hospital for severe COVID-19 who did not require mechanical ventilation were prospectively followed up at 3 months, 6 months, 9 months, and 12 months after discharge from Renmin Hospital of Wuhan University, Wuhan, China.  
Patients with a history of hypertension; diabetes; cardiovascular disease; cancer; and chronic lung disease, including asthma or chronic obstructive pulmonary disease; or a history of smoking were excluded.  
>135 eligible patients, 83 (61%) patients participated in this study. **Findings:**  
> The median age of participants was 60 years (IQR 52–66). Temporal improvement in pulmonary physiology and exercise capacity was observed in most patients; however, persistent physiological and radiographic abnormalities remained in some patients with COVID-19 at 12 months after discharge.  
> A significant reduction in DLCO over the study period was observed, with a median of 77% of predicted (IQR 67–87) at 3 months, 76% of predicted (68–90) at 6 months, and 88% of predicted (78–101) at 12 months after discharge.  
> At 12 months after discharge, radiological changes persisted in 20 (24%) patients.  
> Multivariate logistic regression showed increasing odds of impaired DLCO associated with female sex (odds ratio 8.61) and radiological abnormalities were associated with peak HRCT pneumonia scores during hospitalisation (1.36).  
In most patients who recovered from severe COVID-19, dyspnoea scores and exercise capacity improved over time. |
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| Nature Med. 05MAY2021 | Delayed production of neutralizing antibodies correlates with fatal COVID-19 | Lucas C., et al. USA [gotopaper](#) | Immunology | **Aim:** to study the exact features of antibody responses that govern COVID-19 disease outcomes, analysing the nature of antibody responses in disease severity and mortality in 229 Covid-19 patients.  
> A correlation between anti-spike (S) IgG levels, length of hospitalization and clinical parameters associated with worse clinical progression was observed. Although high anti-S IgG levels correlated with worse disease severity, such correlation was time dependent.  
> Decreased patients did not have higher overall humoral response than discharged patients. However, they mounted a robust, yet delayed, response, measured by anti-S, anti-receptor-binding domain IgG and neutralizing antibody (NAb) levels compared to survivors.  
> Delayed seroconversion kinetics correlated with impaired viral control in deceased patients.  
> Sera from 85% of patients displayed some neutralization capacity during their disease course, but NAb generation before 14 d of disease onset emerged as a key factor for recovery.  

These data indicate that COVID-19 mortality does not correlate with the cross-sectional antiviral antibody levels per se but rather with the delayed kinetics of NAb production. |
| Lancet 05MAY2021 | Impact and effectiveness of mRNA BNT162b2 vaccine against SARS-CoV-2 infections and COVID-19 cases, hospitalisations, and deaths following a nationwide vaccination campaign in Israel: an observational study using national surveillance data | Haas E.J., et al. Israel [gotopaper](#) | Vaccines | **Real-world effectiveness of two doses of BNT162b2 against a range of SARS-CoV-2 outcomes and to evaluate the nationwide public-health impact following the widespread introduction of the vaccine.**  
**Methods**  
> National surveillance data from the first 4 months of the nationwide vaccination campaign to ascertain:  
(i) incident cases of laboratory-confirmed SARS-CoV-2 infections  
(ii) vaccine uptake in residents of Israel aged 16 years and older.  
> Vaccine effectiveness against SARS-CoV-2 outcomes was calculated on the basis of incidence rates in fully vaccinated individuals compared with rates in unvaccinated individuals  
**Findings**  
> By April 3, 2021, 4,714,932 (72.1%) of 6,538,911 people aged 16 years and older were fully vaccinated with two doses of BNT162b2.  
> Vaccine effectiveness at 7 days or longer after the second dose were  
95.3% against SARS-CoV-2 infection,  
91.5% against asymptomatic SARS-CoV-2 infection,  
97.0% against symptomatic COVID-19,  
97.2% against COVID-19-related hospitalisation  
97.5% against severe or critical COVID-19-related hospitalisation  
96.7% against COVID-19-related death.  
> In all age groups, as vaccine coverage increased, the incidence of SARS-CoV-2 outcomes declined.  
> 8006 of 8472 samples tested showed a spike gene target failure, giving an estimated prevalence of the B.1.1.7 variant of 94.5% among SARS-CoV-2 infections.  
**Conclusions**  
Two doses of BNT162b2 are highly effective across all age groups in preventing symptomatic and asymptomatic SARS-CoV-2 infections and COVID-19-related hospitalisations, severe disease, and death, including those caused by the B.1.1.7 SARS-CoV-2 variant. There were marked and sustained declines in SARS-CoV-2 incidence corresponding to increasing vaccine coverage. |
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Methods  
> Phase 2a–b trial (South Africa).  
> Population: HIV− or HIV + in a stable condition. Randomization 1:1 ratio, two doses of either NVX-CoV2373 vaccine (5 μg of recombinant spike protein with 50 μg of Matrix-M1 adjuvant) or placebo.  
Primary end points: Safety and vaccine efficacy against laboratory-confirmed symptomatic Covid-19 at ≥7 days after the second dose  
Findings  
> 4387 participants receiving at least 1 injection of vaccine or placebo.  
> 30% of the participants were seropositive for SARS-CoV-2 at baseline.  
> Among 2684 baseline seronegative participants (94% HIV-negative and 6% HIV-positive), predominantly mild-to-moderate Covid-19 developed in 15 participants in the vaccine group and 29 in the placebo group (vaccine efficacy, 49.4%; 95% CI, 6.1 to 72.8).  
> Vaccine efficacy among HIV-negative participants was 60.1% (95% CI, 19.9 to 80.1).  
> Of 41 sequenced isolates, 38 (92.7%) were the B.1.351 variant.  
> Post hoc vaccine efficacy against B.1.351 was 51.0% (95% CI, −0.6 to 76.2) among the HIV-negative participants.  
> Preliminary local and systemic reactogenicity events were more common in the vaccine group; serious adverse events were rare in both groups.  
Conclusions  
The NVX-CoV2373 vaccine was efficacious in preventing Covid-19, with higher vaccine efficacy observed among HIV-negative participants. Most infections were caused by the B.1.351 variant. |
| NEJM 05MAY2021   | Effectiveness of the BNT162b2 Covid-19 Vaccine against the B.1.1.7 and B.1.351 Variants | Abu-Raddad L.J., Chemaitelly H. Qatar [gotopaper](https://doi.org/10.1056/NEJMoa2109514) | Vaccines | Context  
Qatar launched a mass immunization campaign with BNT16b2 vaccine on December 21, 2020. Vaccination scale-up occurred during Qatar’s 2nd and 3rd waves of SARS-CoV-2 infections, triggered by expansion of B.1.1.7 and B.1.351 variants. Viral genome sequencing (Feb 23 to March 18) indicated that 50.0% of cases of Covid-19 in Qatar were caused by B.1.351 and 44.5% by B.1.1.7. Nearly all cases in which virus was sequenced after March 7 were caused by either B.1.351 or B.1.1.7.  
Aim: Evaluation of BNT162b2 vaccine effectiveness  
Methodology  
(i) Test-negative case–control study design  
(ii) Cohort study design by comparing the incidence of infection among vaccinated persons with the incidence in the national cohort of persons who were antibody-negative  
Findings  
> Effectiveness of the vaccine against any documented infection with B.1.1.7 variant: 89.5% (95% [CI], 85.9 to 92.3) at ≥14 days after dose 2  
> Effectiveness against any documented infection with the B.1.351 variant: 75.0% (95% CI, 70.5 to 78.9).  
> Vaccine effectiveness against severe, critical, or fatal disease due to infection with any SARS-CoV-2 (with B.1.1.7 and B.1.351 variants being predominant within Qatar): 97.4% (95% CI, 92.2 to 99.5)  
> Vaccine effectiveness within the cohort 87.0% (95% CI, 81.8 to 90.7) against the B.1.1.7 variant and 72.1% (95% CI, 66.4 to 76.8) against the B.1.351 variant.  
Conclusions  
The BNT162b2 vaccine was effective against infection and disease in the population of Qatar, despite the B.1.1.7 and B.1.351 variants being predominant within the country  
> Vaccine effectiveness against the B.1.351 variant was approximately 20% lower than the effectiveness (>90%) reported in the clinical trial and in real-world conditions in Israel and the United States.  
> The reduced protection against infection with the B.1.351 variant did not seem to translate into poor protection against the most severe forms of infection (greater than 90%) |
### Methods
> Retrospective analysis of clinical parameters from 85 patients with laboratory-confirmed coronavirus disease 2019 (COVID-19); moreover, kidney histopathology from six additional COVID-19 patients with post-mortem examinations was performed.

### Findings
> We find that 27% (23/85) of patients exhibited AKI. The elderly patients and cases with comorbidities (hypertension and heart failure) are more prone to develop AKI.
> Haematoxylin & eosin staining shows that the kidneys from COVID-19 autopsies have moderate to severe tubular damage.
> In situ hybridization assays illustrate that viral RNA accumulates in tubules.
> Immunohistochemistry shows nucleocapsid and spike protein deposits in the tubules, and immunofluorescence double staining shows that both antigens are restricted to the angiotensin converting enzyme-II-positive tubules.
> Immunohistochemistry shows nucleocapsid and spike protein deposits in the tubules, and immunofluorescence double staining shows that both antigens are restricted to the angiotensin converting enzyme-II-positive tubules.
> SARS-CoV-2 infection triggers the expression of hypoxic damage-associated molecules, including DP2 and prostaglandin D synthase in infected tubules. Moreover, it enhances CD68+ macrophages infiltration into the tubulointerstitium, and complement C5b-9 deposition on tubules is also observed.

These results suggest that SARS-CoV-2 directly infects human kidney to mediate tubular pathogenesis and AKI.

### Aim:
To assess rates of antibody response to mRNA SARS-CoV-2 vaccine among kidney transplant recipients, and to identify factors associated with reduced immunogenicity.

### Methods
> Prospective cohort study including consecutive kidney transplant recipients in a single referral transplant center.
> Anti-spike (anti-S) antibodies test 2-4 weeks following second vaccine dose.

### Primary outcome:
Rate of seropositivity.

### Findings:
> 308 kidney transplant recipients included, only 112 (36.4%) tested positive for anti-S antibodies 2-4 weeks after receiving the second dose of BNT162b2 vaccine.
> Median antibody titers: 15.5 AU/mL
> Factors associated with antibody response:
  (i) higher estimated glomerular filtration rate (eGFR) (odds ratio [OR] 1.025 per ml/min/1.73m², 95% confidence interval [CI] 1.014 - 1.037, p<0.001),
  (ii) lower mycophenolic acid dose (OR 2.347 per 360 mg decrease, 95% CI 1.782 - 3.089, p<0.001),
  (iii) younger age (OR 1.032 per year decrease, 95% CI 1.015 - 1.05, p<0.001)
  (iv) lower calcineurin inhibitors (CNI) blood level (OR 1.987, 95% CI 1.146 - 3.443, p=0.014).
> No serious adverse events to the vaccine were reported.

### Conclusions:
Kidney transplant recipients demonstrated inadequate antibody response to mRNA SARS-CoV-2 vaccination. Immunosuppression level was a significant factor in this response.
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| **Lancet 01MAY2021** | **Tocilizumab in patients admitted to hospital with COVID-19 (RECOVERY): a randomised, controlled, open-label, platform trial** | **RECOVERY Collaborative Group**<br>UK | **Therapeutics** | **Aim:** to evaluate the effects of tocilizumab in adult patients admitted to hospital with COVID-19 with both hypoxia and systemic inflammation.  
**Methods:** Randomised, controlled, open-label, platform trial assessing several possible treatments in patients hospitalised with COVID-19 in the UK. Participants with hypoxia (oxygen saturation <92% on air or requiring oxygen therapy) and evidence of systemic inflammation (C-reactive protein 275 mg/L) were eligible for random assignment in a 1:1 ratio to usual standard of care alone versus usual standard of care plus tocilizumab at an IV dose of 400 mg–800 mg (depending on weight). The primary outcome was 28-day mortality, assessed in the intention-to-treat population.  
**Findings:**  
> 4116 adults of 21 550 patients enrolled into the RECOVERY trial were included in the assessment of tocilizumab, including 3385 (82%) patients receiving systemic corticosteroids.  
> Overall, 621 (31%) of the 2022 patients allocated tocilizumab and 729 (35%) of the 2094 patients allocated to usual care died within 28 days (rate ratio 0·85; 95% CI 0·76–0·94; p=0·0028).  
> Consistent results were seen in all prespecified subgroups of patients, including those receiving systemic corticosteroids. **Patients allocated to tocilizumab were more likely to be discharged from hospital within 28 days** (57% vs 50%; rate ratio 1·22; 1·12–1·33; p<0·0001).  
> Among those not receiving invasive mechanical ventilation at baseline, patients allocated tocilizumab were less likely to reach the composite endpoint of invasive mechanical ventilation or death (35% vs 42%; risk ratio 0·84; 95% CI 0·77–0·92; p<0·0001).  
**In hospitalised COVID-19 patients with hypoxia and systemic inflammation,** tocilizumab improved survival and other clinical outcomes. |
| **JAMA 30APR2021** | **US Case Reports of Cerebral Venous Sinus Thrombosis With Thrombocytopenia After Ad26.COV2.S Vaccination, March 2 to April 21, 2021** | **See L., et al. USA**<br>gotopaper | **Vaccines** | **Aim:** To describe reports of CVST with thrombocytopenia following Ad26.COV2.S vaccine receipt.  
**Methods**  
Case series of 12 US patients with CVST and thrombocytopenia following use of Ad26.COV2.S vaccine under EUA reported to the Vaccine Adverse Event Reporting System (VAERS) from March 2 to April 21, 2021 (with follow-up reported through April 21, 2021).  
**Main Outcomes and Measures:** Clinical course, imaging, laboratory tests, and outcomes after CVST diagnosis obtained from VAERS reports, medical record review, and discussion with clinicians.  
**Findings**  
> Patients characteristics:  
(i) age: 18 to younger than 60 years  
(ii) all White women.  
(iii) seven patients had at least 1 CVST risk factor, including obesity (n = 6), hypothyroidism (n = 1), and oral contraceptive use (n = 1); none had documented prior heparin exposure.  
> Time from Ad26.COV2.S vaccination to symptom onset: 6 to 15 days.  
> 11 patients initially presented with headache; 1 patient initially presented with back pain and later developed headache.  
> Of the 12 patients with CVST, 7 also had intracerebral hemorrhage; 8 had non-CVST thromboses.  
> After diagnosis of CVST, 6 patients initially received heparin treatment.  
> Platelet nadir ranged from 9 ×103/µL to 127 ×103/µL. All 11 patients tested for the heparin-platelet factor 4 HIT antibody by enzyme-linked immunosorbent assay (ELISA) screening had positive results.  
> All patients were hospitalized (10 in an intensive care unit [ICU]). As of April 21, 2021, outcomes were death (n = 3), continued ICU care (n = 3), continued non-ICU hospitalization (n = 2), and discharged home (n = 4).  
**Conclusions**  
The initial 12 US cases of CVST with thrombocytopenia after Ad26.COV2.S vaccination represent serious events. |
### Science 30APR2021

**Title**: Prior SARS-CoV-2 infection rescues B and T cell responses to variants after first vaccine dose

**Authors**: Reynolds C.J., et al.

**Field of expertise**: Vaccines

**Key facts**

**Aim**: to investigate if single dose vaccination, with or without prior infection, confers cross protective immunity to variants.

- Analysis of T and B cell responses after first dose vaccination with the Pfizer/BioNTech mRNA vaccine BNT162b2 in healthcare workers (HCW) followed longitudinally, with or without prior Wuhan-Hu-1 SARS-CoV-2 infection.

  > After one dose, individuals with prior infection showed enhanced T cell immunity, antibody secreting memory B cell response to spike and neutralizing antibodies effective against B.1.1.7 and B.1.351.

  > By comparison, HCW receiving one vaccine dose without prior infection showed reduced immunity against variants.

  > B.1.1.7 and B.1.351 spike mutations resulted in increased, abrogated or unchanged T cell responses depending on human leukocyte antigen (HLA) polymorphisms.

  Single dose vaccination with BNT162b2 in the context of prior infection with a heterologous variant substantially enhances neutralizing antibody responses against variants.

### Lancet 30APR2021

**Title**: Ethnic differences in SARS-CoV-2 infection and COVID-19-related hospitalisation, intensive care unit admission, and death in 17 million adults in England: an observational cohort study using the OpenSAFELY platform

**Authors**: Mathur R., et al.

**Field of expertise**: Public Health / Epidemiology

**Key facts**

**Aim**: to quantify ethnic differences in SARS-CoV-2 infection and COVID-19 outcomes during the first and second waves of the COVID-19 pandemic in England.

**Methods**

- Observational cohort study of adults (aged ≥18 years) who had at least 1 year of continuous registration with primary care practices in England at the start of each study period (3 waves).

- Multivariable Cox regression was used to examine ethnic differences in the outcomes of interest. Models were adjusted for age, sex, deprivation, clinical factors and comorbidities, and household size, with stratification by geographical region.

**Results**

> 17,288,532 adults included: 10,877,978 (62.9%) White, 1,025,319 (5.9%) South Asian, 340,912 (2.0%) Black, 170,484 (1.0%) of mixed ethnicity, 320,788 (1.9%) of other ethnicity, and 4,553,051 (26.3%) of unknown ethnicity.

> In wave 1, the likelihood of being tested for SARS-CoV-2 infection was slightly higher in the South Asian group (adjusted hazard ratio 1.08 [95% CI 1.07–1.09]), Black group (1.08 [1.06–1.09]), and mixed ethnicity group (1.04 [1.02–1.05]) and was decreased in the other ethnicity group (0.77 [0.76–0.78]) relative to the White group.

> The risk of testing positive for SARS-CoV-2 infection was higher in the South Asian group (1.99 [1.94–2.04]), Black group (1.69 [1.62–1.77]), mixed ethnicity group (1.49 [1.39–1.59]), and other ethnicity group (1.20 [1.14–1.28]).

> Compared with the White group, the four remaining high-level ethnic groups had an increased risk of COVID-19-related hospitalisation (South Asian group 1.99 [1.94–2.04], Black group 1.69 [1.62–1.77], mixed ethnicity group 1.49 [1.39–1.59], and other ethnicity group 1.20 [1.14–1.28]).

> In wave 2, the risks of hospitalisation, ICU admission, and death relative to the White group were increased in the South Asian group but attenuated for the Black group compared with these risks in wave 1.

> Disaggregation into 16 ethnicity groups showed important heterogeneity within the five broader categories.

Some minority ethnic populations in England have excess risks of testing positive for SARS-CoV-2 and of adverse COVID-19 outcomes compared with the White population, even after accounting for differences in sociodemographic, clinical, and household characteristics.
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<td><strong>Methods:</strong> &gt; Massive online survey administered through Facebook throughout the United States (576,051 respondents with at least one child in school age); analysis adjusts for non-response and coverage bias. &gt; COVID-19 infection assessed through: i) reporting COVID-19-like illness (CLI; fever, cough/shortness of breath), ii) loss of taste or smell, and/or iii) positive SARS-CoV-2 test result within the previous 14 days. &gt; Questions on type of schooling (in-person or not, full-time or part-time), and on mitigation measures in place at school.</td>
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<td><strong>Findings:</strong> &gt; Living in a household with a child engaged in full-time in-person schooling is associated with a substantial increase in the odds reporting COVID-19-like illness (CLI), loss of taste or smell, or a positive SARS-CoV-2 test result within the previous 14 days. &gt; When the child is engaged in part-time schooling, the association is attenuated but still statistically significant. &gt; There is a negative relationship between the number of mitigation measures implemented and the risk of COVID-19 outcomes among adult household members responding to the survey. &gt; Daily symptom screening is associated with the greater risk reduction; mask mandates and cancelling extra-curricular activities are also associated with risk reduction. &gt; Limits: self-reporting; confounding factors (heterogeneities of economic and racial status)</td>
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<td><strong>Conclusion:</strong> The results of this massive online survey in the US provide evidence that in-person schooling poses a risk to those living in the households of children, but that this risk can be managed through commonly implemented school-based mitigation measures.</td>
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<td>Clin Infect Dis. 29APR2021</td>
<td>Development and validation of the long covid symptom and impact tools, a set of patient-reported instruments constructed from patients' lived experience</td>
<td>Tran V., et al. France gotopaper</td>
<td>Diagnostics</td>
<td><strong>Aim:</strong> To develop and validate patient-reported instruments, based on patients’ lived experiences, for monitoring the symptoms and impact of long covid.</td>
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<td><strong>Design</strong> &gt; The long covid Symptom and Impact Tools (ST and IT) were constructed from the answers to a survey with open-ended questions to 492 patients with long COVID. &gt; Tool validation: adult patients with suspected or confirmed COVID-19 and symptoms &gt;3 weeks after onset. &gt; Construct validity was assessed by examining the relations of the ST and IT scores with health related quality of life (EQ-5D-5L), function (PCFS, post-COVID functional scale), and perceived health (MYMOP2). Reliability was determined by a test-retest. “Patient acceptable symptomatic state” (PASS) was determined by the percentile method.</td>
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<td><strong>Results</strong> &gt; Validation involved 1022 participants (55% confirmed cases, 79% female, and 12.5% hospitalized for COVID-19). &gt; The long COVID ST and IT scores were strongly correlated with the EQ-5D-5L (rs = -0.45 and rs = -0.59 respectively), the PCFS (rs = -0.39 and rs = -0.55), and the MYMOP2 (rs = -0.40 and rs = -0.59). &gt; Reproducibility was excellent with an interclass correlation coefficient of 0.83 (95% CI 0.80-0.86) for the ST score, 0.84 (0.80-0.87) for the IT score. &gt; 793 (77.5%) patients reported an unacceptable symptomatic state, thereby setting the PASS for the long covid IT score at 30 (28 to 33).</td>
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<td><strong>The long covid ST and IT tools provide the first validated and reliable instruments for monitoring the symptoms and impact of long covid.</strong></td>
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<td><strong>Lancet HIV</strong> 29APR21</td>
<td>SARS-CoV-2 seroprevalence, and IgG concentration and pseudovirus neutralising antibody titres after infection, compared by HIV status: a matched case-control observational study</td>
<td>Spinelli A.M., et al. USA gotopaper</td>
<td>Public Health / Epidemiology</td>
<td>We aimed to compare SARS-CoV-2 IgG seroprevalence, disease severity, and neutralising antibody activity after infection among people with and without HIV receiving care in a county hospital system over a 3-month period. <strong>Methods</strong> &gt; In this matched case-control observational study, remnant serum samples were collected between Aug 1 and Oct 31, 2020, from all people living with HIV who underwent routine outpatient laboratory testing in a municipal health-care system <strong>Findings</strong> &gt; 1138 samples from 955 people living with HIV and 1118 samples from 1062 people without HIV were tested. &gt; SARS-CoV-2 IgG seroprevalence was 3·7% (95% CI 2·4 to 5·0) among people with HIV compared with 7·4% (5·7 to 9·2) among people without HIV (adjusted odds ratio 0·50, 95% CI 0·30 to 0·83). &gt; Among 31 people with HIV and 70 people without HIV who had evidence of past infection, the odds of severe COVID-19 were 5·52 (95% CI 1·01 to 64·48) times higher among people living with HIV. &gt; Adjusting for time since PCR-confirmed infection, SARS-CoV-2 IgG concentrations were lower (percentage change −53%, 95% CI −4 to −76), pseudovirus neutralising antibody titres were lower (−67%, −25 to −86), and avidity was similar (7%, −73 to 87) among people living with HIV compared with those without HIV. Although fewer infections were detected by SARS-CoV-2 IgG testing among people living with HIV than among those without HIV, people with HIV had more cases of severe COVID-19. Among people living with HIV with past SARS-CoV-2 infection, lower IgG concentrations and pseudovirus neutralising antibody titres might reflect a diminished serological response to infection, and the similar avidity could be driven by similar time since infection.</td>
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<td><strong>Clin Infect Dis.</strong> 29APR21</td>
<td>SARS-CoV-2 detection on self-collected saliva or anterior nasal specimens compared with healthcare personnel-collected nasopharyngeal specimens</td>
<td>Marx G.E., et al. USA gotopaper</td>
<td>Public Health / Epidemiology</td>
<td>Self-collected saliva or anterior nasal specimens (ANS) for SARS-CoV-2 detection are less invasive but the sensitivity of these specimen types has not been thoroughly evaluated. <strong>Methods</strong> &gt; During September–November 2020, 730 adults undergoing SARS-CoV-2 testing at community testing events and homeless shelters in Denver provided self-collected saliva and ANS specimens before NPS collection and answered a short survey about symptoms and specimen preference. &gt; Subgroup analyses included test outcomes by symptom status and culture results <strong>Findings</strong> &gt; Sensitivity for SARS-CoV-2 detection by rRT-PCR appeared higher for saliva than for ANS (85% vs. 80%) and among symptomatic participants than among those without symptoms (94% vs. 29% for saliva; 87% vs. 50% for ANS). &gt; Among participants with culture-positive SARS-CoV-2 by any specimen type, sensitivity of saliva and ANS by rRT-PCR was 94% and 100%, respectively. &gt; Saliva and ANS were equally preferred by participants; most would undergo NPS again despite being least preferred. Saliva was slightly more sensitive than ANS for SARS-CoV-2 detection by rRT-PCR. Both saliva and ANS reliably detected SARS-CoV-2 among participants with symptoms. Self-collected saliva and ANS offer practical advantages, are preferred by patients, and might be most useful for testing people with COVID-19 symptoms.</td>
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| Lancet Diabetes Endocrinol. 28APR2021 | Associations between body-mass index and COVID-19 severity in 6.9 million people in England: a prospective, community-based, cohort study | Gao M., et al. UK gotopaper | Public Health / Epidemiology | **Aim:** to examine the association between obesity the risk of severe COVID-19, including interactions with demographic and behavioural characteristics, type 2 diabetes, and other health conditions.  
**Methods:** Prospective, community-based, cohort study, using de-identified patient-level data from the QResearch database of general practices in England, UK. Data was extracted from patients aged 20 years and older who were registered at a practice eligible for inclusion in the QResearch database, and with available data on BMI. Outcomes, as a proxy measure of severe COVID-19, were admission to hospital, admission to an intensive care unit (ICU), and death due to COVID-19.  
**Findings:** > Among 6 910 695 eligible individuals (mean BMI 26·78 kg/m² [SD 5·59]), 13 503 (0·20%) were admitted to hospital, 1601 (0·02%) to an ICU, and 5479 (0·08%) died after a positive test for SARS-CoV-2. > J-shaped associations were found between BMI and admission to hospital due to COVID-19 (adjusted hazard ratio [HR] per kg/m² from the nadir at BMI of 23 kg/m² of 1·05 [95% CI 1·05–1·05]) and death (1·04 [1·04–1·05]), and a linear association across the whole BMI range with ICU admission (1·10 [1·09–1·10]). > A significant interaction was found between BMI and age and ethnicity, with higher HR per kg/m² above BMI 23 kg/m² for younger people (20–39 years age group vs 80–100 years group) and Black people than White people (1·07 vs 1·04). > The risk of admission to hospital and ICU due to COVID-19 associated with unit increase in BMI was slightly lower in people with type 2 diabetes, hypertension, and cardiovascular disease than in those without these morbidities.  
**The relative risk due to increasing BMI is particularly notable in people younger than 40 years and of Black ethnicity.** |
## Vaccine side-effects and SARS-CoV-2 infection after vaccination in users of the COVID Symptom Study app in the UK: a prospective observational study

### Authors and expertise
- **Authors:** Menni C., et al. UK
- **Field of expertise:** Vaccines

### Key facts
**Aim:** to investigate the safety and effectiveness of BNT162b2 and ChAdOx1 nCoV-19 vaccines in a UK community setting.

**Methods:**
Prospective observational study examining proportion and probability of self-reported systemic and local side-effects 88 days of vaccination in individuals using the COVID Symptom Study app who received 1 or 2 doses of BNT162b2 or 1 dose of ChAdOx1 nCoV-19.

Infection rates were compared in vaccinated individuals tested for SARS-CoV-2 (PCR, lateral flow tests) and in unvaccinated controls.

**Findings:**
> Between Dec 8, and March 10, 2021, 627,383 individuals reported being vaccinated with 655,590 doses: 282,103 received one dose of BNT162b2, of whom 28,207 received a second dose, and 345,280 received one dose of ChAdOx1 nCoV-19.  
> Systemic side-effects were reported by 13.5% of individuals after the first dose of BNT162b2, by 22.0% after the second dose of BNT162b2, and by 33.7% after the first dose of ChAdOx1 nCoV-19.  
> Local side-effects were reported by 71.9% of individuals after the first dose of BNT162b2, by 68.5% after the second dose of BNT162b2, and by 58.7% after the first dose of ChAdOx1 nCoV-19.  
> Systemic side-effects were more common (1.6 times after the first dose of ChAdOx1 nCoV-19 and 2.9 times after the first dose of BNT162b2) among individuals with previous SARS-CoV-2 infection than among those without known past infection.  
> 3106 of 103,622 vaccinated individuals and 50,340 of 464,356 unvaccinated controls tested positive for SARS-CoV-2 infection.  
> Significant reductions in infection risk were seen starting at 12 days after the first dose, reaching 60% (95% CI 49–68) for ChAdOx1 nCoV-19 and 69% (66–72) for BNT162b2 at 21–44 days and 72% (63–79) for BNT162b2 after 45–59 days.

## Safety and immunogenicity of one versus two doses of the COVID-19 vaccine BNT162b2 for patients with cancer: interim analysis of a prospective observational study

### Authors and expertise
- **Authors:** Monin L., et al. UK
- **Field of expertise:** Vaccines

### Key facts
**Aim:** Assessment of the safety and immunogenicity of the BNT162b2 (Pfizer–BioNTech) vaccine in patients with cancer (interim data)

**Methods:**
> Prospective observational study.  
> Patients with cancer and healthy controls (mostly HCW)  
> Participants who were vaccinated between Dec 8 and Dec 29, 2020, received two 30 μg doses of BNT162b2 administered intramuscularly 21 days apart; patients vaccinated after this date received only one 30 μg dose with a planned follow-up boost at 12 weeks.  
> Blood samples taken before vaccination and at 3 and 5 weeks after the first vaccination.  
> Where possible (rRT-PCR) swab tests were done every 10 days or in cases of symptomatic COVID-19.

**Primary endpoints:** seroconversion to SARS-CoV-2 spike (S) protein in patients with cancer following the first vaccination with the BNT162b2 vaccine, effect of vaccine boosting after 21 days on seroconversion.

**Findings:**
> 151 patients with cancer (95 patients with solid cancer and 56 patients with haematological cancer) and 54 healthy controls.  
> The proportion of positive anti-S lgG titres at 21 days after a single vaccine dose across the three cohorts were 32 (94%); 95% CI 81–98) of 34 healthy controls; 21 (38%; 26–51) of 56 patients with solid cancer, and 8 (18%; 10–32) of 44 patients with haematological cancer.  
> 16 healthy controls, 25 patients with solid cancer, and six patients with haematological cancer received a second dose on day 21. Of the patients with available blood samples 2 weeks following a 21-day vaccine boost, 18 (95%; 95% CI 75–99) of 19 patients with solid cancer, 12 (100%; 76–100) of 12 healthy controls, and three (60%; 23–88) of five patients with haematological cancers were seropositive, compared with ten (30%; 17–47) of 33, 18 (86%; 65–95) of 21, and four (11%; 4–25) of 36, respectively, who did not receive a boost.  
> The vaccine was well tolerated. No vaccine-related deaths reported.

**Conclusion**
In patients with cancer, one dose of the BNT162b2 vaccine yields poor efficacy. Immunogenicity increased significantly in patients with solid cancer within 2 weeks of a vaccine boost at day 21 after the first dose.
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<tr>
<td>Clin Infect Dis. 27APR2021</td>
<td>Vaccinated and convalescent donor-derived SARS-CoV-2-specific T cells as adoptive immunotherapy for high-risk COVID-19 patients</td>
<td>Papayanni P.G., et al. Greece gotopaper</td>
<td>Therapeutics</td>
<td>Aim: to provide the rationale towards the development of a SARS-CoV-2-specific T-cell (CoV-2-ST) bank from convalescent donors as T-cell immunotherapy against severe COVID-19. Methods: SARS-CoV-2-specific T-cell immunity and expansion was tested in unexposed donors, COVID-19 infected individuals (convalescent), asymptomatic PCR-positive subjects, vaccinated individuals, non-ICU hospitalized patients and ICU patients who either recovered and were discharged (ICU recovered) or had a prolonged stay and/or died (ICU critical). CoV-2-STs were generated from all types of donors and underwent phenotypic and functional assessment. Findings: &gt; A causal relationship between the expansion of endogenous CoV-2-STs and the disease outcome was shown; insufficient expansion of circulating CoV-2-STs identified hospitalized patients at high-risk for an adverse outcome. &gt; CoV-2-STs with a similarly functional and non-alloreactive, albeit highly cytotoxic, profile against SARS-CoV-2 could be expanded from both convalescent and vaccinated donors generating clinical-scale, SARS-CoV-2-specific T-cell products with functional activity against both the unmutated virus and its B.1.1.7 variant. &gt; Critical COVID-19 patient-originating CoV-2-STs failed to expand, recapitulating the in vivo failure of CoV-2-specific T-cell immunity to control the infection. &gt; CoV-2-STs generated from asymptomatic PCR+ individuals presented weak responses whereas their counterparts originating from exposed to other seasonal coronaviruses subjects failed to kill the virus, thus disempowering the hypothesis of protective cross-immunity. The authors provide evidence on risk stratification of hospitalized COVID-19 patients and the feasibility of generating powerful CoV-2-ST products from both convalescent and vaccinated donors as an “off-the-shelf” T-cell immunotherapy for high-risk patients.</td>
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<td>Clin Infect Dis. 27APR2021</td>
<td>Age-dependent immune response to the Biontech/Pfizer BNT162b2 COVID-19 vaccination</td>
<td>Müller L., et al. Germany gotopaper</td>
<td>Vaccines</td>
<td>Cohort study with two age groups (young vaccinees &lt;60 years old and elderly vaccinees &gt;80, to compare antibody responses to the first and second dose of the BNT162b2 COVID-19 vaccination. &gt; While the majority of participants in both groups produced specific IgG antibody titers against SARS-CoV-2 spike protein, titers were significantly lower in elderly participants. &gt; The increment of antibody levels after the second immunization was higher in elderly participants, but the absolute mean titer of this group remained lower than the &lt;60 group. &gt; After the second vaccination, 31.3% of the elderly had no detectable neutralizing antibodies in contrast to the younger group, in which only 2.2% had no detectable neutralizing antibodies. These data show differences between the antibody responses raised after the first and second BNT162b2 vaccination, in particular lower frequencies of neutralizing antibodies in the elderly group.</td>
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<td>Lancet 23APR2021</td>
<td>Interim findings from first-dose mass COVID-19 vaccination roll-out and COVID-19 hospital admissions in Scotland: a national prospective cohort study</td>
<td>Vasileiou E., et al. UK gotopaper</td>
<td>Vaccines</td>
<td>Aim: prospective cohort study to investigate the association between the mass roll-out of the first doses of BNT162b2 mRNA and ChAdOx1 nCoV-19 COVID-19 vaccines and hospital admissions for COVID-19. &gt; 1 331,993 people were vaccinated between Dec 8, 2020, and Feb 22, 2021 (mean age 65 years (SD 16·2)). &gt; The first dose of the BNT162b2 mRNA vaccine was associated with a vaccine effect of 91% (95% CI 85–94) for reduced COVID-19 hospital admission at 28–34 days post-vaccination. &gt; Vaccine effect at the same time interval for the ChAdOx1 vaccine was 88% (95% CI 75–94). &gt; Results of combined vaccine effects against hospital admission due to COVID-19 were similar when restricting the analysis to those aged 80 years and older (83%, 95% CI 72–89 at 28–34 days post-vaccination). Mass roll-out of the first doses of the BNT162b2 mRNA and ChAdOx1 vaccines was associated with substantial reductions in the risk of hospital admission due to COVID-19 in Scotland.</td>
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### Aim:
To assess the clinical characteristics, outcomes and factors associated with hospital admission or death in adult outpatients with COVID-19.

### Methods
> Prospective cohort of outpatients with suspected or confirmed COVID-19, registered in Covidom telesurveillance solution for home monitoring of patients with COVID-19 in the Greater Paris area, from March to August 2020.

### Findings:
> Among 43,103 patients, mean age was 42.9 years (SD=14.3); 93.0% (n=40,081) of patients were < 65 years old and 61.9% (n=26,688) were women. Of these 43,103 patients, 67.5% (n=29,104) completed a medical questionnaire on comorbidities and symptoms.
> The main reported comorbidities were asthma (12.8%; n=3,685), hypertension (12.3%; n=3,546) and diabetes (4.8%; n=1,385).
> A small proportion of all eligible patients (4.1% [95% CI: 3.9–4.2]; 1,751/43,103) experienced clinical worsening. The rate of hospitalisation was 4.0% (95% CI: 3.8–4.2; n=1,728) and 0.1% (95% CI: 0.1–0.2; n=64) died.
> Probability of worsening was reduced with anosmia/ageusia.

Clinical worsening was rare among outpatients. Male sex, older age and comorbidities such as chronic renal disease, active cancers or obesity were independently associated with clinical worsening. However, our cohort may include patients younger and healthier than the general population.

### Key facts
- **Aim:** to assess the clinical characteristics, outcomes and factors associated with hospital admission or death in adult outpatients with COVID-19.
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  - A small proportion of all eligible patients (4.1% [95% CI: 3.9–4.2]; 1,751/43,103) experienced clinical worsening. The rate of hospitalisation was 4.0% (95% CI: 3.8–4.2; n=1,728) and 0.1% (95% CI: 0.1–0.2; n=64) died.
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### Lancet 23APR2021

### Title: COVID-19 vaccine coverage in health-care workers in England and effectiveness of BNT162b2 mRNA vaccine against infection (SIREN): a prospective, multicentre, cohort study

### Authors and link:
Hall V.J., et al.
[goto paper](https://doi.org/10.1016/S2666-3890(21)00021-7)

### Field of expertise:
Vaccines

### Key facts
- **Aim:** prospectively determine factors associated with vaccine coverage for BNT162b2 mRNA and ChAdOx1 nCOV-19 and document BNT162b2 effectiveness in a cohort of health-care workers undergoing regular asymptomatic testing.
- **Findings:**
  - 23,324 participants from 104 sites were enrolled. Median age: 46·1 years (IQR 36·0–54·1), 19,692 (84%) were female;
  - 8,203 (35%) assigned to the positive cohort, 15,121 (65%) assigned to the negative cohort.
  - Total follow-up time was 2 months and 1,106,905 person-days (396,318 vaccinated and 710,587 unvaccinated).
  - Vaccine coverage was 89% on Feb 5, 2021, 94% had BNT162b2.
  - Significantly lower coverage was associated with previous infection, gender, age, ethnicity, job role, Index of Multiple Deprivation score.
  - During follow-up, there were 977 new infections in the unvaccinated cohort, an incidence density of 14 infections per 10,000 person-days; the vaccinated cohort had 71 new infections 21 days or more after their first dose (incidence density of 8 infections per 10,000 person-days) and nine infections 7 days after the second dose (incidence density of 0 infections per 10,000 person-days).
  - In the unvaccinated cohort, 543 (56%) participants had typical COVID-19 symptoms and 140 (14%) were asymptomatic on or 14 days before their PCR positive test date, compared with 29 (36%) with typical COVID-19 symptoms and 15 (19%) asymptomatic in the vaccinated cohort.
  - A single dose of BNT162b2 vaccine showed vaccine effectiveness of 70% (95% CI 55–85) 21 days after first dose and 85% (74–96) 7 days after two doses in the study population.

Our findings show that the BNT162b2 vaccine can prevent both symptomatic and asymptomatic infection in working-age adults. This cohort was vaccinated when the dominant variant in circulation was B1.1.7 and shows effectiveness against this variant.
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| JAMA Netw Open 23APR2021 | Association of Maternal Perinatal SARS-CoV-2 Infection With Neonatal Outcomes During the COVID-19 Pandemic in Massachusetts | Angelidou A., et al. USA [gotopaper](#) | Public health / Epidemiology | Ascertain the percentage of neonates who were born to mothers with positive SARS-CoV-2 test results during birth hospitalization, clinical and sociodemographic factors associated with neonatal test positivity, and clinical and virological outcomes for newborns during hospitalization and 30 days after discharge.  
Methods:  
> Multicenter cohort study. Neonates were born to mothers with positive SARS-CoV-2 test results within 14 days before to 72h after delivery, and were followed up for 30 days after hospital discharge.  
Primary outcomes for neonates: (1) positive SARS-CoV-2 test results, (2) indicators of adverse health, and (3) clinical signs and viral testing.  
Findings:  
> Of the 255 neonates who were born to mothers with SARS-CoV-2 infection, 225 (88.2%) were tested for SARS-CoV-2 and 5 (2.2%) had positive results during the birth hospitalization.  
> High maternal social vulnerability was associated with higher likelihood of neonatal test result positivity (adjusted odds ratio, 4.95; 95% CI, 1.53-16.01; P = .008), adjusted for maternal COVID-19 symptoms, delivery mode, and rooming-in practice.  
> Adverse outcomes during hospitalization were associated with preterm delivery indicated by worsening maternal COVID-19 symptoms.  
> Of the 151 newborns with follow-up data, 28 had nonroutine clinical visits, 7 underwent SARS-CoV-2 testing, and 1 had a positive result.  
Conclusion: Newborns exposed to SARS-CoV-2 were at risk for both direct and indirect adverse health outcomes. |
| Science 23APR2021 | Resurgence of SARS-CoV-2: detection by community viral surveillance | Riley S., et al. UK [gotopaper](#) | Public Health / Epidemiology | Aims: To estimate prevalence of COVID-19 in England through a community-wide national representative surveillance program in England (REACT-1 study); to detect resurgence from low prevalence.  
Methods:  
> Repeated random population-based sampling: 2.4 Million people were invited to join the study over 4 rounds from May 2020 to beginning of September 2020; 596,000 tested swabs were obtained (overall response rate of 25%)  
> Estimation of prevalence in this random sample (correcting for variation in response rate), over time, fitting a model of constant exponential growth and decay, and a model with a flexible p-spline.  
> Geographical variation in prevalence investigated by fitting a spatio-temporal logistic model  
Findings:  
> More reliable estimates of prevalence than from routine case data, is affected by test availability and test-seeking behaviour  
> Detection of epidemic resurgence in the summer 2020 (between end of July and mid-August), that led to the announcement of the "rule of six" social distancing measure by the UK government  
> Substantial variations in age patterns over time; the second wave started in young adults;  
> Case data (routine surveillance) consistently underestimates infections at 5-14yo compared to random population based sampling  
> Higher prevalence (x2) in participants of Asian ethnicity, also higher in Black people – higher rates of hospitalization and mortality from COVID-19 for minority ethnic groups in England may therefore reflect their higher rates of infection rather than a poorer prognosis once infected.  
> Spatial heterogeneity in prevalence detected at sub-regional level  
Conclusions:  
> Demonstration of the capability of a large national community surveillance program to detect a resurgence of SARS-CoV-2 infection at low prevalence.  
> The prevalence in the 5-14 age group is higher in this random testing study compared to case data. |
Maternal and Neonatal Morbidity and Mortality Among Pregnant Women With and Without COVID-19 Infection


Public Health / Epidemiology

Evaluation of the risks associated with COVID-19 in pregnancy on maternal and neonatal outcomes compared with not-infected, concomitant pregnant individuals.

Methods:
Cohort study involving 43 institutions in 18 countries, 2 unmatched, consecutive, not-infected women concomitantly enrolled immediately after each infected woman was identified, at any stage of pregnancy or delivery. Women and neonates were followed up until hospital discharge.

Main Outcomes: indices of (maternal and severe neonatal/perinatal) morbidity and mortality (adjusted for country, month entering study, maternal age, and history of morbidity)

Findings:
> A total of 706 pregnant women with COVID-19 diagnosis and 1424 pregnant women without COVID-19 diagnosis enrolled
> Broadly similar demographic characteristics (mean [SD] age, 30.2 [6.1] years).
> Overweight early in pregnancy occurred in 323 women (48.6%) with COVID-19 diagnosis and 554 women (40.2%) without.
> Women with COVID-19 diagnosis were at higher risk for preeclampsia/eclampsia (relative risk [RR], 1.76; 95% CI, 1.27-2.43), severe infections (RR, 3.38; 95% CI, 1.63-7.01), intensive care unit admission (RR, 5.04; 95% CI, 3.13-8.10), maternal mortality (RR, 22.3; 95% CI, 2.88-172), preterm birth (RR, 1.99; 95% CI, 1.30-1.94), medically indicated preterm birth (RR, 1.97; 95% CI, 1.56-2.51), severe neonatal morbidity index (RR, 2.66; 95% CI, 1.69-4.18), and severe perinatal morbidity and mortality index (RR, 2.14; 95% CI, 1.66-2.75).
> Fever and shortness of breath for any duration was associated with increased risk of severe maternal complications (RR, 2.56; 95% CI, 1.92-3.40) and neonatal complications (RR, 4.97; 95% CI, 2.11-11.69).
> Asymptomatic women with COVID-19 diagnosis remained at higher risk only for maternal morbidity (RR, 1.24; 95% CI, 1.00-1.54) and preeclampsia (RR, 1.63; 95% CI, 1.01-2.63).
> Among women who tested positive (98.1% by real-time polymerase chain reaction), 54 (13%) of their neonates tested positive.
> Cesarean delivery (RR, 2.15; 95% CI, 1.18-3.91) but not breastfeeding (RR, 1.10; 95% CI, 0.66-1.85) was associated with increased risk for neonatal test positivity.

Conclusions:
COVID-19 in pregnancy was associated with consistent and substantial increases in severe maternal morbidity and mortality and neonatal complications, compared to healthy pregnant women.

Nature

High-dimensional characterization of post-acute sequelae of COVID-19

Al-Aly Z., et al. USA gotopaper

Clinics

Aim: To identify incident sequelae in 30-day survivors of COVID-19 from the national healthcare databases (US Dept. of Veterans Affairs).

Methods: Approach to comprehensively identify the 6-months outcomes of incident diagnoses (from 379 diagnostic categories), incident medication use (from 380 medication classes), and incident laboratory abnormalities (from 62 laboratory tests) in people who survived the first 30 days of COVID-19. The cohort included 73,435 users.

Findings:
> Beyond the first 30 days of illness, people with COVID-19 exhibit higher risk of death and health resource utilization.
> Increased incidence of several therapeutics including pain medications (opioids and non-opioids), antidepressants, anxiolytics, antihypertensives, and oral hypoglycemics and evidence of laboratory abnormalities in multiple organ systems.
> Analysis of an array of pre-specified outcomes reveals a risk gradient that increased across severity of the acute COVID-19 infection (non-hospitalized, hospitalized, admitted to intensive care).
> Beyond the acute illness, substantial burden of health loss — spanning pulmonary and several extrapulmonary organ systems — is experienced by COVID-19 survivors.

The results provide a roadmap to inform health system planning and development of multidisciplinary care strategies to reduce chronic health loss among COVID-19 survivors.
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> COVID-19 patient sera activates the complement system and kills infected cells by Antibody-Dependent Cellular Cytotoxicity (ADCC).  
> Sera from asymptomatic individuals harbor polyfunctional antibodies. They neutralize the virus, activate ADCC and trigger complement deposition.  
> Antibody levels and functions are lower in asymptomatic individuals than in symptomatic cases. Antibody functions are correlated, regardless of disease severity.  
> Longitudinal samplings show that antibody functions follow similar kinetics of induction and contraction. Overall, asymptomatic SARS-CoV-2 infection elicits polyfunctional antibodies neutralizing the virus and targeting infected cells. |
| PNAS 21APR2021 | Dynamic prioritization of COVID-19 vaccines when social distancing is limited for essential workers | Buckner J.H., et al. USA [gotopaper](#) | Public Health / Epidemiology | Aims: To evaluate the optimal allocation of a limited vaccine supply in the United States across groups defined by age and essential worker status, which constrains opportunities for social distancing.  
Methods:  
> Compartmental model of transmission dynamics capturing key sources of group heterogeneity. The model contains 9 epidemiological status per group, 6 age classes, 2 groups depending on the essential worker status for age classes 20-39 and 40-59 and 4 possible locations (home, work, school, other).  
> Three alternative policy objectives are considered: minimizing infections, years of life lost, or deaths.  
> Assuming vaccines are available for 60% of the population for the first 6 months, the optimal vaccine allocation strategy, that evolves with the epidemiological state of the population, is computed.  
Findings:  
> The model predicts that older essential workers should be targeted first whatever the objective.  
> With the objective of minimizing infection, younger essential workers must be prioritized next.  
> With the objective of reducing mortality, older age classes must be prioritized next.  
> The dynamic optimal policy outperforms an untargeted approach from 17% to 44%, depending on the objective, the vaccine effectiveness and non-pharmaceutical interventions.  
> There are trade-offs in what can be achieved between the objectives. For example, policies that minimize infections result in substantially more deaths than a policy that minimizes deaths.  
> The optimal prioritization is sensitive to several factors, most notably vaccine effectiveness and supply, rate of transmission and the magnitude of initial infections.  
Conclusions:  
> Temporal flexibility of the allocation strategy is important to optimize public health goals.  
> Distinguishing between essential and non-essential workers is important for vaccine allocation. |
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**Primary end points:** vaccine efficacy against moderate-to-severe critical Covid-19 with an onset at least 14 days and at least 28 days after administration.

**Findings:**
> 19,630 SARS-CoV-2–negative participants who received Ad26.COV2.S and 19,691 who received placebo.
> Ad26.COV2.S protected against moderate-to-severe critical Covid-19 with onset at least 14 days after administration (116 cases in the vaccine group vs. 348 in the placebo group; efficacy, 66.9%; adjusted 95% confidence interval [CI], 59.0 to 73.4) and at least 28 days after administration (66 vs. 193 cases; efficacy, 66.1%; adjusted 95% CI, 55.0 to 74.8).
> Vaccine efficacy was higher against severe–critical Covid-19 (76.7% [adjusted 95% CI, 54.6 to 89.1] for onset at ≥14 days and 85.4% [adjusted 95% CI, 54.2 to 96.9] for onset at ≥28 days).
> Despite 86 of 91 cases (94.5%) in South Africa with sequenced virus having the 20H/501Y.V2 variant, vaccine efficacy was 52.0% and 64.0% against moderate to severe–critical Covid-19 with onset at least 14 days and at least 28 days after administration, respectively, and efficacy against severe–critical Covid-19 was 73.1% and 81.7%, respectively.
> Reactogenicity was higher with Ad26.COV2.S than with placebo but was generally mild to moderate and transient.
> The incidence of serious adverse events was balanced between the two groups. Three deaths occurred in the vaccine group (none were Covid-19–related), and 16 in the placebo group (5 were Covid-19–related).

**Conclusion:**
A single dose of Ad26.COV2.S protected against symptomatic Covid-19 and asymptomatic SARS-CoV-2 infection and was effective against severe–critical disease, including hospitalization and death. Safety appeared to be similar to that in other phase 3 trials of Covid-19 vaccines.

Assessement of data from “v-safe after vaccination health checker” surveillance system, the v-safe pregnancy registry, and the Vaccine Adverse Event Reporting System (VAERS) to characterize the initial safety of mRNA Covid-19 vaccines in pregnant persons.

**Findings:**
> 35,691 v-safe participants 16 to 54 years of age identified as pregnant.
> Injection-site pain was reported more frequently among pregnant persons than among nonpregnant women, whereas headache, myalgia, chills, and fever were reported less frequently.
> Among 3958 participants enrolled in the v-safe pregnancy registry, 827 had a completed pregnancy, of which 115 (13.9%) resulted in a pregnancy loss and 712 (86.1%) resulted in a live birth (mostly among participants with vaccination in the third trimester).
> Adverse neonatal outcomes included preterm birth (in 9.4%) and small size for gestational age (in 3.2%); no neonatal deaths were reported.
> Although not directly comparable, calculated proportions of adverse pregnancy and neonatal outcomes in persons vaccinated against Covid-19 who had a completed pregnancy were similar to incidences reported in studies involving pregnant women that were conducted before the Covid-19 pandemic.
> Among 221 pregnancy-related adverse events reported to the VAERS, the most frequently reported event was spontaneous abortion (46 cases).

**Conclusions:**
Preliminary findings did not show obvious safety signals among pregnant persons who received mRNA Covid-19 vaccines.
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<td><a href="#">gotopaper</a></td>
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<td>&gt; Retrospective analysis of data from the Israeli Ministry of Health (28 August 2020–24 February 2021)</td>
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<td>&gt; Temporal dynamics of the number of new COVID-19 cases and hospitalizations after the vaccination campaign, initiated on 20 December 2020.</td>
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<td>To distinguish the possible effects of the vaccination on cases and hospitalizations from other factors, including a third lockdown (8 January 2021) 3 comparison were performed: (1) individuals aged 60 years and older prioritized to receive the vaccine first versus younger age groups (2) the January lockdown versus the September lockdown (3) early-vaccinated versus late-vaccinated cities.</td>
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<td>&gt; 2 months after the initiation of the vaccination campaign, with 85% of individuals older than 60 years already vaccinated with two doses (24 February 2021), there was an approximately 77% drop in cases, a 45% drop in positive test percentage, a 68% drop in hospitalizations and a 67% drop in severe hospitalizations compared to peak values</td>
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<td>&gt; consecutive drops in younger age groups later, according to the order of vaccine prioritization, including earlier drops in some young age groups (16–21 years) prioritized over older age groups (21–35 years).</td>
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<td>&gt; Similar pattern of a larger and faster decline of cases and hospitalizations in older individuals during the previous lockdown implemented in Israel (between 18 September 2020 and 18 October 2020) were not observed.</td>
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<td>Analysis of large-scale, real-world data from Israel demonstrating real-life effectiveness of a national vaccination campaign</td>
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<td>Cell 20APR2021</td>
<td>Transmission, infectivity, and neutralization of a spike L452R SARS-CoV-2 variant</td>
<td>Deng X., et al. USA</td>
<td>Public Health / Epidemiology - Variants</td>
<td>&gt; We identified an emerging SARS-CoV-2 variant by viral whole-genome sequencing of 2,172 nasal/nasopharyngeal swab samples from 44 counties in California, a state in the Western United States</td>
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<td><a href="#">gotopaper</a></td>
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<td>&gt; Named B.1.427/B.1.429 to denote its 2 lineages, the variant emerged in May 2020 and increased from 0% to &gt;50% of 42 sequenced cases from September 2020 to January 2021</td>
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<td>&gt; Showing 18.6–24% increased transmissibility relative to wild-type circulating strains</td>
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<td>&gt; The variant carries 3 mutations in the spike protein, including an L452R substitution.</td>
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<td>&gt; 2-fold increased B.1.427/B.1.429 viral shedding in vivo and increased L452R pseudovirus infection of cell cultures and lung organoids albeit decreased relative to pseudoviruses carrying the N501Y mutation common to variants B.1.1.7, B.1.351, and P.1</td>
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<td>&gt; Antibody neutralization assays revealed 4.0 to 6.7-fold and 2.0-fold decreases in neutralizing titers from convalescent patients and vaccine recipients, respectively</td>
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<td>The increased prevalence of a more transmissible variant in California exhibiting decreased antibody neutralization warrants further investigation</td>
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<td>Lancet Infect Dis. 19APR2021</td>
<td>Safety and immunogenicity of SARS-CoV-2 recombinant protein vaccine formulations in healthy adults: interim results of a randomised, placebo-controlled, phase 1–2, dose-ranging study</td>
<td>Goepfert P.A., et al. gotopaper</td>
<td>Vaccines</td>
<td>Interim safety and immunogenicity results of the first-in-human study of the CoV2 preS dTM vaccine with two different adjuvant formulations. &gt; Phase 1–2, randomised, double-blind study in healthy, SARS-CoV-2-seronegative adults in ten clinical research centres in the USA. &gt; Stratified by age (18–49 years and ≥50 years) &gt; One dose (on day 1) or two doses (on days 1 and 22) of placebo or candidate vaccine, containing low-dose (effective dose 1.3 μg) or high-dose (2.6 μg) antigen with adjuvant AF03 (Sanofi Pasteur) or AS03 (GlaxoSmithKline) or unadjuvanted high-dose antigen (18–49 yrs only).  <strong>Primary endpoints:</strong> safety (up to day 43), and immunogenicity (SARS-CoV-2 neutralising antibodies on 1, 22, and 36.</td>
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<td>Findings &gt; Interim safety analyses included 439 (&gt;99%) of 441 randomly assigned participants (299 aged 18–49 years and 140 aged ≥50 years). &gt; Nab titres analysed in 326 (74%) of 441 participants (235 [79%] of 299 aged 18–49 years and 91 [64%] of 142 aged ≥50 years). &gt; No vaccine-related unsolicited immediate adverse events, serious adverse events, medically attended adverse events classified as severe, or adverse events of special interest. &gt; Solicited local and systemic reactions of any grade after two vaccine doses were reported in 81% (95% CI 61–93; 21 of 26) of participants in the low-dose plus AS03 group, 93% (84–97; 74 of 80) in the low-dose plus AS03 group, 89% (70–98; 23 of 26) in the high-dose plus AS03 group, 95% (88–99; 81 of 85) in the high-dose plus AS03 group, 29% (10–56; five of 17) in the unadjuvanted high-dose group, and 21% (8–40; six of 29) in the placebo group. &gt; A single vaccine dose did not generate neutralising antibody titres above placebo levels in any group at days 22 or 36. &gt; Among participants aged 18–49 years, neutralising antibody titres after two vaccine doses were 13-1 (95% CI 6–40–26-9) in the low-dose plus AF03 group, 20-5 (13-1–32-1) in the low-dose plus AS03 group, 43-2 (20-6–90-4) in the high-dose plus AF03 group, 75-1 (50-5–112-0) in the high-dose plus AS03 group, 5-00 (not calculated, NT) in the unadjuvanted high-dose group, and 5-00 (NT) in the placebo group. &gt; Among participants aged 50 years or older, neutralising antibody titres after two vaccine doses were 8-62 (1-90–39-0) in the low-dose plus AF03 group, 12-9 (7-09–23-4) in the low-dose plus AS03 group, 12-3 (4-35–35-0) in the high-dose plus AF03 group, 52-3 (25-3–108-0) in the high-dose plus AS03 group, and 5-00 (NT) in the placebo group.</td>
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<td>Conclusions: Lower than expected immune responses, especially in the older age groups, and high reactogenicity after dose two probably due to higher than anticipated host-cell protein content and lower than planned antigen doses in the formulations tested, which was discovered during characterisation studies on the final bulk drug substance. Further studies will focus on optimal antigen formulation and dose.</td>
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**Methods**
- Phase 1, double-blind, placebo-controlled, block-randomised trial in a single clinical trial site in Brisbane, QLD, Australia. NCT04495933.
- Healthy adults (aged ≥18 to ≤55 years). No history of SARS-CoV-2 infection; randomly assigned to one of five treatment groups and received two doses via intramuscular injection 28 days apart of either placebo, clamp vaccine at 5 μg, 15 μg, or 45 μg, or one dose of clamp vaccine at 45 μg followed by placebo.
- **Primary safety endpoints:** solicited local and systemic adverse events in the 7 days after each dose and unsolicited adverse events up to 12 months after dosing.
- **Primary immunogenicity endpoints:** were antigen-specific IgG ELISA and SARS-CoV-2 microneutralisation assays assessed at 28 days after each dose.

**Findings:**
- 120 volunteers randomly assigned to groups (n=24 per group).
- 114 (95%) completed the study up to day 57 (mean age 32.5 years [SD 10.4], 65 [54%] male, 55 [46%] female).
- Both solicited reactions and unsolicited adverse events occurred at a similar frequency in participants receiving placebo and the SARS-CoV-2 clamp vaccine.
- Solicited reactions occurred in 19 (79%) of 24 participants receiving placebo and 86 (90%) of 96 receiving the SARS-CoV-2 clamp vaccine at any dose. Unsolicited adverse events occurred in seven (29%) of 24 participants receiving placebo and 35 (36%) of 96 participants receiving the SARS-CoV-2 clamp vaccine at any dose.

**Vaccination with SARS-CoV-2 clamp elicited a similar antigen-specific response irrespective of dose:** 4 weeks after the initial dose (day 29) with 5 μg dose ([GMT] 6400, 95% CI 3683–11 122), with 15 μg dose (7492, 4959–11 319), and the two 45 μg dose cohorts (8770, 5526–13 920 in the two-dose 45 μg cohort; 8793, 5570–13 881 in the single-dose 45 μg cohort); 4 weeks after the second dose (day 57) with two 5 μg doses (102 400, 64 857–161 676), with two 15 μg doses (74 725, 51 300–108 847), with two 45 μg doses (79 586, 55 430–114 268), only a single 45 μg dose (4795, 2858–8043). At day 57, 67 (99%) of 68 participants who received two doses of clamp vaccine at any concentration produced a neutralising immune response, compared with six (25%) of 24 who received a single 45 μg dose and none of 22 who received placebo. Participants receiving two doses of clamp vaccine elicited similar neutralisation titres, irrespective of dose: two 5 μg doses ([GMT] 228, 95% CI 146–356), two 15 μg doses (230, 170–312), and two 45 μg doses (239, 187–307).

**Conclusions:**
- Subunit vaccine MF59-adjuvanted, molecular clamp-stabilised recombinant spike protein elicits strong immune responses with a promising safety profile.
- However, the glycoprotein 41 peptide present in the clamp created HIV diagnostic assay interference, a possible barrier to widespread use highlighting the criticality of potential non-spike directed immunogenicity during vaccine development.
- Studies are ongoing with alternative molecular clamp trimerisation domains to ameliorate this response.

- B.1.1.7 variant is 45% (95% CI 20–60%) more transmissible than the wild-type strain, and became the dominant in Israel within 3.5 weeks.
- **Active surveillance** through focused RT-PCR testing markedly reduces the transmission of B.1.1.7 in nursing homes.
- **Prioritized vaccination** programs seem capable of preventing the spread of the B.1.1.7 variant in the elderly.
- Proactive surveillance combined with prioritized vaccination are achievable, and reduce severe illness and subsequent death.
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| Blood 16APR2021  | Efficacy of the BNT162b2 mRNA COVID-19 Vaccine in Patients with Chronic Lymphocytic Leukemia | Herishanu Y., et al. | Vaccines | The goal of this study was to determine the efficacy of COVID-19 vaccine (BNT162b2 mRNA) in patients with CLL. **Methods**  
- We evaluated humoral immune responses to BNT162b2 mRNA COVID-19 vaccine in patients with CLL and compared responses with those obtained in age-matched healthy controls.  
- Patients received two vaccine doses, 21 days apart, and antibody titers were measured using Elecsys® Anti-SARS-CoV-2S assay after administration of the second dose.  
**Findings**  
- In 167 total patients with CLL the antibody response rate was 39.5%.  
- A comparison between 52 patients with CLL and 52 sex- and age-matched healthy controls, revealed a significantly reduced response rate among patients (52% vs 100%, respectively; adjusted odds ratio=0.010, 95% CI 0.001-0.162; p<0.001).  
- Response rate was highest in patients who obtained clinical remission after treatment (79.2%), followed by 55.2% in treatment-naïve and 16% in patients under treatment at the time of vaccination.  
- None of the patients exposed to anti-CD20 antibodies <12 months prior to vaccination responded.  
- In a multivariate analysis, the independent predictors of response were younger age, females, lack of currently active treatment, IgG levels ≥550 mg/dL and IgM levels ≥40mg/dL.  
- Antibody response to BNT162b2 mRNA COVID19 vaccine in CLL patients with is markedly impaired and affected by disease activity and treatment.  
- In patients treated with either Bruton’s tyrosine kinase inhibitors or venetoclax ± anti-CD20 antibody, responses are relatively low (16.0% and 13.6%, respectively).  
In conclusion, antibody-mediated response to BNT162b2 mRNA COVID-19 vaccine in patients with CLL is markedly impaired and affected by disease activity and treatment. |
| Blood 16APR2021  | Low Neutralizing Antibody Responses Against SARS-CoV-2 in Elderly Myeloma Patients After the First BNT162b2 Vaccine Dose | Terpos E., et al. | Vaccines | We report the development of neutralizing antibodies (NAbs) against SARS-CoV-2 in MM patients (above 18 years) after the first dose of the BNT162b2 vaccine. **Methods**  
- Included 48 MM patients (29males/19females; median age: 83years, range: 59-92years) and 104controls (57males/47females; median age: 83 years, range: 65-95 years), who were vaccinated during the same period, at the same vaccination center (Greece).  
**Findings**  
- After the first dose of the vaccine, on D22, MM patients had lower NAb titers compared to controls:medianNAb inhibition titersand rangewas 20.6% (0-96.7%) for MMpatients versus 32.5% (5.2-97.3%)for controls; P<0.01. More, specifically, only12 (25.0%) MM patients vs 57 (54.8%) controls developed NAb titers ≥30% on D22.  
- The respective number of MM patients and controls who developed NAb titers ≥50% (which corresponds to clinically relevant viral inhibition11) was 4 (8.3%) and 21 (20.2%), respectively.  
- Interestingly, only one (11.1%) out of nine patients with smoldering myeloma had NAb titers of equal or more than 30% (positivity cut-off) versus 11/39 (28.2%) patients with active MM.  
- This observationis of great interest as hypoglobulinemia has been associated withinferior antibody response among patients with chronic lymphocytic leukemia and COVID-19.  
- Our data indicate that the first dose of BNT162b2leads to production of lower levels of NAbsagainst SARS-CoV-2 compared to non-MM controls of similar ageand gender and without malignant disease.  
- This low antibody response of elderly myeloma patients after the first BNT162b2 dose may not be seen in younger patients.  
- Some anti-myeloma therapies have a B-cell depleting activity which in turn may impair immune response to vaccines, whereas both myeloma microenvironment and anti-myeloma treatmentsmay impair T-cell function. |
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| Science 16APR2021 | SARS-CoV-2 within-host diversity and transmission | Lythgoe K.A., et al. | Genomics / Phylogenomics | **Aim:** Characterize SARS-CoV-2 within-host diversity and transmission.  
**Methods:** Deep-sequencing of 1313 clinical samples from the UK (including 16 assumed transmission pairs), transmission bottleneck inference with exact beta-binomial sampling method, phylogenetics.  
**Key facts:**  > Within-host viral diversity is relatively low during acute infection; selection seems to be mostly negative (removal of deleterious mutations)  > Estimation of the bottleneck size for transmission: of 1 to 8 viruses  > Narrow transmission bottleneck, so most often transmission of the majority within-host variant; but sometimes transmission of minority variant (leading to change in consensus sequence, i.e. variation at the host level), and possible transmission of mixed infection.  > Identification of spike mutations present in multiple samples with known phenotypic effect (e.g. L5F, G446V, A879V)  
**Conclusion:** Emergence of vaccine and therapeutic escape mutations likely to be rare during early infection, but observation of immune-escape variants underlines the need for continued vigilance. Key role of open, large and rigorously controlled datasets, integrating genomic, clinical and epidemiological information. |
| NEJM 16APR21 | Pathologic Antibodies to Platelet Factor 4 after ChAdOx1 nCoV-19 Vaccination | Scully M., et al. | Vaccination | **Methods:**  > 23 patients presenting thrombosis and thrombocytopenia 6 to 24 days after receiving the first dose of the ChAdOx1 nCoV-19 vaccine (AstraZeneca).  
**Findings:**  > Median age was 46 years (range, 21 to 77). 16 patients (70%) younger than 50 years. 14 patients (61%) female  > 22 patients presented with acute thrombocytopenia and thrombosis, primarily cerebral venous thrombosis, and 1 patient presented with isolated thrombocytopenia and a hemorrhagic phenotype.  > All the patients had low or normal fibrinogen levels and elevated d-dimer levels at presentation. No evidence of thrombophilia or causative precipitants was identified.  > Testing for antibodies to platelet factor 4 (PF4) was positive in 22 patients (with 1 equivocal result) and negative in 1 patient.  
**Conclusions:** A pathogenic PF4-dependent syndrome, unrelated to the use of heparin therapy, can occur after the administration of the ChAdOx1 nCoV-19 vaccine. |
| Clin Infect Dis. 16APR2021 | Impact of convalescent plasma therapy on SARS CoV-2 antibody profile in COVID-19 patients | Tang J., et al. | Therapeutics | **Aim:** to better understand the impact of convalescent plasma (CP) on antibody response in COVID-19 patients  
**Methods:** Longitudinal analysis of antibody profile on 115 sequential plasma samples from 16 hospitalized COVID-19 patients treated with either CP or standard of care  
**Findings:**  > Differential antibody kinetics was observed for antibody binding, IgM/IgG/IgA distribution, and affinity maturation in ‘survived’ vs. ‘fatal’ COVID-19 patients.  > Surprisingly, CP treatment did not predict survival. Strikingly, marked decline in neutralization titers was observed in the fatal patients prior to death, and convalescent plasma treatment did not reverse this trend.  > Irrespective of CP treatment, higher antibody affinity to the SARS-CoV-2 prefusion spike was associated with survival outcome, while sustained elevated IgA response was associated with fatal outcome in COVID-19 patients.  > Treatment of COVID-19 patients with CPAs should be carefully targeted, and effectiveness of treatment may depend on the clinical and immunological status of COVID-19 patients as well as the quality of the antibodies in the CP. |
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| **PNAS 16APR2021** | A high-throughput microfluidic nanoimmunoassay for detecting anti–SARS-CoV-2 antibodies in serum or ultralow-volume blood samples | Swank Z., et al. Switzerland gotopaper | Diagnostics | **Aim:** development of a sensitive and specific microfluidic nanoimmunoassay (NIA) for the detection of anti–SARS-CoV-2 IgG antibodies in 1,024 samples in parallel.  
**Methods:** To eliminate the need for venipuncture, they developed a low-cost, ultralow-volume whole blood sampling methods based on two commercial devices and repurposed a blood glucose test strip. The glucose test strip permits the collection, shipment, and analysis of 0.6 μL of whole blood easily obtainable from a simple finger prick. High-throughput NIA was conducted using a PDMS microfluidic device.  
**Findings:**  
> The method achieved a specificity of 100% and a sensitivity of 98% based on the analysis of 289 human serum samples (155 positive SARS-CoV-2–infected and 134 negative individuals)  
> A single researcher can achieve a throughput of one or two devices, or 512 to 1,024 samples per day (analyzed in duplicate) in a small research laboratory not dedicated or equipped for high-throughput molecular diagnostics.  
The combination of a high-throughput, highly specific and sensitive NIA and the ability to analyze minute volumes of dried blood samples have enormous potential for SARS-CoV-2 serology, epidemiological studies, vaccine trial, and therapeutic development support. |

| **Lancet Respir Med. 15APR2021** | SARS-CoV-2 seropositivity and subsequent infection risk in healthy young adults: a prospective cohort study | Letizia A.G., et al. USA gotopaper | Public Health / Epidemiology | **Aim:** To investigate the risk of subsequent SARS-CoV-2 infection among young adults seropositive for a previous infection  
**Methods:** 3249 participants (US Marine recruits, aged 18–20 years, following a 2-week unsupervised quarantine at home) were enrolled and were assessed for baseline SARS-CoV-2 IgG seropositivity, defined as a dilution of 1:150 or more on receptor-binding domain and full-length spike protein ELISA.  
**Findings:**  
> Among 189 seropositive participants, 19 (10%) had at least one positive PCR test for SARS-CoV-2 during the 6-week follow-up (1·1 cases per person-year).  
> In contrast, 1079 (48%) of 2247 seronegative participants tested positive (6·2 cases per person-year) IR 0.18.  
> Among seropositive recruits, infection was more likely with lower baseline full-length spike protein IgG titres than in those with higher baseline full-length spike protein IgG titres (hazard ratio 0·45).  
> Infected seropositive participants had viral loads that were about 10-times lower than those of infected seronegative participants.  
> Among seropositive participants, baseline neutralising titres were detected in 45 (83%) of 54 uninfected and in six (32%) of 19 infected participants during the 6 weeks of observation.  
Seropositive young adults had about one-fifth the risk of subsequent infection compared with seronegative individuals. Although antibodies induced by initial infection are largely protective, they do not guarantee effective SARS-CoV-2 neutralisation activity or immunity against subsequent infection. |
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| Brain 15APR2021  | COVID-19 neuropathology at Columbia University Irving Medical Center/New York Presbyterian Hospital | Thakur K.T., et al. USA | Clinics | Aim: to present the clinical, neuropathological, and molecular findings of 41 consecutive patients with SARS-CoV-2 infections who died and underwent autopsy in a medical center. **Findings:**  
> Hospital-associated complications were common, including 8 (20%) with deep vein thrombosis/pulmonary embolism (DVT/PE), 7 (17%) patients with acute kidney injury requiring dialysis, and 10 (24%) with positive blood cultures during admission.  
> Neuropathological examination of 20–30 areas from each brain revealed hypoxic/ischemic changes in all brains, both global and focal; large and small infarcts, many of which appeared hemorrhagic; and microglial activation with microglial nodules accompanied by neuronophagia, most prominently in the brainstem.  
> Sparse T lymphocyte accumulation was observed in either perivascular regions or in the brain parenchyma.  
> qRT-PCR revealed low to very low, but detectable, viral RNA levels in the majority of brains, although they were far lower than those in nasal epithelia.  
> RNAscope and immunocytochemistry failed to detect viral RNA or protein in brains.  

**Microglial activation, microglial nodules and neuronophagia, observed in the majority of brains, do not result from direct viral infection of brain parenchyma, but rather likely from systemic inflammation.** |
| JAMA 15APR2021  | Spike Antibody Levels of Nursing Home Residents With or Without Prior COVID-19 3 Weeks After a Single BNT162b2 Vaccine Dose | Blain, H., et al. France | Vaccines | Older adults living in nursing homes are at higher risk for severe COVID-19, and the immune response to the vaccine may differ from that of younger, healthier adults.  
**Findings:**  
> 102 residents: 60 had no prior SARS-CoV-2 infection (COVID-19), 36 had a positive RT-PCR result and were seropositive for SARS-CoV-2 N-protein IgG in June 2020, and 6 had a positive RT-PCR result or were seropositive for SARS-CoV-2 N-protein IgG.  
> All 36 residents with prior COVID-19 were seropositive for S-protein IgG after 1 vaccine dose vs 29 of 60 residents (49.2%) without prior COVID-19.  
> Among residents with prior COVID-19, the median level of S-protein IgG was 40 000 AU/mL or greater vs 48.0 AU/mL in those without prior COVID-19.  
> One resident with a positive RT-PCR result in April 2020 tested seronegative for N-protein IgG in June 2020 and January 2021; the resident had a robust S-protein IgG level (≥40 000 AU/mL).  
> Five residents were found to be seropositive for N-protein IgG in June 2020 while having repeated negative RT-PCR results in April 2020. All 5 of these residents had high levels of S-protein IgG antibody.  
> Among the 6 residents with a positive RT-PCR result or who were seropositive for N-protein IgG, the levels of S-protein IgG antibody were significantly higher than among the 60 without prior COVID-19 and were not statistically significantly different from the 36 who had a positive RT-PCR result and were seropositive for N-protein IgG.  
**Conclusions:** This preliminary study suggests that a single dose of BNT162b2 vaccine may be sufficient to obtain a high level of S-protein IgG antibody in nursing home residents previously diagnosed with COVID-19 based on RT-PCR results. |
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<tr>
<td>15APR2021</td>
<td>recovered individuals following mRNA vaccination</td>
<td>gotopaper</td>
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<td>&gt; SARS-CoV-2 naïve individuals required both vaccine doses for optimal increases in antibodies, particularly for neutralizing titers against the B.1.351 variant.</td>
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<td>&gt; Memory B cells specific for full-length spike protein and the spike receptor binding domain (RBD) were also efficiently primed by mRNA vaccination and detectable in all SARS-CoV-2 naïve subjects after the second vaccine dose, though the memory B cell response declined slightly with age.</td>
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<td>&gt; In SARS-CoV-2 recovered individuals, antibody and memory B cell responses were significantly boosted after the first vaccine dose; however, there was no increase in circulating antibodies, neutralizing titers, or antigen-specific memory B cells after the second dose.</td>
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<td>&gt; The robust boosting after the first vaccine dose strongly correlated with levels of pre-existing memory B cells in recovered individuals, identifying a key role for memory B cells in mounting recall responses to SARS-CoV-2 antigens.</td>
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<td>Robust serological and cellular priming by mRNA vaccines were demonstrated. COVID-19 recovered subjects may only require a single vaccine dose to achieve peak antibody and memory B cell responses.</td>
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<td>Science Immunol.</td>
<td>SARS-CoV-2 genome-wide T cell epitope mapping reveals immunodominance</td>
<td>Saini S.K., et al. Denmark</td>
<td>Immunology</td>
<td>Aim: to examine the full-spectrum of CD8+ T cell immunity in COVID-19, by experimentally evaluating 3141 major histocompatibility (MHC) class I-binding peptides covering the complete SARS-CoV-2 genome.</td>
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<td>14APR2021</td>
<td>and substantial CD8+ T cell activation in COVID-19 patients</td>
<td>gotopaper</td>
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<td>Results</td>
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<td>&gt; A comprehensive list of 122 immunogenic and a subset of immunodominant SARS-CoV-2 T cell epitopes was reported.</td>
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<td>&gt; Substantial CD8+ T cell recognition was observed in COVID-19 patients, with up to 27% of all CD8+ lymphocytes interacting with SARS-CoV-2-derived epitopes.</td>
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<td>&gt; Most immunogenic regions were derived from ORF1 and ORF3, with ORF1 containing most of the immunodominant epitopes.</td>
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<td>&gt; CD8+ T cell recognition of lower affinity was also observed in healthy donors toward SARS-CoV-2-derived epitopes. This pre-existing T cell recognition signature was partially overlapping with the epitope landscape observed in COVID-19 patients and may drive the further expansion of T cell responses to SARS-CoV-2 infection.</td>
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<td>&gt; The phenotype of the SARS-CoV-2-specific CD8+ T cells revealed a strong T cell activation in COVID-19 patients, while minimal T cell activation was seen in healthy individuals.</td>
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<td>&gt; Patients with severe disease displayed significantly larger SARS-CoV-2-specific T cell populations compared to patients with mild diseases and these T cells displayed a robust activation profile.</td>
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<td>These results further the understanding of T cell immunity to SARS-CoV-2 infection and hypothesize that strong antigen-specific T cell responses are associated with different disease outcomes.</td>
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<td>Clin Infect Dis.</td>
<td>Viral sequencing reveals US healthcare personnel rarely become infected</td>
<td>Braun K.M., et al. USA</td>
<td>Public Health /</td>
<td>Aim: to infer the most likely source of infection in health personnel (HCP) by combining epidemiological data and viral sequences from healthcare and the general community.</td>
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<td>15APR2021</td>
<td>with SARS-CoV-2 through patient contact</td>
<td>gotopaper</td>
<td>Epidemiology</td>
<td>&gt; SARS-CoV-2 infection clusters involving 95 HCP and 137 possible patient contact sequences.</td>
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<td>&gt; The majority of HCP infections could not be linked to a patient or co-worker (55/95; 57.9%) and were genetically similar to viruses circulating concurrently in the community.</td>
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<td>&gt; 10.5% of infections could be traced to a co-worker (10/95). Strikingly, only 4.2% of HCP infections could be traced to a patient source (4/95).</td>
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<td>This study found no evidence for healthcare-associated transmission in the majority of HCP infections evaluated. It appears that HCP most commonly becomes infected with SARS-CoV-2 via community exposure.</td>
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> Levels of sero-neutralization and IgG rates against the ancestral strain decreased significantly over time. After 6 months, 2.8% of the patients had a negative serological status for both anti-S (spike) and anti-NP (Nucleocapsid) IgG.  
> All sera had a persistent and effective neutralizing effect against SARS-CoV-2. IgG levels correlated with sero-neutralization and this correlation was stronger for anti-S than for anti-NP antibodies.  
> The level of sero-neutralization quantified at 6 months correlated with markers of initial severity, notably ICU admission and the need for mechanical invasive ventilation.  
> Sera collected at 6 months showed efficient neutralizing effects against D614G, B.1.1.7 and P.1 variants but a significantly weaker activity against B.1.351 variant.  
> These results indicate a sustained humoral response against the ancestral strain and the D614G, B.1.1.7 and P.1 variants for at least 6 months in patients previously hospitalized for COVID-19. |
| Science 14APR2021 | Genomics and epidemiology of the P.1 SARS-CoV-2 lineage in Manaus, Brazil | Faria N.R., et al. UK [gotopaper](#) | Public Health / Epidemiology - Variants | Genome sequencing of viruses sampled in Manaus between November 2020 and January 2021 revealed the emergence and circulation of a novel SARS-CoV-2 variant of concern: Investigate the emergence of the P.1 lineage and explore epidemiological explanations for the resurgence of COVID-19 in Manaus.  
**Methods**  
> Using genomic data, structure-based mapping of mutations of interest onto the spike protein, and dynamical epidemiology modelling of genomic and mortality data (two-category dynamical model that integrates genomic and mortality data)  
> We sequenced SARS-CoV-2 genomes from 184 samples from patients seeking COVID-19 testing in two diagnostic laboratories in Manaus between November and December 2020, using the ARTIC V3 multiplexed amplicon scheme (24) and the MinION sequencing platformings  
**Findings**  
> Lineage P.1, acquired 17 mutations, including a trio in the spike protein (K417T, E484K and N501Y) associated with increased binding to the human ACE2 receptor.  
> Molecular clock analysis shows that P.1 emergence occurred around mid-November 2020 and was preceded by a period of faster molecular evolution.  
> We estimate that P.1 may be 1.7-2.4-fold more transmissible, and that previous (non-P.1) infection provides 54–79% of the protection against infection with P.1 that it provides against non-P.1 lineages.  
> The B.1.1.7 lineage exhibits similar evolutionary characteristics, which was hypothesized to have occurred in a chronically infected or immunocompromised patient.  
> Our results further show that natural immunity waning alone is unlikely to explain the observed dynamics in Manaus, with support for P.1 possessing altered epidemiological characteristics robust to a range of values assumed for the date of the lineage’s emergence and the rate of natural immunity waning.  
Enhanced global genomic surveillance of variants of concern, which may exhibit increased transmissibility and/or immune evasion, is critical to accelerate pandemic responsiveness. Studies to evaluate real-world vaccine efficacy in response to P.1 are urgently needed. |
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| NEJM 14APR21     | Thrombotic Thrombocytopenia after Ad26.COV2.S Vaccination | Muir K.L., *et al.* USA [gotopaper](#) | Vaccines | **Case report:** of extensive thrombosis associated with severe thrombocytopenia and disseminated intravascular coagulation that resembled autoimmune heparin-induced thrombocytopenia in a patient who had received the Ad26.COV2.S vaccine (Johnson & Johnson/Janssen) (Correspondace)  
  > 48-year-old woman, PCR - for SARS CoV 2. Receiving the Ad26.COV2.S vaccine 14 days before symptom onset  
  > Mild anemia and severe thrombocytopenia. Marked reduction in the platelet count with occasional schistocytes, prolonged activated partial thromboplastin time, and a marked elevation in the d-dimer level, indicating a disseminated intravascular coagulation–like state.  
  > Screening test for antibodies against platelet factor 4 (PF4)--heparin by latex-enhanced immunoassay negative. However, the result of enzyme-linked immunosorbent assay for antibodies against PF4--polyanion was strongly positive  
  **Conclusions:** Rare occurrence of vaccine-induced immune thrombotic thrombocytopenia could be related to adenoviral vector vaccines. |
| Clin Infect Dis. 14APR21 | Nosocomial outbreak of COVID-19 by possible airborne transmission leading to a superspreading event | Chi-Chung Cheng, V., *et al.* China [gotopaper](#) | Public Health / Epidemiology | Description of a nosocomial outbreaks with superspreading of COVID-19 due to a possible airborne transmission  
  **Methods:** Epidemiological analysis, environmental samplings, and whole genome sequencing (WGS) were performed for a hospital outbreak.  
  **Findings:**  
  > Superspreading event involving 12 patients and 9 healthcare workers (HCWs) occurred within 4 days in 3 of 6 cubicles at an old-fashioned general ward with no air exhaust built within the cubicles.  
  > Environmental contamination by SARS-CoV-2 RNA was significantly higher in air grilles than high-touch clinical surfaces.  
  > Six (66.7%) of 9 contaminated air exhaust grilles were located outside patient cubicle.  
  > The clinical attack rate of patients was significantly higher than HCWs (15.4%, 12/78 exposed-patients vs 4.6%, 9/195 exposed-HCWs, p=0.005).  
  > Clinical attack rate of ward-based HCWs was significantly higher than non-ward-based HCWs (8.1%, 7/68 vs 1.8%, 2/109, p=0.045).  
  > The outbreak strains belong to SARS-CoV-2 lineage, B.1.36.27 with the unique S-T470N mutation on WGS.  
  **Conclusion** This nosocomial point source superspreading due to possible airborne transmission demonstrated the need for stringent SARS-CoV-2 screening at admission to healthcare facilities and better architectural design of the ventilation system to prevent such outbreaks. Portable high-efficiency particulate filters were installed in each cubicle to improve ventilation before resumption of clinical service. |
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<td><strong>Methods:</strong> retrospective case-control study across a single healthcare system of non-hospitalized patients, with documented positive SARS-CoV-2 testing, risk factors for severe COVID-19, and referrals for bamlanivimab via emergency use authorization.</td>
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<td><strong>Findings:</strong> &gt;The most reported and documented symptoms of COVID-19 illness at initial presentation were cough (65.8%), fever (42.3%), myalgias (37.7%), and fatigue (34.8%).</td>
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<td>&gt;The 30-day hospitalization rate was significantly lower among patients who received bamlanivimab (7.3% v 20.0%, RR 0.37), and the number needed to treat was 8.</td>
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<td>&gt;On logistic regression, odds of hospitalization were increased in patients not receiving bamlanivimab and with a higher number of pre-specified comorbidities (OR 4.19 CI: 1.31-2.16, p&lt;0.001; OR 1.68, CI: 2.12-8.30, p&lt;0.001, respectively).</td>
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<td><strong>Ambulatory patients with COVID-19 who received bamlanivimab had a lower 30-day hospitalization than control patients in real-world experience.</strong></td>
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<td>Br J Sports Med.</td>
<td>Physical inactivity is associated with a higher risk for severe COVID-19 outcomes: a study in 48 440 adult patients</td>
<td>Sallis R., et al. USA gotopaper</td>
<td>Public Health / Epidemiology</td>
<td><strong>Aim:</strong> To compare hospitalisation rates, ICU admissions and mortality for 48 440 patients with COVID-19 who were consistently inactive, doing some activity or consistently meeting physical activity guidelines.</td>
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<td>&gt; Patients with COVID-19 who were consistently inactive had a greater risk of hospitalisation (OR 2.26; 95% CI 1.81-2.83), admission to the ICU (OR 1.73; 95% CI 1.18-2.55) and death (OR 2.49; 95% CI 1.34-6.7) due to COVID-19 than patients who were consistently meeting physical activity guidelines.</td>
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<td>&gt; Patients who were consistently inactive also had a greater risk of hospitalisation (OR 1.20; 95% CI 1.10-1.32), admission to the ICU (OR 1.10; 95% CI 0.93-1.29) and death (OR 1.32; 95% CI 1.09-1.60) due to COVID-19 than patients who were doing some physical activity.</td>
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<td>Consistently meeting physical activity guidelines was strongly associated with a reduced risk for severe COVID-19 outcomes among infected adults.</td>
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<td>&gt; 504 breast milk samples from 84 women (weekly sampling for 6 weeks from week 2 after one dose of vaccine).</td>
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<td>&gt; Mean levels of anti-SARS-CoV-2–specific IgA antibodies in the breast milk increased rapidly and were significantly elevated at 2 weeks after the first vaccine (2.05 ratio; P &lt; .001), when 61.8% of samples tested positive, increasing to 86.1% at week 4 (1 week after the second vaccine). Mean levels remained elevated for the duration of follow-up. At week 6, 65.7% of samples tested positive.</td>
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<td>&gt; Anti–SARS-CoV-2–specific IgG antibodies increased at week 4 (20.5 U/mL; P = .004), when 91.7% of samples tested positive, increasing to 97% at weeks 5 and 6.</td>
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<td>&gt; No mother or infant experienced any serious adverse event during the study period. Mild vaccine-related adverse effects were observed in vaccinees, and fever with upper respiratory tract symptoms were observed in 4 infants.</td>
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<td>SARS-CoV-2–specific IgA and IgG antibodies in breast milk after vaccination were found. These showed strong neutralizing effects, suggesting infant protection.</td>
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Changes in symptomatology, reinfection, and transmissibility associated with the SARS-CoV-2 variant B.1.1.7: an ecological study

Graham M.S., et al.
Public Health / Epidemiology - Variants

Aim to investigate whether increases in the proportion of infections with this variant are associated with differences in symptoms or disease course, reinfection rates, or transmissibility.

Methods
> Data on types and duration of symptoms were obtained from longitudinal reports from users of the COVID Symptom Study app who reported a positive test for COVID-19.
> We assessed the Spearman correlation between the proportion of B.1.1.7 cases and number of reinfections over time, and between the number of positive tests and reinfections.

Findings
> From Sept 28 to Dec 27, 2020, positive COVID-19 tests were reported by 36920 COVID Symptom Study app users whose region was known and who reported as healthy on app sign-up. We found no changes in reported symptoms or disease duration associated with B.1.1.7.
> For the same period, possible reinfections were identified in 249 (0.7% [95% CI 0.6–0.8]) of 36509 app users who reported a positive swab test before Oct 1, 2020, but there was no evidence that the frequency of reinfections was higher for the B.1.1.7 variant than for pre-existing variants. Reinfection occurrences were more positively correlated with the overall regional rise in cases (Spearman correlation 0.56–0.69 for South East, London, and East of England) than with the regional increase in the proportion of infections with the B.1.1.7 variant (Spearman correlation 0.38–0.56 in the same regions), suggesting B.1.1.7 does not substantially alter the risk of reinfection.

> We found a multiplicative increase in the $R_t$ of B.1.1.7 by a factor of 1.35 (95% CI 1.02–1.69) relative to pre-existing variants. However, $R_t$ fell below 1 during regional and national lockdowns, even in regions with high proportions of infections with the B.1.1.7 variant.

The lack of change in symptoms identified in this study indicates that existing testing and surveillance infrastructure do not need to change specifically for the B.1.1.7 variant. In addition, given that there was no apparent increase in the reinfection rate, vaccines are likely to remain effective against the B.1.1.7 variant.

Genomic characteristics and clinical effect of the emergent SARS-CoV-2 B.1.1.7 lineage in London, UK: a whole-genome sequencing and hospital-based cohort study

Frampton D., et al.
Public Health / Epidemiology - Variants

Describe the emergence of the B.1.1.7 variant of concern (VOC), including virological characteristics and clinical severity in contemporaneous patients with and without the variant.

Methods
> In this cohort study, samples positive for SARS-CoV-2 on PCR that were collected from Nov 9, 2020, for patients acutely admitted to one of two hospitals on or before Dec 20, 2020, in London, UK.
> Poisson regression models to investigate the association between B.1.1.7 infection and severe disease.

Findings
> Of 496 patients with samples positive for SARS-CoV-2 on PCR and who met inclusion criteria, 341 had samples that could be sequenced. 198 (58%) of 341 had B.1.1.7 infection and 143 (42%) had non-B.1.1.7 infection.
> No evidence of an association between severe disease and death and lineage (B.1.1.7 vs non-B.1.1.7) in unadjusted analyses (prevalence ratio [PR] 0.97 [95% CI 0.72–1.31]), or in analyses adjusted for hospital, sex, age, comorbidities, and ethnicity (adjusted PR 1.02 [0.76–1.38]).
> We detected no B.1.1.7 VOC-defining mutations in 123 chronically shedding immunocompromised patients or in 32 remdesivir-treated patients.
> Viral load by proxy was higher in B.1.1.7 samples than in non-B.1.1.7 samples, as measured by cycle threshold value (mean 28.8 [SD 4.7] vs 32.0 [4.8]; p=0.0085) and genomic read depth (1280 [1004] vs 831 [682]; p=0.0011).

Emerging evidence exists of increased transmissibility of B.1.1.7, and we found increased virus load by proxy for B.1.1.7 in our data. We did not identify an association of the variant with severe disease.
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| Nature Commun. 09APR2021 | Seroprevalence and correlates of SARS-CoV-2 neutralizing antibodies from a population-based study in Bonn, Germany | Aziz N.A., et al. Germany gotopaper | Immunology | **Aim:** to estimate the seroprevalence and temporal course of SARS-CoV-2 neutralizing antibodies.  
Anti-SARS-CoV-2 IgG levels were assessed by immunoassay, followed by confirmatory testing of borderline and positive test results with a recombinant spike-based immunofluorescence assay and a plaque reduction neutralization test (PRNT). Borderline or positive individuals were retested after 4-5 months.  
> At baseline, 4771 persons participated (April 24th - June 30th, 2020).  
> **Seroprevalence** was 0.97% (95% CI: 0.72–1.30) by immunoassay and 0.36% (95% CI: 0.21–0.61) when considering only those with two additional positive confirmatory tests.  
> Antibody response magnitude, total number of symptoms experienced, and presence of particular symptoms were **associated with the presence of neutralizing antibodies** in those with a positive immunoassay test result.  
> In those with a **borderline immunoassay result**, the presence of neutralizing antibodies was **extremely rare** and apparently transient.  
> About 20% of PRNT+ individuals lost their neutralizing antibodies **within 5 months**. Neutralizing antibodies are detectable in only one third of those with a positive immunoassay result, and **wane relatively quickly**.  
> The probability of neutralizing antibody loss was inversely related to the magnitude of the IgG response.  
> Self-referral bias can lead to substantial overestimation of seroprevalence. |
| Lancet 09APR2021 | SARS-CoV-2 infection rates of antibody-positive compared with antibody-negative health-care workers in England: a large, multicentre, prospective cohort study (SIREN) | Hall V.J., et al. UK gotopaper | Public Health / Epidemiology | Investigate whether antibodies against SARS-CoV-2 were associated with a decreased risk of symptomatic and asymptomatic reinfection.  
**Methods**  
> The primary outcome was a reinfection in the positive cohort or a primary infection in the negative cohort, determined by PCR tests.  
> A proportional hazards frailty model using a Poisson distribution was used to estimate incidence rate ratios (IRR) to compare infection rates in the two cohorts.  
**Findings**  
> From June 18, 2020, to Dec 31, 2020, 30625 participants were enrolled into the study. 51 participants withdrew from the study, 4913 were excluded, and 25661 participants (with linked data on antibody and PCR testing) were included in the analysis. Data were extracted from all sources on Feb 5, 2021, and include data up to and including Jan 11, 2021.  
> 155 infections were detected in the baseline positive cohort of 8278 participants, collectively contributing 2 047 113 person-days of follow-up. This compares with 1704 new PCR positive infections in the negative cohort of 17383 participants, contributing 2971436 person-days of follow-up.  
> The incidence density was 7.6 reinfections per 100 000 person-days in the positive cohort, compared with 53.7 primary infections per 100 000 person-days in the negative cohort, between June, 2020, and January, 2021.  
> The adjusted IRR was 0.159 for all reinfections (95% CI 0.13–0.19) compared with PCR-confirmed primary infections. The median interval between primary infection and reinfection was more than 200 days.  
A previous history of SARS-CoV-2 infection was associated with an 84% lower risk of infection, with median protective effect observed 7 months following primary infection. This time period is the minimum probable effect because seroconversions were not included. This study shows that previous infection with SARS-CoV-2 induces effective immunity to future infections in most individuals. |
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| NEJM 09APR21     | Thrombosis and Thrombocytopenia after ChAdOx1 nCov-19 Vaccination | Schultz, N., et al. Norway gotopaper | Vaccines | **Case report:** findings in five patients who presented with venous thrombosis and thrombocytopenia 7 to 10 days after receiving the first dose of the AZ vaccine against Covid-19.  
**Findings:**  
> Health care worker, 32 to 54 years of age.  
> All five patients were negative for antibodies to SARS-CoV-2 nucleocapsid protein.  
> All five patients had high levels of antibodies to platelet factor 4–polyanion complexes;  
> No previous exposure to heparin.  
> Platelets in serum from Patients 1, 3, 4, and 5 were clearly activated in the absence of added heparin  
> Four of the patients had severe cerebral venous thrombosis with intracranial hemorrhage, and the outcome was fatal in three.  
**Conclusions:** Findings indicate a shared pathophysiological basis of the condition in these five patients and should raise awareness that a syndrome similar to autoimmune heparin-induced thrombocytopenia may occur in some persons after vaccination with AZ vaccine (five cases in a population of more than 130,000 vaccinated persons) |
| NEJM 09APR21     | Thrombotic Thrombocytopenia after ChAdOx1 nCov-19 Vaccination | Greinacher, A., et al. International gotopaper | Vaccines | **Aim**  
Assessment of clinical and laboratory features of 11 patients in Germany and Austria developing thrombosis or thrombocytopenia after AZ vaccination.  
**Methods**  
ELISA detection of platelet factor 4 (PF4)–heparin antibodies and a modified (PF4-enhanced) platelet-activation test to detect platelet-activating antibodies under various reaction conditions  
Included in this testing were samples from patients who had blood samples referred for investigation of vaccine-associated thrombotic events, with 28 testing positive on a screening PF4–heparin immunosassay.  
**Findings**  
> 11 patients, including 9 women. Median age: 36 years (22 to 49).  
> Patients presented with one or more thrombotic events beginning 5 to 16 days after vaccination. 9 had cerebral venous thrombosis, 3 had splanchnic-vein thrombosis, 3 had pulmonary embolism, and 4 had other thromboses; of these patients, 6 died. 5 patients had disseminated intravascular coagulation.  
> One patient presented with fatal intracranial hemorrhage.  
> None of the patients had received heparin before symptom onset.  
> All 28 patients who tested positive for antibodies against PF4–heparin tested positive on the platelet-activation assay in the presence of PF4 independent of heparin. Platelet activation was inhibited by high levels of heparin, Fc receptor–blocking monoclonal antibody, and immune globulin (10 mg per milliliter). Additional studies with PF4 or PF4–heparin affinity purified antibodies in 2 patients confirmed PF4-dependent platelet activation.  
**Conclusions** Vaccination with ChAdOx1 nCov-19 can result in the rare development of immune thrombotic thrombocytopenia mediated by platelet-activating antibodies against PF4, which clinically mimics autoimmune heparin-induced thrombocytopenia. |
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<th>Authors and link</th>
<th>Field of expertise</th>
<th>Key facts</th>
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<td>Lancet Respir Med.</td>
<td>Inhaled budesonide in the treatment of early COVID-19 (STOIC): a phase 2, open-label, randomised controlled trial</td>
<td>Ramakrishnan S., et al., UK</td>
<td>Therapeutics</td>
<td>Aim: to evaluate the efficacy of the widely used inhaled glucocorticoid budesonide in individuals with early COVID-19 in the community. Methods: open-label, parallel-group, phase 2, randomised controlled trial (Steroids in COVID-19; STOIC) of inhaled budesonide, compared with usual care, in adults within 7 days of the onset of mild COVID-19 symptoms. Primary endpoint: COVID-19-related urgent care visit, including emergency department assessment or hospitalisation. Secondary outcomes: self-reported clinical recovery (symptom resolution). Findings: &gt; For the per-protocol population (n=139), the primary outcome occurred in ten (14%) of 70 participants in the budesonide group and one (1%) of 69 participant in the usual care group. &gt; For the Intention-to-treat population, the primary outcome occurred in 11 (15%) participants in the usual care group and two (3%) participants in the budesonide group. &gt; Clinical recovery was 1 day shorter in the budesonide group compared with the usual care group (median 7 days in the budesonide group vs 8 days in the usual care group) &gt; The mean total score change in the CCQ and FLUPro over 14 days was significantly better in the budesonide group compared with the usual care group &gt; Budesonide was safe, with only five (7%) participants reporting self-limiting adverse events. Early administration of inhaled budesonide reduced the likelihood of needing urgent medical care and reduced time to recovery after early COVID-19.</td>
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<td>JAMA</td>
<td>Symptoms and Functional Impairment Assessed 8 Months After Mild COVID-19 Among Health Care Workers</td>
<td>Havervall S., et al., Sweden</td>
<td>Public Health / Epidemiology</td>
<td>Aim: to investigated COVID-19–related long-term symptoms in healthcare professionals Methods: The COMMUNITY (COVID-19 Biomarker and Immunity) study investigates long-term immunity after mild COVID-19. Between April 15, 2020, and May 8, 2020, health care professionals at Danderyd Hospital, Stockholm, Sweden, were invited to participate Findings: &gt; Seropositive participants who reported no or mild prior symptoms had a median age of 43 years and 83% were women &gt; Comparing seropositive vs seronegative participants, 26% vs 9% reported at least 1 moderate to severe symptom lasting for at least 2 months (RR, 2.9) and 15% vs 3% reported at least 1 moderate to severe symptom lasting for at least 8 months (RR, 4.4). &gt; The most common moderate to severe symptoms lasting for at least 2 months in the seropositive group were anosmia, fatigue, ageusia, and dyspnea. &gt; Of the seropositive participants, 8% reported that their long-term symptoms moderately to markedly disrupted their work life, compared with 4% of the seronegative participants (RR, 1.8). &gt; 15% reported their long-term symptoms moderately to markedly disrupted their social life, compared with 6% of the seronegative participants (RR, 2.5). &gt; 12% reported that their long-term symptoms moderately to markedly disrupted their home life, compared with 5% of the seronegative participants (RR, 2.3) A considerable portion of low-risk individuals with mild COVID-19 reported a diversity of long-term symptoms, and these symptoms disrupted work, social, and home life.</td>
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**NEJM 08APR21**

**Antibody Responses in Seropositive Persons after a Single Dose of SARS-CoV-2 mRNA Vaccine**

Krammer F., et al.
USA
Vaccines

- Immune response to one dose of BNT162b2 or mRNA-1273 in persons with previous Covid-19.

**Methods:**
110 PARIS study participants with or without documented preexisting SARS-CoV-2 immunity. 67 seronegative participants and 43 seropositive participants receiving their first spike mRNA vaccine dose in 2020

**Findings:**
- The majority of seronegative participants had variable and relatively low SARS-CoV-2 IgG responses within 9 to 12 days after vaccination. In contrast, participants with SARS-CoV-2 antibodies at baseline before the first vaccine injection rapidly developed uniform, high antibody titers within days after vaccination.
- The antibody titers of vaccinees with preexisting immunity were 10 to 45 times as high as those of vaccinees without preexisting immunity.
- No increase in antibody titers was observed in the Covid-19 survivors who received the second vaccine dose (3-fold in non-infected participants).
- No substantial difference was noted in the dynamics of antibody responses elicited by the Pfizer and Moderna vaccines after the first dose.
- Vaccine recipients with preexisting immunity had systemic side effects at higher frequencies than those without preexisting immunity (fatigue, headache, chills, muscle pain, fever, and joint pain, in order of decreasing frequency).

**Conclusion:**
A single dose of mRNA vaccine elicited rapid immune responses in seropositive participants, with postvaccination antibody titers that were similar to or exceeded titers found in seronegative participants who received two vaccinations. Whether a single dose of mRNA vaccine provides effective protection in seropositive persons requires investigation.

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**NEJM 07APR21**

**Neutralizing Response against Variants after SARS-CoV-2 Infection and One Dose of BNT162b2**

Lustig Y., et al.
Israel
Vaccines - Variants

- Aim: to investigate whether one dose of the BNT162b2 vaccine would increase neutralizing activity against the B.1.1.7, B.1.351, and P.1 variants in persons previously infected with SARS-CoV-2.

**Methods:** Microneutralization assay with isolates of the original virus (sublineage B.1) and the B.1.1.7, B.1.351, and P.1 variants on 6 HCW previously infected with the original variant of SARS-CoV-2 and vaccinated (3 time points: 1-12 weeks after natural infection, immediately before vaccination, and 1-2 weeks after vaccination).

**Findings:**
- Time point 1: Samples obtained had neutralizing activity against the original virus and the B.1.1.7 and P.1 variants, with geometric mean titers (GMT) of 456, 256, and 71, respectively, but had little or no neutralizing activity against the B.1.351 variant (GMT 8).
- Time point 2: GMT were 81, 40, 36, and 7 for the original virus and the B.1.1.7, P.1, and B.1.351 variants, respectively.
- Time point 3: GMT were 9195, 8192, 2896, and 1625 for the original virus and the B.1.1.7, P.1, and B.1.351 variants, respectively — that is, the titers after vaccination were 114, 203, 81, and 228 times as high as the titers immediately before vaccination.

This study showed that one vaccine dose substantially increased neutralizing activity against all variants tested, highlighting the importance of vaccination even in previously infected patients.
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| NEJM 07APR2021  | Cross-Reactive Neutralizing Antibody Responses Elicited by SARS-CoV-2 501Y.V2 (B.1.351) | Moyo-Gwete T., et al. South Africa | Vaccines - Variants | **Aim:** Assessment of the immune response to 501Y.V2 (B.1.351) and its cross-reactivity with other variants. Samples were collected when 501Y.V2 prevalence was 90% in Cape Town.  
**Findings:**  
> 501Y.V2 elicited high-titer binding and neutralizing antibody responses.  
> Titers of binding antibodies to RBD and the full spike protein of the original variant were highly correlated with titers of binding antibodies to the corresponding proteins of 501Y.V2.  
> Plasma samples (46) had higher titers to the spike protein of 501Y.V2 than to the spike protein of the original variant (mean of 1.7 times as high), but high-level binding to the original variant remained.  
> 53 of 57 tested samples maintained neutralization activity against the original variant, with a geometric mean titer of 203 (95% CI, 141-292), approximately one third of the titer against the 501Y.V2 variant. When limiting the analysis to 22 sequencing-confirmed infection with 501Y.V2 with positive titers of binding antibodies, the same pattern was observed.  
> Testing a subset of 10 plasma samples against the 501Y.V3 (P.1) variant revealed high levels of neutralization, with some samples showing higher potency against 501Y.V3 (P.1) than against 501Y.V2, possibly due to the very different N-terminal domains.  
501Y.V2 elicits robust neutralizing antibody responses against both the original variant and 501Y.V3 (P.1), indicating high levels of cross-reactivity. |
| NEJM 07APR2021  | Neutralization of SARS-CoV-2 Variants B.1.429 and B.1.351 | Shen X., et al. USA | Vaccines - Variants | **Aim:** to measure the neutralizing activity against SARS-CoV-2 variant B.1.429 (California) and B.1.351 (South Africa) of serum specimens obtained from 14 convalescent persons and from 49 recipients of mRNA-1273 (26) or protein nanoparticle vaccine NVX-CoV2373 (23).  
**Findings**  
> As compared with the D614G variant, **B.1.429** was approximately 2 to 3 times less sensitive to neutralization by convalescent serum and by serum samples obtained from vaccinated persons  
> **B.1.351** was approximately 9 to 14 times less sensitive to neutralization.  
**Neutralisation assays with pseudoviruses:**  
- **B.1.429** was neutralized by convalescent serum and by vaccine serum. The geometric mean ID50 titers against B.1.429 were **3.1 times (1.4-8.8) lower** than those against D614G for convalescent serum and were **2.0 and 2.5 times (0.7-8.6) lower** than against D614G for serum from persons who had received the mRNA-1273 and NVX-CoV2373 vaccines, respectively.  
- The geometric mean ID50 titer against **B.1.351** was **13.1 times lower** than against D614G for convalescent serum and **9.7 times and 14.5 times lower** than against D614G for serum from persons who had received the mRNA-1273 and NVX-CoV2373 vaccines, respectively.  
These results suggest that vaccine-elicited neutralizing antibodies are likely to remain effective against the **B.1.429** variant. The magnitude of resistance seen with the **B.1.351** variant is of greater concern with respect to current vaccines. |
## Antibody Persistence through 6 Months after the Second Dose of mRNA-1273 Vaccine for Covid-19

**Doria-Rose N., et al.**  
USA  
**gotopaper**  
**Vaccines**

The durability of protection of mRNA-1273 vaccine from Moderna currently unknown.

**Findings:**  
> mRNA1273 elicited binding and neutralizing antibodies in 33 healthy adult participants 180 days after the second dose of 100 μg (day 209).  
> S protein binding antibodies had geometric mean end-point titers of 92,451 (95% CI, 57,148 to 149,562) in participants 18 to 55 years of age, 62,424 (95% CI, 36,765 to 105,990) in those 56 to 70 years of age, and 49,373 (95% CI, 25,171 to 96,849) in those 71 years of age or older.

> All the participants had detectable neutralization activity, with 1050 GMTs of 406 (95% CI, 286 to 578), 171 (95% CI, 95 to 307), and 131 (95% CI, 69 to 251) depending on age.

> The estimated half-life of binding antibodies after day 43 for all the participants ranged between 52 and 109 days depending on the method use for assessment. The neutralizing antibody half-life estimates was between 68 and 202 days.

> Antibodies that were elicited by mRNA-1273 persisted through 6 months after the second dose, as detected by three distinct serologic assays.

Ongoing studies are monitoring immune responses beyond 6 months as well as determining the effect of a booster dose to extend the duration and breadth of activity against emerging viral variants.

**Conclusion:**  
Our data show antibody persistence and thus support the use of this vaccine in addressing the Covid-19 pandemic.

## Post-Discharge Thromboembolic Outcomes and Mortality of Hospitalized COVID-19 Patients: The CORE-19 Registry

**Giannis D., et al.**  
USA  
**gotopaper**  
**Public Health / Epidemiology**

Thromboembolic events including venous thromboembolism (VTE), arterial thromboembolism (ATE), and mortality from sub-clinical thrombotic events occur frequently in COVID-19 inpatients.

**Methods:**  
Prospective registry included consecutive COVID-19 patients hospitalized within our multi hospital system from March 1st - May 31st 2020  
Primary outcome = a composite of adjudicated VTE, ATE, and all-cause mortality (ACM)  
Principal safety outcome = major bleeding (MB)

**Findings:**  
> Among 4,906 patients (53.7% male) mean age was 61.7 years. Comorbidities included hypertension (38.6%), diabetes (25.1%), obesity (18.9%), and cancer history (13.1%)

> Post-discharge thromboprophylaxis was prescribed in 13.2%. VTE rate was 1.55%, ATE 1.71%, ACM 4.83%, and MB 1.73%.

> The composite primary outcome rate was 7.13% and was significantly associated with advanced age (OR: 3.66, 95%CI: 2.84-4.71), prior VTE (OR: 2.99, 95%CI:2.00-4.47), ICU stay (OR: 2.22, 95%CI:1.78-2.93), chronic kidney disease (CKD) (OR: 2.10, 95%CI:1.47-3.0), peripheral arterial disease (OR: 2.04, 95%CI:1.10-3.80), carotid occlusive disease (OR: 2.02,95%CI: 1.30-3.14), IMPROVE-DD VTE score ≥4 (OR: 1.51, 95%CI: 1.06-2.14), and coronary artery disease(OR: 1.50, 95%CI: 1.04-2.17).

> Post-discharge anticoagulation was significantly associated with reducing the primary outcome (OR: 0.54, 95%CI: 0.47-0.81).

**Conclusions:**  
Post-discharge VTE, ATE, and ACM occur frequently following COVID-19 hospitalization. Advanced age, cardiovascular risk factors, CKD, IMPROVE-DD VTE score ≥4, and ICU stay increase risk. Post-discharge anticoagulation reduced risk by 46%.
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<th>Authors and link</th>
<th>Field of expertise</th>
<th>Key facts</th>
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| *Lancet Psychiatry* 06APR2021 | 6-month neurological and psychiatric outcomes in 236 379 survivors of COVID-19: a retrospective cohort study using electronic health records | Taquet M., et al. UK gotopaper | Public Health / Epidemiology | **Aim:** to provide robust estimates of incidence rates and relative risks of neurological and psychiatric diagnoses in patients in the 6 months following a COVID-19 diagnosis.  
> Among 236 379 patients diagnosed with COVID-19, the estimated incidence of a neurological or psychiatric diagnosis in the following 6 months was 33·62% (95% CI 33·17–34·07), with 12·84% (12·36–13·33) receiving their first such diagnosis.  
> For patients who had been admitted to an intensive therapy unit (ITU), the estimated incidence of a diagnosis was 46·42% (44·78–48·09) and for a first diagnosis was 25·79% (23·50–28·25).  
> The whole COVID-19 cohort had estimated incidences of 0·56% (0·50–0·63) for intracranial haemorrhage, 2·10% (1·97–2·23) for ischaemic stroke, 0·11% (0·08–0·14) for parkinsonism, 0·67% (0·59–0·75) for dementia, 17·39% (17·04–17·74) for anxiety disorder, and 1·40% (1·30–1·51) for psychotic disorder, among others.  
> In the group with ITU admission, estimated incidences were 2·66% (2·24–3·16) for intracranial haemorrhage, 6·92% (6·17–7·76) for ischaemic stroke, 0·26% (0·15–0·45) for parkinsonism, 1·74% (1·31–2·30) for dementia, 19·15% (17·90–20·48) for anxiety disorder, and 2·77% (2·31–3·33) for psychotic disorder.  
> Most diagnostic categories were more common in patients who had COVID-19 than in those who had influenza (hazard ratio [HR] 1·44, 95% CI 1·40–1·47, for any diagnosis; 1·78, 1·68–1·89, for any first diagnosis) and those who had other respiratory tract infections (1·16, 1·14–1·17, for any diagnosis; 1·32, 1·27–1·36, for any first diagnosis).  
> HRs were higher in patients who had more severe COVID-19 (e.g., those admitted to ITU compared with those who were not: 1·58, 1·50–1·67, for any diagnosis; 2·87, 2·45–3·35, for any first diagnosis).  
**Substantial neurological and psychiatric morbidity were observed in the 6 months after COVID-19 infection. Risks were greatest in, but not limited to, patients who had severe COVID-19.** |
| *JAMA Netw Open* 01APR2021 | Mortality and Readmission Rates Among Patients With COVID-19 After Discharge From Acute Care Setting With Supplemental Oxygen | Banerjee J., et al. USA gotopaper | Clinics | **Aim:** to assess outcomes of patients with COVID-19 pneumonia discharged via the expected practice approach to home or quarantine housing with supplemental home oxygen.  
**Methods:** retrospective cohort study of 621 patients with COVID-19 discharged with supplemental home oxygen (at least 3 L per minute of oxygen) from emergency department and inpatient encounters at 2 large urban medical centers.  
Main Outcomes and Measures : All-cause mortality and all-cause 30-day return admission.  
**Findings:**  
> A total of 621 patients with COVID-19 pneumonia (404 male [65.1%] and 217 female [34.9%]) were discharged with home oxygen.  
> Median age of these patients was 51 years (interquartile range, 45–61 years), with 149 (24.0%) discharged from the emergency department and 472 (76%) discharged from inpatient encounters.  
> The all-cause mortality rate was 1.3% (95% CI, 0.6%–2.5%) and the 30-day return hospital admission rate was 8.5% (95% CI, 6.2%–10.7%) with a median follow-up time of 26 days (interquartile range, 15–55 days).  
> **No deaths occurred in the ambulatory setting.**  
**Ambulatory management of COVID-19 with home oxygen has an acceptable safety profile, and the expected practice approach may help optimize outcomes.** |
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| Nature Med. 01APR2021 | Antibody responses to the BNT162b2 mRNA vaccine in individuals previously infected with SARS-CoV-2 | Ebinger J.E., et al. USA gotopaper     | Vaccines           | **Background:** Detectable presence of anti-SARS-CoV-2 antibodies and virus-specific T cells suggest possible alternate vaccination strategies for previously infected individuals. As thus, individuals with prior infection might have naturally acquired immunity that could be sufficiently enhanced by a single dose rather than a double dose of administered vaccine.  
**Methods:** Cohort of BNT162b2 (Pfizer–BioNTech) mRNA vaccine recipients (n = 1,090). Antibody levels were measured at three time points: before or up to 3 d after dose 1; within 7–21 d after dose 1; and within 7–21 d after dose 2.  
**Findings:**  
> For both IgG(N) (representing response to prior infection) and IgG(S-RBD) (representing response to either prior infection or vaccine), individuals with prior SARS-CoV-2 infection had higher antibody levels at all time points.  
> IgG(S-RBD) levels were not significantly different among previously infected individuals after a single dose and infection-naive individuals who had received two doses.  
> ACE2 binding inhibition was significantly higher among previously infected individuals than infection-naive individuals after a single vaccine dose, with no between-group difference seen after the second dose.  
> Post-vaccine symptoms were more prominent for those with prior infection after the first dose, but symptomology was similar between groups after the second dose.  
**Conclusions:** Individuals previously infected with SARS-CoV-2 developed vaccine-induced antibody responses after a single dose of the BNT162b2 mRNA vaccine similar to those seen after a two-dose vaccination in infection-naive individuals. |
| Am J Obstet Gynecol 25MAR2021 | COVID-19 vaccine response in pregnant and lactating women: a cohort study | Gray K.J., et al. USA gotopaper         | Vaccines           | **Aim:** to evaluate the immunogenicity and reactogenicity of COVID-19 mRNA vaccination in pregnant and lactating women compared to: (1) non-pregnant controls and (2) natural COVID-19 infection during pregnancy.  
> 131 reproductive-age vaccine recipients (84 pregnant, 31 lactating, and 16 non-pregnant)  
**Findings:**  
> Vaccine-induced antibody titers were equivalent in pregnant and lactating compared to non-pregnant women (median [IQR] 5.59 [4.68-5.89] pregnant, 5.74 [5.06-6.22] lactating, 5.62 [4.77-5.98] non-pregnant, p = 0.24).  
> All titers were significantly higher than those induced by SARS-CoV-2 infection during pregnancy (p < 0.0001).  
> Vaccine-generated antibodies were present in all umbilical cord blood and breastmilk samples.  
> Neutralizing antibody titers were lower in umbilical cord compared to maternal sera, but it was not achieve statistically significant (median [IQR] 104.7 [61.2-188.2] maternal sera, 52.3 [11.7-69.6] cord sera, p=0.05).  
> The second vaccine dose (boost dose) increased SARS-CoV-2-specific IgG, but not IgA, in maternal blood and breastmilk.  
> No differences were noted in reactogenicity across the groups.  
**COVID-19 mRNA vaccines generated robust humoral immunity in pregnant and lactating women, significantly greater than the response to natural infection. Immune transfer to neonates occurred via placenta and breastmilk.** |
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> Over a mean follow-up of 140 days, nearly a third of individuals who were discharged from hospital after acute covid-19 were readmitted (14 060 of 47 780) and more than 1 in 10 (5875) died after discharge, with these events occurring at rates 4 and 8 times greater, respectively, than in the matched control group.  
> Rates of respiratory disease (P<0.001), diabetes (P<0.001), and cardiovascular disease (P<0.001) were also significantly raised in patients with covid-19, with 770 (95% CI 758-783), 127 (122-132), and 126 (121-131) diagnoses per 1000 person years, respectively.  
> Rate ratios were greater for individuals aged <70 than for those aged ≥70, and in ethnic minority groups compared with the white population. Largest differences was seen for respiratory disease (10.5 (95% CI 9.7-11.4) for age < 70 years v 4.6 (4.3 to 4.8) for age ≥70, and 11.4 (9.8-13.3) for non-white v 5.2 (5.0-5.5) for white individuals).  
Individuals discharged from hospital after covid-19 had increased rates of multiorgan dysfunction compared with the expected risk in the general population. The increase in risk was not confined to the elderly and was not uniform across ethnicities. |
| Lancet 30MAR2021 | Efficacy of ChAdOx1 nCoV-19 (AZD1222) vaccine against SARS-CoV-2 variant of concern 202012/01 (B.1.1.7): an exploratory analysis of a randomised controlled trial | Emary K.R.W., et al. UK | Vaccines | Background: A new variant of SARS-CoV-2, B.1.1.7, emerged as the dominant cause of COVID-19 disease in the UK from November, 2020. We report a post-hoc analysis of the efficacy of the adenoviral vector vaccine, ChAdOx1 nCoV-19 (AZD1222), against this variant.  
Methods:  
> Volunteers (aged ≥18 years), enrolled during the phase 2/3 vaccine efficacy studies in the UK receiving randomly ChAdOx1 nCoV-19 or a meningooccal conjugate control (MenACWY) vaccine  
> Upper airway swabs on a weekly basis and recording of COVID-19 disease symptoms if any  
> Swabs were tested by nucleic acid amplification test (NAAT) for SARS-CoV-2 and positive samples were sequenced  
> Assessment of neutralising antibody responses against the B.1.1.7 lineage and a canonical non-B.1.1.7 lineage (Victoria).  
Findings:  
> 8534 participants, 6636 (78%) aged 18–55 years and 5065 (59%) female.  
> 520 participants developed SARS-CoV-2 infection.  
> 1466 NAAT positive nose and throat swabs were collected from these participants during the trial.  
> Of these, 401 swabs from 311 participants were successfully sequenced.  
> Laboratory virus neutralisation activity by vaccine-induced antibodies was lower against the B.1.1.7 variant than against the Victoria lineage (geometric mean ratio 8·9, 95% CI 7·2–11·0).  
> Clinical vaccine efficacy against symptomatic NAAT positive infection was 70-4% (95% CI 43·6–84·5) for B.1.1.7 and 81-5% (67·9–89·4) for non-B.1.1.7 lineages.  
Conclusion: ChAdOx1 nCoV-19 showed reduced neutralisation activity against the B.1.1.7 variant compared with a non-B.1.1.7 variant in vitro, but the vaccine showed efficacy against the B.1.1.7 variant of SARS-CoV-2. |
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<tr>
<td><strong>Cell</strong> 30MAR2021</td>
<td>Antibody evasion by the P.1 strain of SARS-CoV-2</td>
<td>Dejnirattisai W. et al. UK</td>
<td>Virology</td>
<td>Examination of an isolate of P.1 variant cultured from a throat swab taken from an infected patient in Manaus, Brazil in December 2020 and comparison of its interactions with serum and antibodies with those of three other viruses, an early isolate, B.1.1.7 and B.1.351.</td>
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<td><strong>Nature Commun.</strong> 30MAR2021</td>
<td>Peginterferon Lambda-1a for treatment of outpatients with uncomplicated COVID-19: a randomized placebo-controlled trial</td>
<td>Jagannathan P., et al. USA</td>
<td>Therapeutics</td>
<td>Aim: to determine whether a single, 180 mcg subcutaneous dose of Peginterferon Lambda-1a (Lambda) within 72 hours of diagnosis could shorten the duration of viral shedding (primary endpoint) or symptoms (secondary endpoint)</td>
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**Findings:**

- Reduction in the neutralizing capacity of immune serum to P.1 similar to the reduction seen with B.1.1.7, but not as severe as that seen with B.1.351.
- Increased affinity of P.1 137 RBD for ACE2
- Investigation of the structural basis of this through crystallography.
- Neutralization by a panel of potent monoclonal antibodies which block RBD/ACE2 interaction: mAb 222, which contacts both K417 and N501, is resistant to the 141 501Y and 417T/N mutations found in the P.1/B1.351 strains.
- Dissection of the basis for this via a series of high resolution structures of RBD-Fab complexes and based on this restoration of neutralization of certain antibodies by swapping the light chain.

**Conclusion:**
P.1 can escape neutralization by a number of monoclonal antibodies including some being developed for prophylactic or therapeutic use, while other antibodies with epitopes away from the mutated RBD residues retain broad neutralization.

**Aim:**

- to determine whether a single, 180 mcg subcutaneous dose of Peginterferon Lambda-1a (Lambda) within 72 hours of diagnosis could shorten the duration of viral shedding (primary endpoint) or symptoms (secondary endpoint) |

**Methods:**

- randomized, single-blind, placebo-controlled trial in 120 outpatients with mild to moderate COVID-19, of whom 110 (91.7%) completed 28 days of follow up. Participants were recruited within 72 h of diagnosis.

**Findings:**

- 60 patients receiving Lambda and 60 receiving placebo, the median time to cessation of viral shedding was 7 days (hazard ratio [HR] = 0.81; 95% confidence interval [CI] 0.56 to 1.19).
- Symptoms resolved in 8 and 9 days in Lambda and placebo, respectively, and symptom duration did not differ significantly between groups (HR 0.94; 95% CI 0.64 to 1.39).
- Both Lambda and placebo were well-tolerated, though liver transaminase elevations were more common in the Lambda vs. placebo arm (15/60 vs 5/60; p = 0.027).

A single dose of subcutaneous Peginterferon Lambda-1a neither shortened the duration of SARS-CoV-2 viral shedding nor improved symptoms in outpatients with uncomplicated COVID-19.
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<tr>
<td>Nature 29MAR2021</td>
<td>Escape of SARS-CoV-2 501Y.V2 from neutralization by convalescent plasma</td>
<td>Cele S., et al. International</td>
<td>Therapeutics - Variants</td>
<td>Live virus neutralization assay to compare neutralization of a non-VOC variant versus the 501Y.V2 variant using plasma collected from adults hospitalized with COVID-19 from two South African infection waves, with the second wave dominated by 501Y.V2 infections. Findings: &gt; Sequencing demonstrated that infections in first wave plasma donors were with viruses harbouring none of the 501Y.V2-defining mutations, except for one with the E484K mutation in the receptor binding domain. &gt; 501Y.V2 virus was effectively neutralized by plasma from second wave infections and first wave virus was effectively neutralized by first wave plasma. &gt; In cross-neutralization, 501Y.V2 virus was poorly neutralized by first wave plasma, with a 15.1-fold drop relative to 501Y.V2 neutralization by second wave plasma across participants. &gt; Second wave plasma cross-neutralization of first wave virus was more effective, showing only a 2.3-fold decline relative to first wave plasma neutralization of first wave virus. Conclusion: Effective neutralization of first wave virus by 501Y.V2 infection elicited plasma provides preliminary evidence that vaccines based on VOC sequences could retain activity against other circulating SARS-CoV-2 lineages.</td>
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<td>Nature Commun. 29MAR2021</td>
<td>A haemagglutination test for rapid detection of antibodies to SARS-CoV-2</td>
<td>Townsend A., et al. USA</td>
<td>Diagnostics</td>
<td>Aim: to describe a quantitative Haemagglutination test (HAT) for the detection of antibodies to the receptor binding domain of the SARS-CoV-2 spike protein. Methods: simple HA test for the detection of Abs to the receptor binding domain (RBD) of the SARS-CoV-2 spike protein. In order to link the SARS-CoV-2 RBD to red cells, they selected the single domain antibody (nanobody) IH46, specific for a conserved epitope on glycophorin A, via a short (GSG)2 linker to produce the fusion protein IH4-RBD-6H. Findings: &gt; HAT functions as a viable test for the presence of antibodies to the RBD of the SARS-CoV-2 spike protein in stored serum/plasma samples, using O−ve red cells as indicators &gt; The HAT has a sensitivity of 90% and specificity of 99% for detection of antibodies after a PCR diagnosed infection.</td>
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<td>Clin Infect Dis. 27MAR2021</td>
<td>Assessing asymptomatic, pre-symptomatic and symptomatic transmission risk of SARS-CoV-2</td>
<td>Wu P., et al. China</td>
<td>Public Health / Epidemiology</td>
<td>Methods: &gt; Detailed information on transmission events and symptom status based on laboratory-confirmed patient data and contact tracing data from four provinces and one municipality in China &gt; Estimated the variation in risk of transmission over time, and the severity of secondary infections, by symptomatic status of the infector. Findings: &gt; 393 symptomatic index cases with 3136 close contacts and 185 asymptomatic index cases with 1078 close contacts included into the study &gt; The secondary attack rate among close contacts of symptomatic and asymptomatic index cases were 4.1% (128/3136) and 1.1% (12/1078), respectively, corresponding to a higher transmission risk from symptomatic cases than from asymptomatic cases (OR: 3.79, 95% CI: 2.06, 6.95) &gt; Approximately 25% (32/128) and 50% (6/12) of the infected close contacts were asymptomatic from symptomatic and asymptomatic index cases &gt; Pre-symptomatic transmission of COVID-19 accounted for 38% of all infections occurred from exposure to symptomatic cases. &gt; Infected contacts of asymptomatic index cases were more likely to be asymptomatic and less likely to be severe. Asymptomatic and pre-symptomatic transmission play an important role in spreading infection, although asymptomatic cases pose a lower risk of transmission than symptomatic cases. Early case detection and effective test-and-trace measures are important to reduce transmission.</td>
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| Nature Commun. 26MAR2021 | N-protein presents early in blood, dried blood and saliva during asymptomatic and symptomatic SARS-CoV-2 infection | Shan D., et al. Germany/USA [gotopaper](#) | Diagnostics | Aim: to describe the development of a SARS-CoV-2 antigen test using Simoa technology to quantify N-protein in serum/plasma, dried blood microsamples (DBS), and saliva. 
Methods: SARS-CoV-2 N-protein and anti-SARS-CoV-2 spike IgG were quantified directly in serum and plasma from venous collection, capillary blood acquired by finger-stick DBS devices (DBS), and saliva samples. 
Findings: 
> Compared to molecular testing, >90% PPA of SARS-CoV-2-positive patients and >98% negative percent agreement (NPA) were observed in all matrices within 7 days of positive PCR test, both for asymptomatic and symptomatic patients. 
> N-protein load decreases as anti-SARS-CoV-2 spike-IgG increases, and N-protein levels correlate with RT-PCR Ct-values in saliva, and between matched saliva and capillary blood samples. 
> N-protein levels in saliva are higher but more variable than levels in capillary blood. 
The Simoa N-protein antigen test represents a robust SARS-CoV-2 detection tool, effectively detecting SARS-CoV-2 infection via antigen levels in blood or saliva, using non-invasive, swab-independent collection methods, with potential at home/point of care sampling. |
Methods: 
> Examined sensitivity of the two variants to SARS-CoV-2 antibodies present in sera and nasal swabs from individuals infected with previously circulating strains or who were recently vaccinated, in comparison with a D614G reference virus. 
> New rapid neutralization assay, based on reporter cells that become positive for GFP after overnight infection. 
Results: 
> Sera from 58 convalescent individuals collected up to 9 months after symptoms, similarly neutralized B.1.1.7 and D614G. In contrast, after 9 months, convalescent sera had a mean 6-fold reduction in neutralizing titers, and 40% of samples lacked any activity against B.1.351. 
> Sera from 19 individuals vaccinated twice with Pfizer Cominarty, up to 6 weeks after vaccination, were similarly potent against B.1.1.7 but less efficacious against B.1.351, when compared to D614G. 
> Neutralizing titers increased after the second vaccine dose, but were 14-fold lower against B.1.351. Sera from convalescent or vaccinated individuals similarly bound the three spike proteins in a flow cytometry-based serological assay. 
Neutralizing antibodies were rarely detected in nasal swabs from vaccinees. Faster-spreading SARS-CoV-2 variants acquired a partial resistance to neutralizing antibodies generated by natural infection or vaccination, most frequently detected in individuals with low antibody levels. Our results indicate that B1.351, but not B.1.1.7, may increase the risk of infection in immunized individuals. |
> Changes in VOC frequency inferred from genetic data correspond closely to changes inferred by S-gene target failures (SGTF) in community-based diagnostic PCR testing. 
> Analysis of trends in SGTF and non-SGTF case numbers in local areas across England shows that the VOC has higher transmissibility than non-VOC lineages, even if the VOC has a different latent period or generation time. 
> The SGTF data indicate a transient shift in the age composition of reported cases, with a larger share of under 20 year olds among reported VOC than non-VOC cases. 
> Time-varying reproduction numbers for the VOC and cocirculating lineages were estimated using SGTF and genomic data. The best supported models did not indicate a substantial difference in VOC transmissibility among different age groups. 
> There is a consensus among all analyses that the VOC has a substantial transmission advantage with a 50% to 100% higher reproduction number. |
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| Lancet Infect Dis. 25MAR2021 | Performance and operational feasibility of antigen and antibody rapid diagnostic tests for COVID-19 in symptomatic and asymptomatic patients in Cameroon: a clinical, prospective, diagnostic accuracy study | Boum Y., et al. Cameroon gotopaper | Diagnostics | **Aim:** to assess the performance of four antibody-based rapid diagnostic tests and one antigen-based rapid diagnostic test for detecting SARS-CoV-2 infection in the community in Cameroon.  
**Methods:** prospective, diagnostic accuracy study with 1195 individuals aged at least 21 years who were either symptomatic and suspected of having COVID-19 or asymptomatic and presented for screening. Peripheral blood for SARS-CoV-2 antibodies were tested using the Innovita, Wondfo, SD Biosensor, and Runkun tests, and nasopharyngeal swabs for SARS-CoV-2 antigen using the SD Biosensor test. Antigen rapid diagnostic tests were compared with Abbott PCR testing, and antibody rapid diagnostic tests were compared with Biomerieux immunoassays. Two diagnostic algorithms that incorporated rapid diagnostic tests for symptomatic and asymptomatic patients using simulation modelling were tested.  
**Findings:** > 347 patients (29%) tested SARS-CoV-2 PCR-positive, 223 (19%) rapid diagnostic test antigen-positive, and 478 (40%) rapid diagnostic test antibody-positive. > Antigen-based rapid diagnostic test sensitivity was 80.0% in the first 7 days after symptom onset, but Antibody-based rapid diagnostic tests had only 26.8% sensitivity. > Antibody rapid diagnostic test sensitivity increased to 76.4% 14 days after symptom onset. > Among asymptomatic participants, the sensitivity of antigen-based and antibody-based rapid diagnostic tests were 37.0% and 50.7%, respectively. > An antigen-based retrospective algorithm applied to symptomatic patients showed 94.0% sensitivity and 91.0% specificity in the first 7 days after symptom onset. > For asymptomatic participants, the algorithm showed a sensitivity of 34% and a specificity of 92.0%.  
**Rapid diagnostic tests had good overall sensitivity for diagnosing SARS-CoV-2 infection. Rapid diagnostic tests could be incorporated into efficient testing algorithms as an alternative to PCR to decrease diagnostic delays and onward viral transmission.** |
| JAMA Netw Open 24MAR2021 | Comparison of Time to Clinical Improvement With vs Without Remdesivir Treatment in Hospitalized Patients With COVID-19 | Garibaldi B.T., et al. USA gotopaper | Therapeutics | **Aim:** to examine whether remdesivir administered with or without corticosteroids for treatment of COVID-19 is associated with more rapid clinical improvement in a racially/ethnically diverse population.  
**Exposures:** No Remdesivir, Remdesivir treatment with or without corticosteroid administration.  
**Primary outcome:** rate of clinical improvement (hospital discharge or decrease of 2 points on the World Health Organization severity score)  
**Secondary outcome:** mortality at 28 days; Clinical improvement and time to death associated with combined remdesivir and corticosteroid treatment.  
> Of 2483 consecutive admissions, 342 individuals received remdesivir, 184 of whom also received corticosteroids. Remdesivir patients were matched with admitted patients who did not receive Remdesivir.  
> For these 342 patients: median age was 60 years (46-69), 55.3% were men, 80.7% self-identified as non-White race/ethnicity.  
> Remdesivir recipients had a **shorter time to clinical improvement** than matched controls without remdesivir treatment (median, 5.0 days [4.0-8.0] vs 7.0 days [4.0-10.0]; adjusted hazard ratio (HR), 1.47 [95% CI, 1.22-1.79]).  
> Remdesivir recipients had a **28-day mortality rate of 7.7% compared with 14.0%** among matched controls, but this difference was **not statistically significant in the time-to-death analysis** (adjusted HR, 0.70; 95% CI, 0.38-1.28).  
> The addition of corticosteroids to remdesivir was **not associated with a reduced hazard of death** at 28 days (adjusted HR, 1.94; 95% CI, 0.67-5.57).  
In this study of adults hospitalized with COVID-19, receipt of remdesivir was associated with faster clinical improvement. Remdesivir plus corticosteroid administration did not reduce the time to death compared with remdesivir administered alone. |
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| Lancet 23MAR2021 | Dynamics of SARS-CoV-2 neutralising antibody responses and duration of immunity: a longitudinal study | Wan N.C., et al. Singapore [gotopaper](#) | Immunology | **Aim:** to investigate the peak levels and dynamics of neutralising antibody waning and IgG avidity maturation over time, and correlate this with clinical parameters, cytokines, and T-cell responses.  
**Methods:** longitudinal study of patients who had recovered from COVID-19 up to day 180 post-symptom onset by monitoring changes in neutralising antibody levels using a previously validated surrogate virus neutralisation test.  
**Findings:** Five distinctive patterns of neutralising antibody dynamics were identified as follows:  
- Negative: individuals who did not, at our intervals of sampling, develop neutralising antibodies at the 30% inhibition level (19 [12%] of 164 patients).  
- Rapid waning: individuals who had varying levels of neutralising antibodies from around 20 days after symptom onset, but seroreverted in less than 180 days (44 [27%] of 164 patients).  
- Slow waning: individuals who remained neutralising antibody-positive at 180 days post-symptom onset (52 [29%] of 164 patients).  
- Persistent: although with varying peak neutralising antibody levels, these individuals had minimal neutralising antibody decay (52 [32%] of 164 patients).  
- Delayed response, a small group that showed an unexpected increase of neutralising antibodies during late convalescence (at 90 or 180 days after symptom onset; three [2%] of 164 patients).  
Neutralising antibody response dynamics in patients who have recovered from COVID-19 vary greatly, and prediction of immune longevity can only be accurately determined at the individual level. |
| JAMA Netw Open 22MAR2021 | Association of Age With SARS-CoV-2 Antibody Response | Yang H.S., et al. USA [gotopaper](#) | Immunology | **Aim:** To investigate the association of age with the quantity and quality of SARS-CoV-2 antibody responses.  
**Methods:** Cross-sectional study evaluating 31,426 SARS-CoV-2 antibody tests from pediatric and adult patients. Data were collected from a New York City hospital from April 9 to August 31, 2020.  
**Findings:** Among 31,426 antibody test results, the seroprevalence in the pediatric (197 [16.5%; 95% CI, 14.4%-18.7%]) and adult (5630 [18.6%; 95% CI, 18.2%-19.1%]) patient populations was similar.  
The SARS-CoV-2 IgG level showed a negative correlation with age in the pediatric population (r = −0.45, P < .001) and a moderate but positive correlation with age in adults (r = 0.24, P < .001).  
Patients aged 19 to 30 years exhibited the lowest IgG levels (eg, aged 25-30 years vs 1-10 years: 99 [44-180] relative fluorescence units [RFU] vs 443 [188-851] RFU).  
Children exhibited higher median (IQR) IgG levels, TAB levels, and SNAB activity compared with adolescents (eg, IgG levels: 473 RFU vs 191 RFU; P < .001) and young adults (eg, IgG levels: 473 RFU vs 85 RFU; P < .001).  
Children had higher antibody binding avidity compared with young adults, but the difference was not significant.  
**This study suggests that SARS-CoV-2 viral specific antibody response profiles are distinct in different age groups. Age-targeted strategies for disease screening and management as well as vaccine development may be warranted.** |
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<tr>
<td>Nature Commun. 22MAR2021</td>
<td>SARS-CoV-2 infection induces sustained humoral immune responses in convalescent patients following symptomatic COVID-19</td>
<td>Wu J., et al. China gotopaper</td>
<td>Immunology</td>
<td>Aim: to quantify immunoglobulin M (IgM) and G (IgG) antibodies recognizing the SARS-CoV-2 receptor-binding domain (RBD) of the spike (S) or the nucleocapsid (N) protein, and neutralizing antibodies during a period of 6 months from disease onset in 349 symptomatic COVID-19 patients. &gt; The positivity rate and magnitude of IgM-S and IgG-N responses increase rapidly. &gt; High levels of IgM-S/N and IgG-S/N at 2-3 weeks after disease onset are associated with virus control and IgG-S titers correlate closely with the capacity to neutralize SARS-CoV-2. &gt; Although specific IgM-S/N can become undetectable 12 weeks after disease onset in most patients, IgG-S/N titers have an intermediate contraction phase, but stabilize at relatively high levels over the 6 month observation period. &gt; At late time points, the positivity rates for binding and neutralizing SARS-CoV-2-specific antibodies are still &gt;70%. These data indicate sustained humoral immunity in recovered patients who had symptomatic COVID-19, suggesting prolonged immunity.</td>
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<td>Cell 20MAR2021</td>
<td>SARS-CoV-2 variants B.1.351 and P.1 escape from neutralizing antibodies</td>
<td>Hoffmann M., et al. Germany gotopaper</td>
<td>Viral variants</td>
<td>Aim: to test sensitivity of SARS-CoV-2 variants B.1.1.7 (UK), B.1.351 (South Africa) and P.1 (Brazil) to cell entry inhibitors and antibodies, by using pseudoparticles. &gt; B.1.1.7, B.1.351 and P.1 do not show augmented host cell entry. &gt; Entry of all variants into human cells is susceptible to blockade by the entry inhibitors soluble ACE2, Camostat, EK-1 and EK-1-C4. &gt; Entry of the B.1.351 and P.1 variant is partially (Casirivimab) or fully (Bamlanivimab) resistant to antibodies used for COVID-19 treatment. &gt; Entry of these variants was less efficiently inhibited by plasma from convalescent COVID-19 patients and sera from BNT162b2 vaccinated individuals. These results suggest that SARS-CoV-2 may escape neutralizing antibody responses.</td>
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<td>Lancet 20MAR2021</td>
<td>Seroprevalence and humoral immune durability of anti-SARS-CoV-2 antibodies in Wuhan, China: a longitudinal, population-level, cross-sectional study</td>
<td>He Z., et al. China gotopaper</td>
<td>Immunology</td>
<td>Seroprevalence and kinetics of anti-SARS-CoV-2 antibodies at population level in Wuhan to inform the development of vaccination strategies. Methods Longitudinal cross-sectional study, population-stratified, cluster random sampling method (100 communities from the 13 districts of Wuhan). Households systematically selected. A venous blood sample taken for immunological testing (pan-immunoglobulins, IgM, IgA, and IgG antibodies against SARS-CoV-2 nucleocapsid protein and neutralising antibodies). Findings &gt; 9542 individuals from 3556 families had sampled for analyses. &gt; 532 participants were positive for pan-immunoglobulins against SARS-CoV-2 (baseline seroprevalence of 6.92%). &gt; 437 of 532 (82.1%) participants who were positive for pan-immunoglobulins were asymptomatic. &gt; 69 (13-0%) of 532 individuals were positive for IgM antibodies, 84 (15-8%) were positive for IgA antibodies, 532 (100%) were positive for IgG antibodies, and 212 (39-8%) were positive for neutralising antibodies at baseline. &gt; On the basis of data from 335 individuals who attended all three follow-up visits and who were positive for pan-immunoglobulins, neutralising antibody levels did not significantly decrease over the study period &gt; Neutralising antibody titres were lower in asymptomatic individuals than in confirmed cases and symptomatic individuals. &gt; Although titres of IgG decreased over time, the proportion of individuals who had IgG antibodies did not decrease substantially Conclusion 6.92% of a cross-sectional sample of the population of Wuhan developed antibodies against SARS-CoV-2, with 39-8% of this population seroconverting to have neutralising antibodies.</td>
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<td>Nature Commun. 19MAR2021</td>
<td>Favipiravir antiviral efficacy against SARS-CoV-2 in a hamster model</td>
<td>Driouich J.S., et al. France <a href="#">gotopaper</a></td>
<td>Therapeutics</td>
<td>Asessment of antiviral efficacy of favipiravir (Syrian hamster model)  &lt;br&gt; <strong>Findings:</strong>&lt;br&gt; &gt; In vitro efficacy of favipiravir  &lt;br&gt; - Vero E6 cells: Infectious titer reductions (fold change in comparison with untreated cells) ≥ 2 with 125 µM of favipiravir and between 11 and 342 with 500 µM.  &lt;br&gt; - Caco-2 cells (no CPE with SARS-CoV-2 BavPat1 strain) infectious titer reductions around 5 with 125 µM of favipiravir and between 144 and 7721 with 500 µM.  &lt;br&gt; &gt; In vivo efficacy of favipiravir  &lt;br&gt; - intranasally infection of Syrian hamsters with different inoculums, receiving favipiravir at the day of infection up to 2 dpi. Doses of favipiravir: 18.75, 37.5, and 75 mg/day. Effect of favipiravir in reducing infectious titers is dose dependent, in particular when low virus inocula were used to infect animal. Significant differences in virus replication in clarified lung homogenates between treated and untreated animals.  &lt;br&gt; - Antiviral effect of favipiravir correlates with incorporation of a large number of mutations into viral genomes and decrease of viral infectivity.  &lt;br&gt; - Antiviral efficacy is achieved with plasma drug exposure comparable with those previously found during human clinical trials (the highest dose of favipiravir tested is associated with signs of toxicity in animals)  &lt;br&gt; Pharmacokinetic and tolerance studies are required to determine whether similar effects can be safely achieved in humans.  &lt;br&gt; <strong>Conclusion:</strong>&lt;br&gt; High doses of favipiravir are associated with antiviral activity against SARS-CoV-2 infection in a hamster model. The better antiviral efficacy was observed using a preventive strategy, suggesting that favipiravir could be more appropriate for a prophylactic use.</td>
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<td>Nature Commun. 19MAR2021</td>
<td>Exposure to SARS-CoV-2 generates T-cell memory in the absence of a detectable viral infection</td>
<td>Wang Z., et al. China <a href="#">gotopaper</a></td>
<td>Immunology</td>
<td>Aim: to test SARS-CoV-2-specific T-cell immunity in virus-exposed individuals.  &lt;br&gt; COVID-19 patients: NAT+, hospitalised and recovered, samples taken 48–86 days after disease onset;  &lt;br&gt; Asymptomatic patients: NAT+, with no signs of symptoms  &lt;br&gt; Close contacts: NAT−, no SARS-CoV-2 specific antibodies, in contact with patients between 5 days before disease onset and hospitalisation.  &lt;br&gt; &gt; Virus-specific CD4+ and CD8+ T-cell memory was observed in recovered COVID-19 patients (in 94.44% and 88.33% of patients, respectively) and close contacts (in 57.97% and 14.49%, respectively).  &lt;br&gt; &gt; The size and quality of the memory T-cell pool of COVID-19 patients are larger and better than those of close contacts.  &lt;br&gt; &gt; However, the proliferation capacity, size and quality of T-cell responses in close contacts are readily distinguishable from healthy donors, suggesting close contacts are able to gain T-cell immunity against SARS-CoV-2 despite lacking a detectable infection.  &lt;br&gt; &gt; Asymptomatic and symptomatic COVID-19 patients contain similar levels and qualities of SARS-CoV-2-specific T-cells.  &lt;br&gt; &gt; CD4+ T memory and CD8+ T memory may have contracted to a stable plateau 48–86 days after symptom onset.  &lt;br&gt; &gt; Virus-specific memory CD4+ T cell pool correlated with the titers of IgG against the S RBD region and the N protein, whereas no apparent correlation between CD8+ T cells and IgG titers was observed.  &lt;br&gt; This study demonstrates the versatility and potential of memory T cells from COVID-19 patients and close contacts, which may be important for host protection.</td>
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<td>JAMA Netw Open 19MAR2021</td>
<td>Association of Vitamin D Levels, Race/Ethnicity, and Clinical Characteristics With COVID-19 Test Results</td>
<td>Meltzer DO., et al. USA gotopaper</td>
<td>Public Health / Epidemiology</td>
<td>Aim: To examine whether COVID-19 test results are associated with differences in vitamin D levels of 30 ng/mL or greater, including for White individuals and for Black individuals.</td>
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Findings:

> Population based cohort study: two cohorts of adults (≥18 yrs) registered at a general practice (1 Feb - 1 Sept 2020)
> Adjusted hazard ratios (HR) for SARS-CoV-2 infection, covid-19 related admission to hospital or intensive care, or death from covid-19, by presence of children in the household.

> Among 9,334,392 adults aged ≤65 yrs, during wave 1, living with children was not associated with materially increased risks of recorded SARS-CoV-2 infection, covid-19 related hospital admission or intensive care, or death from covid-19.
> In wave 2, among adults aged ≤65 yrs, living with children of any age was associated with an increased risk of recorded SARS-CoV-2 infection (HR 1.06 (95% CI 1.05 to 1.08) for living with children aged 0-11 years; 1.22 (1.20 to 1.24) for living with children aged 12-18 years) and covid-19 related hospital admission (1.18 (1.06 to 1.31) for living with children aged 0-11; 1.26 (1.12 to 1.40) for living with children aged 12-18).

Living with children aged 0-11:
> was associated with reduced risk of death from both covid-19 and non-covid-19 causes in both waves; living with children of any age was also associated with lower risk of dying from non-covid-19 causes.
> For adults ≤65 yrs during wave 2, was associated with an increased absolute risk of having SARS-CoV-2 infection recorded of 40-60 per 10,000 people, from 810 to between 850 and 870, and an increase in hospital admissions of 1-5 per 10,000 people, from 160 to between 161 and 165.

Living with children aged 12-18 yrs was associated with an increase of 160-190 per 10,000 in the number of SARS-CoV-2 infections and an increase of 2.6 per 10,000 in the number of hospital admissions.

In contrast to wave 1, evidence existed of increased risk of reported SARS-CoV-2 infection and covid-19 outcomes among adults living with children during wave 2. However, this did not translate into a materially increased risk of covid-19 mortality, and absolute increases in risk were small.
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<tr>
<td>Lancet 17MAR2021</td>
<td>Assessment of protection against reinfection with SARS-CoV-2 among 4 million PCR-tested individuals in Denmark in 2020: a population-level observational study</td>
<td>Hansen CH., et al. Denmark</td>
<td>Public Health / Epidemiology</td>
<td>Using national PCR-test data from 2020 (4 million individuals (69% of the population) underwent 10·6 million tests), we estimated protection towards repeated infection with SARS-CoV-2.</td>
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**Methods**

> Analysis of infection rates during the second surge of the COVID-19 epidemic (Sept 1 - Dec 31, 2020), by comparing infection rates between individuals with positive and negative PCR tests during the first surge (March - May, 2020)
>
> Alternative cohort analysis, comparing infection rates throughout the year between those with and without a previous confirmed infection at least 3 months earlier, irrespective of date.

**Findings**

> During the first surge (before June, 2020), 533381 people were tested, of whom 11727 (2·20%) were PCR positive, and 525339 were eligible for follow-up in the second surge, of whom 11068 (2·11%) had tested positive during the first surge.
>
> Among eligible PCR-positive individuals from the first surge of the epidemic, 72 (0·65% [95% CI 0·51–0·82]) tested positive again during the second surge compared with 16819 (3·27% [3·22–3·32]) of 514271 who tested negative during the first surge.
>
> Protection against repeat infection was 80·5% (95% CI 75·4–84·5).
>
> In the alternative cohort analysis, among those aged ≥65, observed protection against repeated infection was 47·1% (95% CI 24·7–62·8).
>
> No difference in estimated protection against repeated infection by sex (male 78·4% [72·1–83·2] vs female 79·1% [73·9–83·3]) or evidence of waning protection over time (3–6 months of follow-up 79·3% [74·4–83·3] vs ≥7 months of follow-up 77·7% [70·9–82·9]).

*These findings could inform decisions on groups to vaccinate and advocate for vaccination of previously infected individuals, as natural protection, especially among older people, cannot be relied on.*
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| JAMA 17MAR2021   | Four-Month Clinical Status of a Cohort of Patients After Hospitalization for COVID-19 | COMEBAC Study Group France | Public Health / Epidemiology - Long Covid | **Aim:** to describe the consequences at 4 months in patients hospitalized for COVID-19.  
**Findings**  
> 478 were evaluated by telephone (mean age, 61 years [SD, 16 years]; 201 men, 277 women).  
> 244 patients (51%) declared at least 1 symptom that did not exist before COVID-19: fatigue in 31%, cognitive symptoms in 21%, and new-onset dyspnea in 16%. There was further evaluation in 177 patients (37%), including 97 of 142 former ICU patients.  
> The median 20-item Multidimensional Fatigue Inventory score (n = 130) was 4.5 (interquartile range IR, 3.0-5.0) for reduced motivation and 3.7 (IR, 3.0-4.5) for mental fatigue (possible range, 1 [best] to 5 [worst]).  
> The median 36-item Short-Form Health Survey score (n = 145) was 25 (IR, 25.0-75.0) for the subscale “role limited owing to physical problems” (possible range, 0 [best] to 100 [worst]).  
> Computed tomographic lung-scan abnormalities were found in 108 of 171 patients (63%), mainly subtle ground-glass opacities. Fibrotic lesions were observed in 33 of 171 patients (19%), involving less than 25% of parenchyma in all but 1 patient. Fibrotic lesions were observed in 19 of 49 survivors (39%) with acute respiratory distress syndrome.  
> Among 94 former ICU patients, anxiety, depression, and posttraumatic symptoms were observed in 23%, 18%, and 7%, respectively.  
> The left ventricular ejection fraction was less than 50% in 8 of 83 ICU patients (10%). New-onset chronic kidney disease was observed in 2 ICU patients.  
> Serology was positive in 172 of 177 outpatients (97%).  
**Four months after hospitalization for COVID-19, a cohort of patients frequently reported symptoms not previously present, and lung-scan abnormalities were common among those who were tested.** |
| Nature 16MAR2021 | Clofazimine broadly inhibits coronaviruses including SARS-CoV-2 | Yuan S., et al. China | Therapeutics | Clofazimine is an anti-leprosy drug with a favourable safety profile  
**In vitro & in vivo studies**  
> We show that clofazimine possesses pan-coronaviral inhibitory activity, and can antagonize SARS-CoV-2 and MERS-CoV replication in multiple in vitro systems.  
> The FDA-approved molecule was found to inhibit viral spike-mediated cell fusion and viral helicase activity.  
> In a hamster model of SARS-CoV-2 pathogenesis, prophylactic or therapeutic administration of clofazimine significantly reduced viral load in the lung and faecal viral shedding, and also mitigated inflammation associated with viral infection.  
> Combinatorial application of clofazimine and remdesivir exhibited antiviral synergy in vitro and in vivo, and restricted upper respiratory tract viral shedding.  
**Since clofazimine is orally bioavailable and has a comparatively low manufacturing cost, it is an attractive clinical candidate for outpatient treatment and remdesivir-based combinatorial therapy for hospitalized COVID-19 patients, particularly in developing countries. Taken together, our data provide evidence that clofazimine may have a role in the control of the current pandemic SARS-CoV-2, and, possibly most importantly, emerging CoVs of the future.** |
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<td>NEJM 16MAR2021</td>
<td>Efficacy of the ChAdOx1 nCoV-19 Covid-19 Vaccine against the B.1.351 Variant</td>
<td>Madhi S.A., et al. International gotopaper</td>
<td>Vaccines - Variants</td>
<td>Efficacy of ChAdOx1 against emerging SARS-CoV-2 variants of concern, including the B.1.351 (S01Y.V2) variant first identified in South Africa. <strong>Methods:</strong> &gt; Multicenter, double-blind, randomized, controlled trial in HIV- in South Africa. &gt; Participants age: 18 to 65 years of age &gt; Two doses of vaccine containing 5×10^10 viral particles or placebo (0.9% sodium chloride solution) 21 to 35 days apart. &gt; Serum samples obtained from 25 participants after the second dose were tested by pseudovirus and live-virus neutralization assays against the original D614G virus and the B.1.351 variant. <strong>Primary end points:</strong> safety and efficacy of the vaccine against laboratory-confirmed symptomatic coronavirus 2019 illness (Covid-19) more than 14 days after the second dose. <strong>Findings:</strong> &gt; 2026 HIV-negative adults enrolled (median age, 30 years); &gt; 1010 and 1011 participants received at least one dose of placebo or vaccine, respectively. &gt; Both the pseudovirus and the live-virus neutralization assays showed greater resistance to the B.1.351 variant in serum samples obtained from vaccine recipients than in samples from placebo recipients. &gt; In the primary end-point analysis, mild-to-moderate Covid-19 developed in 23 of 717 placebo recipients (3.2%) and in 19 of 750 vaccine recipients (2.5%), for an efficacy of 21.9% (95% confidence interval [CI], −49.9 to 59.8). &gt; Among the 42 participants with Covid-19, 39 cases (92.9%) were caused by the B.1.351 variant; vaccine efficacy against this variant, analyzed as a secondary end point, was 10.4% (95% CI, −76.8 to 54.8). &gt; The incidence of serious adverse events was balanced between the vaccine and placebo groups. <strong>Conclusion:</strong> A two-dose regimen of the ChAdOx1 nCoV-19 vaccine did not show protection against mild-to-moderate Covid-19 due to the B.1.351 variant.</td>
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<td>Cell Rep. 16MAR2021</td>
<td>Virological and immunological features of SARS-CoV-2-infected children who develop neutralizing antibodies</td>
<td>Cotugno N., et al. Italy gotopaper</td>
<td>Immunology</td>
<td>Aim: to define the humoral and cellular responses in SARS-CoV-2-infected children. <strong>Methods:</strong> Analysis of anti-SARS-CoV-2 antibodies and their neutralizing activity (PRNT) in 66 COVID-19-infected children at 7 (±2) days after symptom onset. Analysis of Ag-specific T and B cells defined as CD4+CD40L+ and SARS-CoV-2 Spike (S1+S2)-positive switched B cells. <strong>Findings:</strong> &gt; Individuals with specific humoral responses presented faster virus clearance and lower viral load associated with a reduced in vitro infectivity. &gt; The frequencies of SARS-CoV-2-specific CD4+CD40L+ T cells and Spike-specific B cells were associated with the anti-SARS-CoV-2 antibodies and the magnitude of neutralizing activity. &gt; The plasma proteome confirmed the association between cellular and humoral SARS-CoV-2 immunity, and PRNT+ patients show higher viral signal transduction molecules (SLAMF1, CD244, CLEC4G). <strong>Cellular and humoral anti-SARS-CoV-2 responses in children, which may drive future vaccination trial end points and quarantine measures policies.</strong></td>
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<td>Clin Infect Dis.</td>
<td>Reinfection Rates among Patients who Previously Tested Positive for COVID-19: a Retrospective Cohort Study</td>
<td>Sheehan M.M., et al. USA gotopaper</td>
<td>Public Health / Epidemiology</td>
<td>Methods&lt;br&gt;&amp; Retrospective cohort study of one multi-hospital health system included 150,325 patients tested for COVID-19 infection via PCR from March 12, 2020 to August 30, 2020&lt;br&gt;&amp; Testing performed up to February 24, 2021 in these patients was included for analysis&lt;br&gt;&amp; Main outcome = reinfection (defined as infection ≥ 90 days after initial testing)&lt;br&gt;&lt;br&gt;Findings&lt;br&gt;&amp; Protection offered from prior infection was 81.8% (95% confidence interval 76.6 to 85.8), and against symptomatic infection was 84.5% (95% confidence interval 77.9 to 89.1)&lt;br&gt;&amp; Prior infection in patients with COVID-19 was highly protective against reinfection and symptomatic disease.&lt;br&gt;&amp; This protection increased over time, suggesting that viral shedding or ongoing immune response may persist beyond 90 days and may not represent true reinfection.&lt;br&gt;&amp; As vaccine supply is limited, patients with known history of COVID-19 could delay early vaccination to allow for the most vulnerable to access the vaccine and slow transmission.</td>
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<td>Nature</td>
<td>Increased mortality in community-tested cases of SARS-CoV-2 lineage B.1.1.7</td>
<td>Davies N.G., et al. UK gotopaper</td>
<td>Variants</td>
<td>Aim: to determine if variant B.1.1.7 leads to changes in disease severity by analysing a dataset linking 2,245,263 positive SARS-CoV-2 community tests and 17,452 COVID-19 deaths in England (1 Sept 2020 - 14 Feb 2021).&lt;br&gt;&amp; For 1,146,534 (51%) of these tests, the presence or absence of B.1.1.7 can be identified because of mutations in this lineage preventing PCR amplification of the spike gene target (S gene target failure, SGTF).&lt;br&gt;&amp; Based on 4,945 deaths with known SGTF status, we estimate that the hazard of death associated with SGTF is 55% (95% CI 39–72%) higher after adjustment for age, sex, ethnicity, deprivation, care home residence, local authority of residence and test date.&lt;br&gt;&amp; These data correspond to the absolute risk of death for a 55–69-year-old male increasing from 0.6% to 0.9% (95% CI 0.8–1.0%) within 28 days after a positive test in the community.&lt;br&gt;&amp; Correcting for misclassification of SGTF and missingness in SGTF status, we estimate a 61% (42–82%) higher hazard of death associated with B.1.1.7. This analysis suggests that B.1.1.7 is not only more transmissible than preexisting SARS-CoV-2 variants, but may also cause more severe illness.</td>
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<td>Nature Commun.</td>
<td>Evaluating the impact of curfews and other measures on SARS-CoV-2 transmission in French Guiana</td>
<td>Andronico A., et al. French Guiana gotopaper</td>
<td>Public Health / Epidemiology</td>
<td>Methods&lt;br&gt;&amp; Deterministic mathematical model to describe the transmission of SARS-CoV-2 and subsequent disease progression (applying age-specific probabilities to the demographic structure and expected contact patterns in French Guiana, ...)&lt;br&gt;&lt;br&gt;Findings&lt;br&gt;&amp; The combination of these interventions coincided with a reduction in the basic reproduction number of SARS-CoV-2 from 1.7 to 1.1, which was sufficient to avoid hospital saturation.&lt;br&gt;&amp; We estimate that thanks to the young demographics, the risk of hospitalisation following infection was 0.3 times that of metropolitan France and that about 20% of the population was infected by July&lt;br&gt;&amp; Our model projections are consistent with a recent seroprevalence study. The study show-cases how mathematical modelling can be used to support healthcare planning in a context of high uncertainty.</td>
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<td>Clin Infect Dis. 12MAR2021</td>
<td>Household SARS-CoV-2 transmission and children: a network prospective study</td>
<td>Soriano-Arandes A., Spain [gotopaper]</td>
<td>Public Health / Epidemiology</td>
<td><strong>Aim:</strong> describe the epidemiological and clinical characteristics of children with COVID-19 in Catalonia (Spain) and investigate the dynamics of household transmission. Prospective, observational, multicenter study performed during summer and school periods (1 July-31 October, 2020) on COVID-19 patients &lt;16 years.  &gt; The study included 1040 COVID-19 patients &lt;16 years. 47.2% were asymptomatic, 10.8% had comorbidities, and 2.6% required hospitalization. No deaths were reported.  &gt; Viral transmission was common among household members (62.3%).  &gt; More than 70% (756/1040) of pediatric cases were secondary to an adult, whereas 7.7% (80/1040) were index cases.  &gt; The Secondary Attack Rate (SAR) was significantly lower in households with COVID-19 pediatric index cases during the school period relative to summer (p=0.02), and when compared to adults (p=0.006).  &gt; No individual or environmental risk factors associated with the SAR were identified.  <strong>Children are unlikely to cause household COVID-19 clusters or be major drivers of the pandemic even if attending school.</strong></td>
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<td>Nature 11MAR2021</td>
<td>Sensitivity of SARS-CoV-2 B.1.1.7 to mRNA vaccine-elicited antibodies</td>
<td>Collier A.C. et al. [International [gotopaper]</td>
<td>Vaccines - Variants</td>
<td><strong>Assessment of immune responses following vaccination with mRNA-based vaccine BNT162b2.</strong>  <strong>Methods</strong>  37 participants (median age 62 years; 35% female) measurement of neutralising antibody responses following first and second immunisations using pseudoviruses expressing the wild-type Spike protein or the 8 amino acid mutations found in the B.1.1.7 spike protein.  <strong>Findings</strong>  &gt; The GMT against wild type (WT) following the second dose of vaccine is substantially higher than after the first dose (318 vs 77). Correlation between total Spike IgG titres and serum neutralisation titres  &gt; Broad range of T cell responses (IFN-Gamma). No correlation with serum neutralization titers  &gt; Vaccine sera exhibited a broad range of neutralising titres against the wild-type pseudoviruses that were modestly reduced against B.1.1.7 variant. Reduction also evident in sera from some convalescent patients.  &gt; Decreased B.1.1.7 neutralisation also observed with monoclonal antibodies targeting the N-terminal domain (9 out of 10), the RBM (5 out of 31), but not in RBM neutralising mAbs binding outside the RBM.  &gt; Introduction of the E484K mutation in a B.1.1.7 background to reflect a newly emergent Variant of Concern (VOC 202102/02) led to a more substantial loss of neutralising activity by vaccine-elicited antibodies and mAbs (19 out of 31) over that conferred by the B.1.1.7 mutations alone.  <strong>Conclusion:</strong>  &gt; Pseudovirus bearing S protein with the full set of mutations present in the B.1.1.7 variant result in small reduction in neutralisation by sera from BNT162B2 vaccinees (more marked following the first dose than the second dose). This could be related to increased breadth/ potency/concentration of antibodies following the boost dose.  &gt; E484K emergence on a B.1.1.7 background represents a threat to the vaccine BNT162b</td>
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**Methods:**  
> Randomized, double-blind, placebo-controlled phase 1 clinical trial of Ad26.COV2.S (NCT04436276).  
> Twenty-five participants; interim analysis at day 71. A single clinical site in Boston  
> 1 or 2 intramuscular injections with 5 x 10^10 viral particles or 1 x 10^11 viral particles of Ad26.COV2.S vaccine or placebo (day 1 and day 57).  
**Main Outcomes and Measure:** Humoral immune responses included binding and neutralizing antibody responses at multiple time points following immunization. Cellular immune responses included immunospot-based and intracellular cytokine staining assays to measure T-cell responses.  
**Findings:**  
> Binding and neutralizing antibodies emerged rapidly by day 8 after initial immunization in 90% and 25% of vaccine recipients, respectively.  
> By day 57, binding and neutralizing antibodies were detected in 100% of vaccine recipients after a single immunization.  
> On day 71, the geometric mean titers of spike-specific binding antibodies were 2432 to 5729 and the geometric mean titers of neutralizing antibodies were 242 to 449 in the vaccinated groups.  
> A variety of antibody subclasses, Fc receptor binding properties, and antiviral functions were induced. CD4+ and CD8+ T-cell responses were induced.  
**Conclusion:** Ad26.COV2.S induces rapid binding and neutralization antibody responses as well as cellular immune responses. |
| PNAS             | A safe and highly efficacious measles virus-based vaccine expressing SARS-CoV-2 stabilized prefusion spike | Lu M., et al. USA | Vaccines          | Evaluation of a SARACoV 2 Measles virus (rMeV) vaccine efficacy in cotton rat, IFNAR−/−/mice, IFNAR−/−/hCD46 mice, and golden Syrian hamsters. Recombinant attenuated vaccine candidates expressing various forms of the SARS-CoV-2 spike (S) protein and its receptor binding domain (RBD).  
**Findings:**  
> rMeV expressing stabilized prefusion S protein (rMeV-preS) was more potent in inducing SARS-CoV-2-specific neutralizing antibodies than rMeV expressing full-length S protein (rMeV-S).  
> rMeVs expressing different lengths of RBD (rMeV-RBD) were the least potent.  
> Animals immunized with rMeV-preS produced higher levels of neutralizing antibody than found in convalescent sera from COVID-19 patients and a strong Th1-biased T cell response.  
> rMeV-preS also provided complete protection of hamsters from challenge with SARS-CoV-2, preventing replication in lungs and nasal turbinates, body weight loss, cytokine storm, and lung pathology.  
**Conclusion:** rMeV-preS is a safe and highly efficacious vaccine candidate, supporting its further development as a SARS-CoV-2 vaccine. |
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<td><strong>BMJ</strong> 10MAR2021</td>
<td>Risk of mortality in patients infected with SARS-CoV-2 variant of concern 202012/1: matched cohort study</td>
<td>Challen R., et al. UK <a href="#">gotopaper</a></td>
<td>Public Health / Epidemiology</td>
<td>To establish whether there is any change in mortality from infection with a new variant of SARS-CoV-2, designated a variant of concern (VOC-202012/1) in December 2020, compared with circulating SARS-CoV-2 variants. <strong>Methods</strong>  &gt; Matched cohort study (participants were matched on age, sex, ethnicity, index of multiple deprivation, lower tier local authority region, and sample date of positive specimens, and differed only by detectability of the spike protein gene using the TaqPath ass)  &gt; Community based (pillar 2) covid-19 testing centres in the UK using the TaqPath assay (a proxy measure of VOC-202012/1 infection)  &gt; 54,906 matched pairs of participants who tested positive for SARS-CoV-2 in pillar 2 between 1 October 2020 and 29 January 2021, followed-up until 12 February 2021 &gt; Main outcome measure: Death within 28 days of the first positive SARS-CoV-2 test result. <strong>Findings</strong>  &gt; The mortality hazard ratio associated with infection with VOC-202012/1 compared with infection with previously circulating variants was 1.64 (95% confidence interval 1.32 to 2.04), corresponding to 64% increased risk of death, in patients who tested positive for covid-19 in the community.  &gt; In this comparatively low risk group, this represents an increase in deaths from 2.5 to 4.1 per 1000 detected cases. Increased risk of mortality is increased by infection with VOC-202012/01 is highly probable. If this finding applies to other populations, infection with VOC-202012/1 could cause substantial additional mortality compared with previously circulating variants. Healthcare capacity planning and national and international control policies are all impacted by this finding, which supports further coordinated and stringent measures to reduce deaths.</td>
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<td>Nature Med. 10MAR2021</td>
<td>Attributes and predictors of long COVID</td>
<td>Sudre C.H., et al. UK <a href="#">gotopaper</a></td>
<td>Clinics - Long Covid</td>
<td>Analysis of prevalence, risk factors and early predictors of long COVID.  &gt; 4,182 incident cases of COVID-19 in which individuals self-reported their symptoms prospectively in the COVID Symptom Study app.  &gt; 558 (13.3%) participants reported symptoms lasting ≥28 days, 189 (4.5%) for ≥8 weeks and 95 (2.3%) for ≥12 weeks  &gt; Long COVID was characterized by symptoms of fatigue, headache, dyspnea and anosmia and was more likely with increasing age and body mass index and female sex  &gt; Experiencing more than five symptoms during the first week of illness was associated with long COVID (odds ratio = 3.53 (2.76–4.50)).  &gt; A simple model to distinguish between short COVID and long COVID at 7 days is presented, which could be used to identify individuals at risk of long COVID.</td>
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<td>Nature 09MAR2021</td>
<td>Emergence of a SARS-CoV-2 variant of concern with mutations in spike glycoprotein</td>
<td>Tegally H., et al. South Africa <a href="#">gotopaper</a></td>
<td>Virology</td>
<td>B.1.351 lineage (VOC 501Y.V2):  &gt; Shows marked hypermutation: 6 non-synonymous mutations in the spike protein by to 15/10/20, then 3 more by 30/11/20, plus deletion of 3 amino acids  &gt; Mutations N501Y, E484K and K417N are at key residues of the RBD – the two latters are key for neutralizing antibody binding  &gt; E484 and N501 pattern of nucleotide variation suggest evolution under positive selection  &gt; B.1.351 most likely evolved by mutation on circulating intermediate mutants  &gt; B.1.351 likely emerged in Nelson Madela Bay in early August and became dominant in Easter Cape, Western Cape and KwaZulu-Natal Provinces within weeks  &gt; It has a selective advantage, from increased transmissibility and/or immune escape</td>
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BBV152 is a whole-virion inactivated SARS-CoV-2 vaccine (3 μg or 6 μg) formulated with a toll-like receptor 7/8 agonist molecule (IMDG) adsorbed to alum (Algel).

**Methods**

- Double-blind, randomised, multicentre, phase 2 clinical trial NCT04471519 to evaluate the immunogenicity and safety of BBV152 in healthy adults and adolescents (aged 12–65 years) at nine hospitals in India.
- Phase 1 trial data allowed to choose phase II formulations of BBV152: 3 μg and 6 μg with Algel-IMDG administered on day 0 and day 28
- Participants with positive SARS-CoV-2 nucleic acid and serology tests were excluded.

**Primary outcome:** SARS-CoV-2 wild-type neutralising antibody titres and seroconversion rates at 4 weeks after the second dose

**Secondary outcome:** Cell-mediated responses (T-helper-1 profiling at 2 weeks after the second dose)

**Safety** assessed in all participants who received at least one dose of the vaccine

**Findings**

- 380 participants enrolled and randomly assigned to the 3 μg with Algel-IMDG group (n=190) or 6 μg with Algel-IMDG group (n=190).
- GMTs; PRNT50 at day 56 were significantly higher in the 6 μg with Algel-IMDG group (197.0 [95% CI 155.6–249.4]) than the 3 μg with Algel-IMDG group (100.9 [74.1–137.4]; p=0.0041).
- Seroconversion based on PRNT50 at day 56 was reported in 171 (92.9% [95% CI 88.2–96.2]) of 184 participants in the 3 μg with Algel-IMDG group and 174 (98.3% [95.1–99.6]) of 177 participants in the 6 μg with Algel-IMDG group.
- GMTs (MNT50) at day 56 were reported in 162 (88.0% [95% CI 82.4–92.3]) of 184 participants in the 3 μg with Algel-IMDG group and 171 (96.6% [92.8–98.5]) of 177 participants in the 6 μg with Algel-IMDG group.
- The 3 μg with Algel-IMDG and 6 μg with Algel-IMDG formulations elicited T-cell responses that were biased to a Th1 phenotype at day 42.
- No significant difference in the proportion of participants who had a solicited local or systemic adverse reaction in the 3 μg with Algel-IMDG group (38 [20.0%; 95% CI 14.7–26.3]) of 190) and the 6 μg with Algel-IMDG group (40 [21.1%; 15.5–27.5] of 190) was observed on days 0–7 and days 28–35; no serious adverse events were reported in the study.

**Conclusion**

BBV152 induced high neutralising antibody responses that remained elevated in all participants at 3 months after the second vaccination. The 6 μg with Algel-IMDG formulation has been selected for the phase 3 efficacy trial.

**Findings**

> Coexposure to airborne pollen enhances susceptibility to respiratory viral infections, regardless of the allergy status.
> We hypothesized this could be also true for SARS-CoV-2 infections.

**Methods**

- Test for relationships between SARS-CoV-2 infection rates and pollen concentrations, along with humidity, temperature, population density, and lockdown effects
- Our unique dataset derives from 130 sites in 31 countries and across five continents (8,019 data points)

**Findings**

- Pollen, some-times in synergy with humidity and temperature, explained, on average, 44% of the infection rate variability
- Lockdown halved infection rates under similar pollen concentrations
- As we cannot completely avoid pollen exposure, we suggest wide dissemination of pollen–virus coexposure information to encourage high-risk individuals to wear particle filter masks during high springtime pollen concentrations
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| Nature 08MAR2021 | Antibody Resistance of SARS-CoV-2 Variants B.1.351 and B.1.1.7 | Wang P., et al. USA gotopaper | Virology | **Background:** Authorized therapeutic or preventive interventions against COVID are directed toward the initial SARS-CoV-2 that emerged in 2019. The recent emergence of new SARS-CoV-2 variants B.1.1.7 in the UK11 and B.1.351 in South Africa is of concern because of their purported ease of transmission and extensive mutations in the spike protein.  
**Findings:** Monoclonal antibodies: neutralizing activity of 12 RBD mAbs against authentic B.1.1.7 and B.1.351 viruses, as compared to the original SARS-CoV-2 strain (WT), in Vero E6 cells  
> neutralization of B.1.1.7: only the activities of 910-3022 and S3095 are significantly impaired.  
> neutralization of B.1.351: the activities of 910-30, 2-1520, LY-CoV555 (bamlanivimab)1,23, C12124, and REGN10933 (casirivimab)2-720,27, REGN10987 (imdevimab), C13524, and S309 retain their activities against B.1.351  
Convalescent plasma from 20 patients more than one month after documented SARS-CoV-2 infection in the Spring of 2020. > Most (16 of 20) plasma samples lost >2.5-fold neutralizing activity against B.1.351, while maintaining activity against B.1.1.7. Only plasma from 4 patients retain neutralizing activities similar to those against the WT.  
Vaccinee sera obtained from 12 participants of a Phase 1 clinical trial of Moderna SARS-Co2 mRNA-1273 Vaccine conducted at the NIH. > Each vaccinee serum sample was assayed for neutralization against B.1.1.7, B.1.351, and WT viruses. No loss of neutralizing activity against B.1.1.7, whereas every sample lost activity against B.1.351. |
| Blood Advances 08MAR2021 | Heterogeneous NLRP3 inflammasome signature in circulating myeloid cells as a biomarker of COVID-19 severity | Courjon J., et al. France gotopaper | Immunology | The NLRP3 inflammasome can play a crucial role during innate immunity activation, but NLRP3 response during SARS-CoV-2 infection in patients is unknown.  
**Aim:** Prospectively monitoring of caspase-1 activation levels in peripheral myeloid cells from healthy donors and patients with mild to critical COVID-19.  
> The caspase-1 activation potential in response to NLRP3 inflammasome stimulation was opposed between nonclassical monocytes and CD66b+CD16dim granulocytes in severe and critical COVID-19 patients.  
> CD66b+CD16dim granulocytes had decreased nigericin-triggered caspase-1 activation potential associated with an increased percentage of NLRP3 inflammasome impaired immature neutrophils and a loss of eosinophils in the blood.  
> In patients who recovered from COVID-19, nigericin-triggered caspase-1 activation potential in CD66b+CD16dim cells was restored and the proportion of immature neutrophils was similar to control.  
NLRP3 inflammasome activation potential differs among myeloid cells. It could be used as a biomarker of COVID-19 patient evolution. |
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Impact on antibody neutralization of a panel of authentic SARS-CoV-2 variants including a B.1.1.7 isolate, chimeric strains with South African or Brazilian spike genes and isogenic recombinant viral variants with designed mutations or deletions at positions 69-70, 417, 478, 501, 614 and/or 681 of the spike protein, using using monoclonal antibodies (mAbs), animal immune sera, human convalescent sera and human sera from recipients of the BNT162b2 mRNA vaccine

**Findings:**
> in vitro experiments using a B.1.1.7 isolate and engineered variants in the backbone of the WA1/2020 strain establish that mutations in the spike can impact the potency of antibody neutralization
> Some neutralizing mAbs targeting the base of the RBD or NTD showed reduced activity against the B.1.1.7 isolate, whereas others targeting the RBM or NTD failed to inhibit infection of Wash SA-B.1.351, Wash BR-B.1.1.248 or variants containing the E484K mutation
> E484K substitution as a vulnerability for multiple neutralizing mAbs
> Several other highly neutralizing mAbs (such as COV2-2196, COV2-2381, COV2-2025 and S2E12) showed intact or mildly diminished inhibitory activity against the suite of variant viruses we tested, possibly because they bind the RBM at sites other than the E484K residue
> Cocktails of mAbs binding different epitopes of the spike protein overcame virus resistance to individual mAbs
> Studies with human sera from convalescent patients and recipients of the BNT162b2 mRNA vaccine and animal sera after immunization with a vaccine encoding a similar spike gene, demonstrate a lower potency of neutralization against E484K and N501Y-containing viruses
> Convalescent and vaccine-induced immune sera neutralized infection of the chimeric SARS-CoV-2 strains encoding the Brazilian spike (B.1.1.248) better than the South African spike (B.1.351) even though both viruses encoded E484 and N501 mutations

**Conclusion:**
Adjustments to some therapeutic antibody cocktails or existing spike sequences in vaccines might be necessary, corroborating in vivo studies are needed.

| JAMA 04MAR2021 | Effect of Ivermectin on Time to Resolution of Symptoms Among Adults With Mild COVID-19A Randomized Clinical Trial | Lopez-Medina E., et al. Colombia/USA [gotopaper] | Therapeutics | **Aim:** To determine whether ivermectin is an efficacious treatment for mild COVID-19.

Double-blind, randomized trial conducted at a single site in Cali, Colombia, on adult patients with mild disease and symptoms for 7 days or fewer (enrolment July 15- November 30, followed up through December 21, 2020)

Patients were randomized to receive ivermectin, 300 μg/kg of body weight per day for 5 days (n = 200) or placebo (n = 200).

**Primary outcome:** time to resolution of symptoms within a 21-day follow-up period.

**Results**
> 398 patients randomized in primary analysis population (median age, 37yo; 58% women)
> Median time to resolution of symptoms was 10 days (IQR, 9-13) in the ivermectin group compared with 12 days (IQR, 9-13) in the placebo group [hazard ratio, 1.07 (95% CI, 0.87 to 1.32]; P = .53 by log-rank test].
> By day 21, 82% in the ivermectin group and 79% in the placebo group had resolved symptoms.
> The most common solicited adverse event was headache in 104 patients (52%) given ivermectin and 111 (56%) who received placebo.
> The most common serious adverse event was multiorgan failure, occurring in 4 patients (2 in each group).

**Conclusion**
Among adults with mild COVID-19, a 5-day course of ivermectin, compared with placebo, did not significantly improve the time to resolution of symptoms.
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| Blood 03MAR2021 | The SARS-CoV-2 receptor-binding domain preferentially recognizes blood group A | Wu S.C., et al. | Virology           | > The RBD of SARS-CoV-2 shares sequence similarity with an ancient lectin family known to bind blood group antigens  
> Examined SARS-CoV-2 RBD binding with RBCs isolated from blood group A, B, or O individuals |
|                  |                                                                      | USA gotopaper   |                    | Methods   |
|                  |                                                                      |                 |                    | > SARS-CoV receptor-binding domain (RBD) was cloned and purified  
> SARS-CoV-2 RBD was incubated with HEK293T cells, HEK293 T cells expressing angiotensin-converting enzyme 2 (ACE2), or red blood cells (RBCs), followed by detection with anti-His antibody (Anti-His-Tag mAb-Alexa Fluor 647) and flow cytometric analysis  
> Anti-A antibody was similarly used to detect the A antigen on blood group A RBCs |
|                  |                                                                      |                 |                    | Findings   |
|                  |                                                                      |                 |                    | > SARS-CoV-2 RBD binds the blood group A expressed on respiratory epithelial cells, directly linking bloodgroup A and SARS-CoV-2 |
|                  |                                                                      |                 |                    |           |
|                  |                                                                      |                 |                    |           |
| Lancet Respir Med. 04MAR2021 | Sarilumab in patients admitted to hospital with severe or critical COVID-19: a randomised, double-blind, placebo-controlled, phase 3 trial | Lescure FX., et al. | Therapeutics | Aim: to assess safety and efficacy of sarilumab, an interleukin-6 receptor inhibitor, in patients with severe (requiring supplemental oxygen by nasal cannula or face mask) or critical (requiring greater supplemental oxygen, mechanical ventilation, or extracorporeal support) COVID-19.  
60-day, randomised, double-blind, placebo-controlled, multinational phase 3 trial. Patients were randomly assigned (2:2:1 with permuted blocks of five) to receive intravenous sarilumab 400 mg, sarilumab 200 mg, or placebo.  
Primary endpoint: time to clinical improvement of two or more points (seven point scale ranging from 1 [death] to 7 [discharged from hospital]) in the modified intention-to-treat population.  
Secondary endpoint: proportion of patients alive at day 29. |
|                  |                                                                      | USA gotopaper   |                    | Findings   |
|                  |                                                                      |                 |                    | > 420 patients were randomly assigned and 416 received placebo (n=84 [20%]), sarilumab 200 mg (n=159 [38%]), or sarilumab 400 mg (n=173 [42%]).  
> At day 29, no significant differences were seen in median time to an improvement of two or more points between placebo (12.0 days [95% CI 9.0 to 15.0]) and sarilumab 200 mg (10.0 days [9.0 to 12.0]; hazard ratio [HR] 1.03 [95% CI 0.75 to 1.40]; log-rank p=0.96) or sarilumab 400 mg (10.0 days [9.0 to 13.0]; HR 1.14 [95% CI 0.84 to 1.54]; log-rank p=0.34), or in proportions of patients alive (77% [92%] of 84 patients in the placebo group; 143% [90%] of 159 patients in the sarilumab 200 mg group; difference Δ7·7 [9.0 to 12.0]; p=0.69 vs placebo; 159% [92%] of 173 patients in the sarilumab 400 mg group; difference Δ1.17 [0.0 to 7.3]; p=0.85 vs placebo).  
> At day 29, there were non-significant survival differences between sarilumab 400 mg (88%) and placebo (79%; difference +8 9% [95% CI Δ7·7 to 25·5]; p=0.25) for patients who had critical disease.  
> No unexpected safety signals were seen.  
> The rates of treatment-emergent adverse events were 65% (55 of 84) in the placebo group, 65% (103 of 159) in the sarilumab 200 mg group, and 70% (121 of 173) in the sarilumab 400 mg group, and of those leading to death 11% (nine of 84) were in the placebo group, 11% (17 of 159) were in the sarilumab 200 mg group, and 10% (18 of 173) were in the sarilumab 400 mg group.  
This trial did not show efficacy of sarilumab in patients admitted to hospital with COVID-19 and receiving supplemental oxygen. |
**Azithromycin for community treatment of suspected COVID-19 in people at increased risk of an adverse clinical course in the UK (PRINCIPLE): a randomised, controlled, open-label, adaptive platform trial**

**Authors and link**
PRINCIPLE Trial Collaborative Group
UK [gootopaper](https://doi.org/10.1016/S0140-6736(21)00275-5)

**Field of expertise**
Therapeutics

**Key facts**

**Aim:** To assess the effectiveness of azithromycin to treat suspected COVID-19 among people in the community who had an increased risk of complications.

Open-label, multi-arm, adaptive platform randomised trial, we randomly assigned people aged 65 years and older, or 50 years and older with at least one comorbidity, who had been unwell for 14 days or less with suspected COVID-19.

**Treatments:** usual care plus azithromycin 500 mg daily for three days, usual care plus other interventions, or usual care alone.

**Coprimary endpoints:** Within 28 days from randomisation: time to first self-reported recovery, and hospital admission or death related to COVID-19.

**Findings:**
- 2120 participants were included in the Bayesian primary analysis, 500 participants in the azithromycin plus usual care group, 823 in the usual care alone group, and 797 in other intervention groups.
- 402/500 (80%) participants in the azithromycin plus usual care group and 631/823 (77%) in the usual care alone group reported feeling recovered within 28 days.
- We found little evidence of a meaningful benefit in the azithromycin plus usual care group in time to first reported recovery versus usual care alone (hazard ratio 1·08, 95% Bayesian credibility interval [BCI] 0·95 to 1·23), equating to an estimated benefit in median time to first recovery of 0·94 days (95% BCI −0·56 to 2·43).
- The probability that there was a clinically meaningful benefit of at least 1·5 days in time to recovery was 0·23. 16/500 (3%) participants in the azithromycin plus usual care group and 28/823 (3%) participants in the usual care alone group were hospitalised (absolute benefit in percentage 0·3%, 95% BCI −1·7 to 2·2).
- No deaths in either study group. Safety outcomes were similar in both groups.

These findings do not justify the routine use of azithromycin for reducing time to recovery or risk of hospitalisation for people with suspected COVID-19 in the community.

**Human Safety, Tolerability, and Pharmacokinetics of Molnupiravir, a Novel Broad-Spectrum Oral Antiviral Agent with Activity Against SARS-CoV-2**

**Authors and link**
Painter W. P., et al.
USA [gootopaper](https://doi.org/10.1128/AAC.02528-21)

**Field of expertise**
Therapeutics

**Key facts**

> Molnupiravir, EIDD-2801/MK-4482, prodrug of the active antiviral ribonucleoside analog 14ß-d-N4-hydroxycytidine (NHC; EIDD-1931)
> Single and multiple doses of molnupiravir were evaluated in this first-in-human, phase 1, randomized, double-blind, placebo-controlled study in healthy volunteers, which included evaluation of the effect of food on pharmacokinetics.

**Findings**
- EIDD-1931 appeared rapidly in plasma, with a median time of maximum observed concentration of 1.00 to 1.75 hours, and declined with a geometric half-life of approximately 1 hour, with a slower elimination phase apparent following multiple doses or higher single doses (7.1 hours at 24th highest dose tested). Mean maximum observed concentration and area under the concentration versus time curve increased in a dose-proportional manner, and there was no accumulation following multiple doses. When administered in a fed state, there was a decrease in the rate of absorption, but no decrease in overall exposure.
- Molnupiravir was well tolerated. Fewer than half of subjects reported an adverse event, the incidence of adverse events was higher following administration of placebo, and 93.3% of adverse events were mild. One discontinued early due to rash. There were no serious adverse events and there were no clinically significant findings in clinical laboratory, vital signs, or electrocardiography.
- Plasma exposures exceeded expected efficacious doses based on scaling from animal models; therefore, dose escalations were discontinued before a maximum tolerated dose was reached.
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> This variant has an estimated 43–90% (range of 95% CI 38–130%) higher reproduction number than pre-existing variants. Its relative growth rate has declined slightly over time but it remains among the highest of any lineage as a function of lineage age  
> No increased or decreased severity of the disease associated to VOC 202012/01 was identified by the increased transmissibility model. A fitted two-strain dynamic transmission model shows that VOC 202012/01 will lead to large resurgences of COVID-19 cases.  
> VOC 202012/01 has spread globally and exhibits a similar transmission increase in Denmark (55%), Switzerland (74%), and the United States (59%).  
Without stringent control measures, COVID-19 hospitalisations and deaths across England in 2021 will exceed those in 2020. |
> Lineage B.1.35 1 is defined by nine changes in the spike protein relative to the Wuhan-1 D614G spike. These changes include N501Y, which confers enhanced affinity for ACE2 and clusters of substitutions in two immunodominant regions of spike, suggesting escape from neutralization.  
> Class 1 antibodies are most frequently elicited in SARS-CoV-2 infection and include an antibody response to an epitope only accessible in the RBD ‘up’ conformation. Class 2 antibodies use more diverse VH-genes and bind to RBD ‘up’ and RBD ‘down’ conformations of spike.  
> An analysis of 3 class 1 antibodies showed reduced binding capacities and neutralisation to 501Y.V2 pseudovirus. 3 class 2 antibodies failed to bind 501Y.V2 RBD and were unable to neutralize the 501Y.V2 pseudovirus as well.  
> This pseudovirus also exhibits substantial to complete escape from neutralization, but not binding, by convalescent plasma.  
Conclusion: The prospect of reinfection with antigenically distinct variants and foreshadows reduced efficacy of spike-based vaccines. |
| JAMA 01MAR2021 | Binding and Neutralization Antibody Titers After a Single Vaccine Dose in Health Care Workers Previously Infected With SARS-CoV-2 | Saadat S., et al. USA gotopaper | Therapeutics | Background:  
> Persons who have had COVID-19 are thought to have protective immunity and memory responses for at least 6 months. However, neither recall responses nor ideal vaccine dosing regimens have been studied in those previously infected with SARS-CoV-2.  
Methods:  
> HCW cohort. stratified into 3 groups: SARS-CoV-2 IgG-antibody negative (Ab-negative); IgG-positive asymptomatic COVID-19 (asymptomatic); and IgG-positive with history of symptomatic COVID-19 (symptomatic)  
> Participants were vaccinated with Pfizer-BioNTech or Moderna  
Findings:  
> 59 volunteers enrolled: 17 in the Ab-negative, 16 in the asymptomatic, and 26 in the symptomatic group.  
> At 0, 7, and 14 days, median reciprocal half-maximal binding titers were higher in each of the asymptomatic (208, 29 364, and 34 033) and symptomatic (302, 32 301, and 35 460) groups compared with the Ab-negative group (<50, <50, and 924) (P < .001 for each).  
> At 0 and 14 days, median reciprocal ID99 virus neutralization titers of each of the asymptomatic (80 and 40 960) and symptomatic (320 and 40 960) groups were higher than the Ab-negative group (<20 and 80) (P < .001 for each).  
Conclusions: Health care workers with previous COVID-19 infection (laboratory-confirmed serology testing) had higher antibody titer responses to a single dose of mRNA vaccine than those not previously infected. |
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| Nature Commun. 26FEB2021 | Association between antecedent statin use and decreased mortality in hospitalized patients with COVID-19 | Gupta A., et al. USA [gotopaper](#) | Therapeutics | **Background:**
> Statins are known to have anti-inflammatory and antithrombotic properties but their benefit has not been assessed in COVID-19.

**Methods:**
> Retrospective analysis of patients admitted with COVID-19 from February 1st through May 12th, 2020 with study period ending on June 11th, 2020.
> Antecedent of statin use
> Multivariable logistic regression model to predict the propensity of receiving statins, adjusting for baseline sociodemographic and clinical characteristics, and outpatient medications.
> The primary endpoint includes in-hospital mortality within 30 days.

**Findings:**
> 2626 patients enrolled, of whom 951 (36.2%) were antecedent statin users.
> Among 1296 patients (648 statin users, 648 non-statin users) identified with 1:1 propensity-score matching, statin use is significantly associated with lower odds of the primary endpoint in the propensity-matched cohort (OR 0.47, 95% CI 0.36–0.62, p < 0.001).

**Conclusion:**
Antecedent statin use in patients hospitalized with COVID-19 is associated with lower inpatient mortality.

| Nature 26FEB2021 | SARS-CoV-2 spike D614G change enhances replication and transmission | Zhou B., et al. International [gotopaper](#) | Virology | **Aim:** to understand if the S-614G has a fitness advantage that improves replication and/or transmission in humans.

The S-614G variant:
> has enhanced binding to human host cell surface receptor ACE2
> has increased replication in primary human bronchial and nasal airway epithelial cultures and in a human ACE2 knock-in mouse model
> has markedly increased replication and transmissibility in hamster and ferret models of SARS-CoV-2 infection.

The S-614G substitution results in subtle increases in binding and replication in vitro, and it provides a real competitive advantage in vivo, particularly during the transmission bottleneck.

> prospective representative serological study were included based on IgG seroconversion to the Spike S1 protein of SARS-CoV-2 (N=353) with up to three consecutive serum samples per seroconverted participant (N=738)

**Findings**
> While SARS-CoV-2-specific IgM and IgA antibodies declined rapidly after the first month post onset of disease, specific IgG was still present in 92% (95% confidence interval, CI, 89-95) of the participants after 7 months.
> The estimated 2-fold decrease of IgG antibodies was 158 days (95% CI 136-189).
> Concentrations sustained better in persons reporting significant symptoms compared to asymptomatic persons or those with mild upper respiratory complaints only.
> Similarly, avidity of IgG antibodies for symptomatic persons showed a steeper increase over time compared with persons with mild or no symptoms (p=0.022).

IgG antibodies sustain in 92% of the participants after 7 months post onset of symptoms whereas IgM and IgA antibodies wane. Concentrations are higher in symptomatic persons and avidity increases with time.

SARS-CoV-2-specific IgG antibodies persist and show increasing avidity over time, indicative of underlying immune maturation. These data support development of immune memory against SARS-CoV-2 providing insight into protection of the general unvaccinated part of the population.
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**Methods:** Case series of 1116 patients aged younger than 21 years hospitalized between March 15 and October 31, 2020, at 66 US hospitals in 31 states.  
**Findings:**  
> Of 1116 patients (median age, 9.7 years; 45% female), 539 (48%) were diagnosed with MIS-C and 577 (52%) with COVID-19.  
> Compared with patients with COVID-19, patients with MIS-C were more likely to be 6 to 12 years old (40.8% vs 19.4%; absolute risk difference [RD], 21.4%; adjusted risk ratio [aRR], 1.51 vs 0-5 years) and non-Hispanic Black (32.3% vs 21.5%; RD, 10.8%; aRR, 1.43 vs White).  
> Patients with MIS-C had higher neutrophil to lymphocyte ratio (median, 6.4 vs 2.7), higher C-reactive protein level (median, 152 mg/L vs 33 mg/L), and lower platelet count (<150 ×103 cells/μL [212/523 (41%) vs 84/486 (17%)].  
> A total of 398 patients (73.8%) with MIS-C and 253 (43.8%) with COVID-19 were admitted to the intensive care unit, and 10 (1.9%) with MIS-C and 8 (1.4%) with COVID-19 died during hospitalization.  
> Among patients with MIS-C with reduced left ventricular systolic function (34.2%) and coronary artery aneurysm (13.4%), an estimated 91.0% and 79.1%, respectively, normalized within 30 days. |
| JAMA Intern Med. 24FEB2021 | Association of SARS-CoV-2 Seropositive Antibody Test With Risk of Future Infection | Harvey R.A., et al. UK [gotopaper](#) | Diagnostics        | **Aim:** To evaluate evidence of SARS-CoV-2 infection based on diagnostic nucleic acid amplification test (NAAT) among patients with positive vs negative test results for antibodies in an observational descriptive cohort study of clinical laboratory and linked claims data.  
**Methods:** The study created cohorts from a deidentified data set composed of commercial laboratory tests, medical and pharmacy claims, electronic health records, and hospital chargemaster data. The cohort included 3 257 478 unique patients.  
**Findings:** From 3 257 478 unique patients with an index antibody test; 56% were female with a median (SD) age of 48 (20) years. Of these, 2 876 773 (88.3%) had a negative index antibody result, and 378 606 (11.6%) had a positive index antibody result.  
> Patients with a negative antibody test result were older than those with a positive result (mean age 48 vs 44 years).  
> Of index-positive patients, 18.4% converted to seronegative over the follow-up period.  
> During the follow-up periods, the ratio of positive NAAT results among individuals who had a positive antibody test at index vs those with a negative antibody test at index was 2.85 at 0 to 30 days, 0.67 at 31 to 60 days, 0.29 at 61 to 90 days, and 0.10 at more than 90 days.  
**Patients with positive antibody test results were initially more likely to have positive NAAT results, consistent with prolonged RNA shedding, but became markedly less likely to have positive NAAT results over time, suggesting that seropositivity is associated with protection from infection.** |
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<td>NEJM 24FEB2021</td>
<td>BNT162b2 mRNA Covid-19 Vaccine in a Nationwide Mass Vaccination Setting</td>
<td>Dagan N., et al. Israel <a href="#">gotopaper</a></td>
<td>Vaccines</td>
<td>Evaluation of the effectiveness of the BNT162b2 mRNA vaccine based on data from Israel’s largest health care organization. <strong>Findings</strong>  &gt; Each study group (vaccinated and control) included 596,618 persons.  &gt; Estimated vaccine effectiveness for the study outcomes at days 14-20 after the first dose and at ≥7 days after the second dose was as follows:  - for documented infection, 46% (95% confidence interval [CI], 40 to 51) and 92% (95% CI, 88 to 95);  - for symptomatic Covid-19, 57% (95% CI, 50 to 63) and 94% (95% CI, 87 to 98);  - for hospitalization, 74% (95% CI, 56 to 86) and 87% (95% CI, 55 to 100);  - for severe disease, 62% (95% CI, 39 to 80) and 92% (95% CI, 75 to 100). &gt; Estimated effectiveness in preventing death from Covid-19 was 72% (95% CI, 19 to 100) for days 14-20 after the first dose. &gt; Estimated effectiveness in specific subpopulations assessed for documented infection and symptomatic Covid-19 was consistent across age groups, with potentially slightly lower effectiveness in persons with multiple coexisting conditions. <strong>BNT162b2 mRNA vaccine is effective for a wide range of Covid-19–related outcomes, a finding consistent with that of the randomized trial.</strong></td>
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<td>Clin Infect Dis. 24FEB2021</td>
<td>Persistence of antibodies to SARS-CoV-2 in relation to symptoms in a nationwide prospective study</td>
<td>den Hartog G., et al. Netherlands <a href="#">gotopaper</a></td>
<td>Public Health / Epidemiology</td>
<td>Aim: to study changes in Immunoglobulin (Ig) isotype seropositivity and IgG binding strength of SARS-CoV-2-specific serum antibodies up to 7 months following onset of symptoms in a nationwide sample. <strong>Methods:</strong> prospective representative serological study in the Netherlands were included based on IgG seroconversion to the Spike S1 protein of SARS-CoV-2 (N=353), with up to three consecutive serum samples per seroconverted participant (N=738). IgM, IgA and IgG antibody concentrations to S1, and increase in IgG were determined. <strong>Findings:</strong>  &gt; While SARS-CoV-2-specific IgM and IgA Abs declined rapidly after the first month post onset of disease, specific IgG was still present in 92% of the participants after 7 months.  &gt; The estimated 2-fold decrease of IgG antibodies was 158 days.  &gt; Concentrations sustained better in persons reporting significant symptoms compared to asymptomatic persons or those with mild upper respiratory complaints only.  &gt; SARS-CoV-2-specific IgG antibodies persist and show increasing avidity over time, indicative of underlying immune maturation.</td>
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<td>Cell 23FEB2021</td>
<td>Extremely potent human monoclonal antibodies from COVID-19 convalescent patients</td>
<td>Andreano E., et al. Italy <a href="#">gotopaper</a></td>
<td>Therapeutics</td>
<td>&gt; 453 neutralizing antibodies were identified by single cell sorting 4,277 SARS-CoV-2 spike protein specific memory B cells from 14 COVID-19 survivors.  &gt; The most potent neutralizing antibodies recognized the spike protein receptor binding domain, followed in potency by antibodies recognizing the S1 domain, the spike protein trimer and the S2 subunit.  &gt; Only 1.4% of the antibodies neutralized the authentic virus with a potency of 1-10 ng/mL.  &gt; The most potent monoclonal antibody, engineered to reduce the risk of antibody dependent enhancement and prolong half-life, neutralized the authentic wild type virus and emerging variants containing D614G, E484K and N501Y substitutions.  &gt; Prophylactic and therapeutic efficacy in the hamster model was observed at 0.25 and 4 mg/kg respectively in absence of Fc-functions.</td>
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| Cell 23FEB2021 | No higher infectivity but immune escape of SARS-CoV-2 501Y.V2 variants | Li Q., *et al.* China [gotopaper](#) | Variants | > Experiments with 18 pseudotyped viruses showed that the 501Y.V2 variants do not confer increased infectivity in multiple cell types except for murine ACE2-overexpressing cells, where a substantial increase in infectivity was observed.  
> The susceptibility of the 501Y.V2 variants to 12 of 17 neutralizing monoclonal antibodies was substantially diminished.  
> Neutralization ability of the sera from convalescent patients and immunized mice was also reduced for these variants.  
> The neutralization resistance was mainly caused by E484K and N501Y mutations in the receptor-binding domain of Spike.  
> The enhanced infectivity in murine ACE2-overexpressing cells suggests the possibility of spillover of the 501Y.V2 variants to mice.  
> The neutralization resistance detected for the 501Y.V2 variants suggests the potential for compromised efficacy of monoclonal antibodies and vaccines. |
Methods: Neutralization of a B.1.351 viral isolate and compare it to 127 neutralization of Victoria, an early Wuhan related isolate. Neutralization assays were performed on a large panel of monoclonal Abs convalescent sera from early in the pandemic, sera from patients suffering from B.1.1.7 and finally from 130 recipients of the Oxford-AstraZenca and Pfizer-BioNTech vaccines.  
Findings:  
> The receptor binding domain mutations provide tighter ACE2 binding and widespread escape from monoclonal Ab neutralization largely driven by E484K although K417N and N501Y act together against some important antibody classes.  
> In a number of cases it would appear that convalescent and some vaccine serum offers limited protection against this variant.  
> Neutralization of B.1.351 by sera from naturally infected or vaccinated individuals is significantly reduced, leading in some cases to a complete inability to neutralize B.1.351 virus. |
| Lancet Infect Dis. 23FEB2021 | Identification and validation of clinical phenotypes with prognostic implications in patients admitted to hospital with COVID-19: a multicentre cohort study | Gutiérrez-Gutiérrez B., *et al.* Spain [gotopaper](#) | Clinics | Aim: to determine whether clinical phenotypes of patients with COVID-19 can be derived from clinical data, to assess the reproducibility of these phenotypes and correlation with prognosis, and to derive and validate a simplified probabilistic model for phenotype assignment.  
Methods: data from two cohorts: the COVID-19@Spain cohort, a retrospective cohort including 4035 consecutive adult patients admitted to 127 hospitals in Spain, and the COVID-19@HULP cohort, including 2226 consecutive adult patients admitted to a teaching hospital in Madrid. The authors developed a simplified probabilistic model for phenotype assignment, including 16 variables.  
Findings:  
> Three distinct phenotypes were derived in the derivation cohort:  
A: Younger patients with, less frequently male, had mild viral symptoms, and had normal inflammatory parameters (516 [19%] patients).  
B: patients with obesity, lymphocytopenia, and moderately elevated inflammatory parameters (1955 [73%]).  
C: older patients with more comorbidities and even higher inflammatory parameters than phenotype B (116 [8%]).  
> 30-day mortality rates were 2·5% for A patients, 30·5% for B patients and 60·7% for C patients.  
> The predicted phenotypes in the internal validation cohort and external validation cohort showed similar mortality rates to the assigned phenotypes (internal validation cohort: 5·3% for phen A, 31·3% for phen B, and 59·5% for phen C; external validation cohort: 3·7% for phen A, 23·7% for phen B, and 51·4% for phenotype C). |
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| Lancet 19FEB2021 | Single-dose administration and the influence of the timing of the booster dose on immunogenicity and efficacy of ChAdOx1 nCoV-19 (AZD1222) vaccine: a pooled analysis of four randomised trials | Voysey M., et al. UK [gottopaper](#) | Vaccines | - Prespecified pooled analysis of trials of ChAdOx1 nCoV-19 (Single blinded: one phase 1/2; UK; one phase 2/3; UK; one phase 3, Brazil. Double-blinded: one phase 1/2, South Africa)
- Exploratory analyses of the impact on immunogenicity and efficacy of extending the interval between priming and booster doses.
- Immunogenicity and protection afforded by the first dose, before a booster dose has been offered.

**FINDINGS**

> 24 422 participants across the four studies (Apr 23–Dec 6, 2020), 17 178 included in the primary analysis (8597 receiving ChAdOx1 nCoV-19, 8581 receiving control vaccine). 332 NAAT-positive infections met the primary endpoint of symptomatic infection >14 days after the second dose.

> Overall vaccine efficacy >14 days after the second dose was 66·7% (95% CI 57·4–74·0), with 84/8597 (1·0%) cases in the ChAdOx1 nCoV-19 group and 248/8581 (2·9%) in the control group.

> There were no hospital admissions for COVID-19 in the ChAdOx1 nCoV-19 group after the initial 21-day exclusion period, and 15 in the control group.

> 108/12 282 (0·9%) participants in the ChAdOx1 nCoV-19 group and 127/11 962 (1·1%) in the control group had serious adverse events. There were 7 deaths considered unrelated to vaccination (2 in the ChAdOx1 nCoV-19 group and 5 in the control group), including one COVID-19-related death in one participant in the control group.

> Exploratory analyses showed that vaccine efficacy after a single standard dose from day 22 to day 90 after vaccination was 76·0% (59·3–85·9). Modelling analysis indicated that protection did not wane during this initial 3-month period.

> Antibody levels were maintained during this period with minimal waning by day 90 (geometric mean ratio [GMR] 0·66 [95% CI 0·59–0·74]).

> In the participants who received two standard doses, after the second dose, efficacy was higher in those with a longer prime-boost interval (vaccine efficacy 81·3% [95% CI 60·3–91·2] at ≥12 weeks) than in those with a short interval (vaccine efficacy 55·1% [33·0–69·9] at <6 weeks).

> Immunogenicity: binding antibody responses >2-fold higher after an interval of ≥12 or more weeks compared with an interval of <6 weeks in those who were aged 18–55 years (GMR 2·32 [2·01–2·68]).

The results of this primary analysis of two doses of ChAdOx1 nCoV-19 were consistent with those seen in the interim analysis of the trials and confirm that the vaccine is efficacious, with results varying by dose interval. A 3-month dose interval might have advantages over a programme with a short dose interval.
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**Methods:** retrospective cohort of 9109 vaccine-eligible HCWs, comparing vaccinated versus unvaccinated.  
**Findings:**  
> there were 170 SARS-CoV-2 infections among HCWs in the period between Dec 19, 2020, and Jan 24, 2021, of which 99 (58%) HCWs reported symptoms. Of the 170 HCWs who became infected, 89 (52%) were unvaccinated, 78 (46%) tested positive after the first dose, and 3 (2%) tested positive after the second dose.  
>Among the 125 infections that could be traced, 87 (70%) were community acquired and there were no nosocomial clusters.  
>Compared with a SARS-CoV-2 infection rate of 7·4 per 10,000 person-days in unvaccinated HCWs, infection rates were 5·5 per 10,000 person-days and 3·0 per 10,000 person-days on days 1–14 and 15–28 after the first dose of the vaccine, respectively.  
>Adjusted rate reductions of SARS-CoV-2 infections were 30% (95% CI 2–50) and 75% (72–84) for days 1–14 and days 15–28 after the first dose, respectively  
>Data show substantial early reductions in SARS-CoV-2 infection and symptomatic COVID-19 rates following first vaccine dose administration. |
**Methods:** Cases reported were screened for laboratory and clinical findings of potential reinfection followed by requests for medical records and laboratory specimens.  
**Findings:**  
> Among 73 potential reinfection patients with available records, 30 patients had recurrent COVID-19 symptoms explained by alternative diagnoses with concurrent SARS-CoV-2 positive RT-PCR.  
>24 patients remained asymptomatic after recovery but had recurrent or persistent RT-PCR.  
>19 patients had recurrent COVID-19 symptoms with concurrent SARS-CoV-2 positive RT-PCR but no alternative diagnoses. These 19 patients had symptom recurrence a median of 57 days after initial symptom onset.  
>Six of these patients had paired specimens available for further testing, but none had laboratory findings confirming reinfections.  
>No confirmation of SARS-CoV-2 reinfection within 90 days of the initial infection based on the clinical and laboratory characteristics of cases in this investigation. |
| Cell 18FEB2021 | Reduced neutralization of SARS-CoV-2 B.1.1.7 variant by convalescent and vaccine sera | Supasa P., et al. UK | Variants | Analysis of the ability of B.1.1.7 to evade antibody responses elicited by natural SARS-CoV-2 infection or vaccination, by mapping the impact of N501Y by structure/function analysis of a large panel of well-characterised monoclonal antibodies.  
> B.1.1.7 is harder to neutralize than parental virus, compromising neutralization by some members of a major class of public antibodies through light chain contacts with residue 501.  
> Original strain convalescent and vaccine sera show reduced B.1.1.7 neutralization  
> N501Y enhances RBD: ACE2 binding affinity 7-fold  
> N501Y compromises neutralisation by many antibodies with public V-region IGHV3-53  
> Widespread escape from monoclonal antibodies or antibody responses generated by natural infection or vaccination was not observed. |
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| Nature Commun. 18FEB2021 | Interleukin-3 is a predictive marker for severity and outcome during SARS-CoV-2 infections | Bénard A., et al. Germany [gotopaper](#) | Clinics | **Aim:** To identify IL-3 as an independent prognostic marker for the outcome during SARS-CoV-2 infections  
**Methods:** Prospective multicentric study. In total, 105 (32 non-severe; 32 severe; 41 recovered) patients positive for SARS-CoV-2 PCR from oral swabs, oral fluid, or BALF were enrolled. Blood samples were collected at the onset of symptoms (≤24 h), and 1, 2, 3, 4, 5, 6, or 7 days later; or after recovery from SARS-CoV-2 infection (time of recovery = 16 days ± 2 days).  
- A mouse model of pulmonary HSV-1 infection was used to characterize the IL-3 mechanism  
**Findings:**  
> Patients with severe COVID-19 exhibit reduced circulating plasmacytoid dendritic cells (pDCs) and low plasma IFNα and IFNβ levels when compared to non-severe COVID-19 patients.  
> In a mouse model of pulmonary HSV-1 infection, treatment with recombinant IL-3 reduces viral load and mortality. Mechanistically, IL-3 increases innate antiviral immunity by promoting the recruitment of circulating pDCs into the airways by stimulating CXCL12 secretion from pulmonary CD123+ epithelial cells.  
> Low plasma IL-3 levels are associated with increased severity, viral load, and mortality during SARS-CoV-2 infections.  
IL-3 might be a predictive disease marker for SARS-CoV-2 infections and recombinant IL-3, or CD123 receptor agonists, may therefore have the potential as novel therapeutic agents in SARS-CoV-2 infected patients. |
> All the 20 serum samples neutralized USA-WA1/2020 (pseudovirus wild-type) and all mutant viruses at titers of 1:40 or greater.  
> As compared with neutralization of USA-WA1/2020, neutralization of ΔΔ242-244+D614G virus was similar and neutralization of the B.1.351-spike virus was weaker by approximately two thirds.  
> Results suggest that virus with mutant residues in the receptor-binding site (K417N, E484K, and N501Y) is more poorly neutralized than virus with ΔΔ242-244, located in the N-terminal domain of the spike protein.  
It is unclear what effect a reduction in neutralization would have on BNT162b2-elicited protection from Covid-19 caused by the B.1.351 lineage. |
| NEJM 17FEB2021 | Serum Neutralizing Activity Elicited by mRNA-1273 Vaccine — Preliminary Report | Wu K., et al. USA [gotopaper](#) | Vaccines - variants | Pseudoviruses bearing the Wuhan-Hu-1 strain, the D614G substitution, the B.1.1.7 and B.1.351 variants and others were tested against sera from mRNA-1273-vaccinated individuals.  
> Both the full panel of mutations in S and a subset of mutations affecting the receptor-binding domain (RBD) region of the B.1.1.7 variant had no significant effect on neutralization.  
> A decrease in titers of neutralizing antibodies against the B.1.351 variant and a subset of its mutations affecting the RBD was observed.  
> In serum samples obtained 1 week after the participants received the second dose of vaccine, we detected reductions by a factor of 2.7 in titers of neutralizing antibodies against the partial panel of mutations and by a factor of 6.4 against the full panel of mutations.  
> Levels of neutralization against the other tested variants that were similar to those against the Wuhan-Hu-1 (D614) isolate.  
Protection against the B.1.351 variant conferred by the mRNA-1273 vaccine remains to be determined. |
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<td>NEJM 18FEB2021</td>
<td>Early High-Titer Plasma Therapy to Prevent Severe Covid-19 in Older Adults</td>
<td>Libster R., et al. Argentina <a href="#">gotopaper</a></td>
<td>Therapeutics</td>
<td>Randomized, double-blind, placebo-controlled trial of convalescent plasma with high IgG titers against SARS-CoV-2 in older adult patients within 72 hours after the onset of mild Covid-19 symptoms. Primary end point: severe respiratory disease, defined as a respiratory rate of 30 breaths per minute or more, an oxygen saturation of less than 93% while the patient was breathing ambient air, or both. The trial was stopped early at 76% of its projected sample size because cases of Covid-19 in the trial region decreased considerably. Findings &gt; A total of 160 patients underwent randomization &gt; In the intention-to-treat population, severe respiratory disease developed in 13 of 80 patients (16%) who received convalescent plasma and 25 of 80 patients (31%) who received placebo (relative risk, 0.52; 95% confidence interval [CI], 0.29 to 0.94; ( P=0.03 )), with a relative risk reduction of 48%. &gt; A modified intention-to-treat analysis that excluded 6 patients who had a primary end-point event before infusion of convalescent plasma or placebo showed a larger effect size (relative risk, 0.40; 95% CI, 0.20 to 0.81). &gt; No solicited adverse events were observed. Early administration of high-titer convalescent plasma against SARS-CoV-2 to mildly ill infected older adults reduced the progression of Covid-19.</td>
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<td>NEJM 18FEB2021</td>
<td>A Randomized Trial of Convalescent Plasma in Covid-19 Severe Pneumonia</td>
<td>Simonovich V.A., et al. Argentina <a href="#">gotopaper</a></td>
<td>Therapeutics</td>
<td>Aim: to gather further evidence of whether convalescent plasma improves clinical outcomes. Randomized trial on hospitalized adult patients with severe Covid-19 pneumonia in a 2:1 ratio to receive convalescent plasma or placebo. Primary outcome: the patient’s clinical status 30 days after the intervention, as measured on a six-point ordinal scale ranging from total recovery to death. Findings &gt; A total of 228 patients were assigned to receive convalescent plasma and 105 to receive placebo. Median time from the onset of symptoms to enrollment in the trial was 8 days, hypoxemia was the most frequent severity criterion for enrollment. &gt; The infused convalescent plasma had a median titer of 1:3200 of total SARS-CoV-2 antibodies (interquartile range, 1:800 to 1:3200). No patients were lost to follow-up. &gt; At day 30, no significant difference was noted between the convalescent plasma group and the placebo group in the distribution of clinical outcomes according to the ordinal scale (odds ratio, 0.83; 95% confidence interval [CI], 0.52 to 1.35; ( P=0.46 )). &gt; Overall mortality was 10.96% in the convalescent plasma group and 11.43% in the placebo group, for a risk difference of −0.46 percentage points (95% CI, −7.8 to 6.8). &gt; Total SARS-CoV-2 antibody titers tended to be higher in the convalescent plasma group at day 2 after the intervention. &gt; Adverse events and serious adverse events were similar in the two groups. No significant differences were observed in clinical status or overall mortality between patients treated with convalescent plasma and those who received placebo.</td>
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| Nature Commun. 16FEB2021 | Modelling safe protocols for reopening schools during the COVID-19 pandemic in France | Di Domenico L., et al. France gotopaper | Public Health / Epidemiology | **Aim:** to explored scenarios of partial, progressive, or full school reopening, through a stochastic age-structured transmission model.  
> Under a scenario with stable epidemic activity if schools were closed, reopening pre-schools and primary schools would lead to up to 76% [67, 84]% occupation of ICU beds if no other school level reopened, or if middle and high schools reopened later.  
> Immediately reopening all school levels may overwhelm the ICU system. Priority should be given to pre- and primary schools allowing younger children to resume learning and development. Full attendance in middle and high schools is not recommended for stable or increasing epidemic activity.  
> Large-scale test and trace is required for epidemic control. |
> Retrospective exploratory analysis using the Hospital Episode Statistics administrative dataset (between March 1 and May 31, 2020)  
> Multilevel logistic regression was used to model the relationship between death and several covariates: age, sex, deprivation (Index of Multiple Deprivation), ethnicity, frailty (Hospital Frailty Risk Score), presence of comorbidities (Charlson Comorbidity Index items), and date of discharge (whether alive or deceased).  
**Findings:**  
> 91,541 adult patients with COVID-19 were discharged during the study period, among which 28,200 (30.8%) in-hospital deaths occurred  
> Significant predictors of in-hospital death included older age, male sex, [1.457 [1.408–1.509]], greater deprivation (1.002 [1.001–1.003]), Asian (1.211 [1.128–1.299]) or mixed ethnicity (1.317 [1.080–1.605]), vs White ethnicity, and most of the assessed comorbidities, including moderate or severe liver disease (5.433 [4.618–6.392]).  
> Later date of discharge was associated with a lower odds of death (0.977 [0.976–0.978]); adjusted in-hospital mortality improved significantly in a broadly linear fashion, from 52.2% in the first week of March to 16.8% in the last week of May > might reflect the impact of changes in hospital strategy and clinical processes  
**Conclusion:**  
> The reasons for the observed improvements in mortality should be thoroughly investigated to inform the response to future outbreaks.  
> The higher mortality rate reported for certain ethnic minority groups in community-based studies compared with our hospital-based analysis might partly reflect differential infection rates in those at greatest risk, propensity to become severely ill once infected, and health-seeking behaviours. |
| Pediatrics 12FEB2021 | Factors Associated With Severe SARS-CoV-2 Infection | Ouldali N., et al. France gotopaper | Clincs | **Aim:** to analyze the clinical spectrum of hospitalized pediatric SARS-CoV-2 infection and predictors of severe disease evolution.  
**Main outcome:** proportion of children with severe disease, defined by hemodynamic or ventilatory (invasive or not) support requirement.  
> 397 hospitalized children with SARS-CoV-2 infection, with several clinical patterns (paucisymptomatic children, admitted for surveillance, lower respiratory tract infection or multisystem inflammatory syndrome).  
> Children <90 days old accounted for 37% of cases (145 of 397), but only 4 (3%) had severe disease.  
> Excluding children with multisystem inflammatory syndrome in children (n = 29) and hospitalized for a diagnosis not related to SARS-CoV-2 (n = 62), 23 of 306 (11%) children had severe disease, including 6 deaths.  
> Factors independently associated with severity were age ≥10 years (odds ratio [OR] = 3.4), hypoxemia (OR = 8.9), C-reactive protein level ≥80 mg/L (OR = 6.6).  
Young age was not an independent factor associated with severe SARS-CoV-2 infection, and children <90 days old were at the lowest risk of severe disease evolution. |
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| Nature Med. 12FEB2021 | Humoral signatures of protective and pathological SARS-CoV-2 infection in children | Bartsch Y.C., et al. USA [gotopaper](#) | Immunology | Aim: identifying immune mechanisms that result in disparate clinical phenotypes in children (largely asymptomatic disease, with rare reports of multisystem inflammatory syndrome in children (MIS-C)).  
> Using systems serology, in 25 children with acute mild COVID-19 we observed a functional phagocyte and complement-activating IgG response to SARS-CoV-2, similar to the acute responses generated in adults with mild disease. Conversely, IgA and neutrophil responses were significantly expanded in adults with severe disease.  
> Weeks after the resolution of SARS-CoV-2 infection, children who develop MIS-C maintained highly inflammatory monocyte-activating SARS-CoV-2 IgG antibodies, distinguishable from acute disease in children but with antibody levels similar to those in convalescent adults.  
These data provide insights into the potential mechanisms of IgG and IgA that might underlie differential disease severity in children infected with SARS-CoV-2 |
| Cell 12FEB2021 | Human neutralizing antibodies against SARS-CoV-2 require intact Fc effector functions for optimal therapeutic protection | Winkler ES., et al. USA [gotopaper](#) | Immunology | Aim: to define correlates of protection of neutralizing human monoclonal antibodies (mAbs) in SARS-CoV-2-infected animals.  
Methods: A K18-hACE2 transgenic mouse model of SARS-CoV-2 pathogenesis and a Fc region genetic variant form of IgG (LALA-73 PG) of a potent RBD-binding neutralizing mAb that cannot engage FcγRs or complement were used to define the role of Fc effector functions in antibody protection.  
Findings:  
> Fc effector functions are dispensable when neutralizing mAbs are administered as prophylaxis, but are required for optimal protection when given as post-exposure therapy.  
> When administered after SARS-CoV-2 infection, intact but not LALA-PG mAbs reduce viral burden and lung disease. Fc engagement by Abs decreases immune cell activation and levels of inflammatory cytokines  
> Neutralizing mAbs require monocytes and CD8+ T cells for maximal clinical and virological benefit. In hamsters, Fc effector functions of a neutralizing mAb are required to prevent weight loss, control viral infection, and limit inflammation.  
> Fc effector functions of neutralizing antibodies are necessary for optimal therapeutic outcome after SARS-CoV-2 infection |
| BMI 11FEB2021 | Early initiation of prophylactic anticoagulation for prevention of coronavirus disease 2019 mortality in patients admitted to hospital in the United States: cohort study | Rentsch CT., et al. UK/USA [gotopaper](#) | Therapeutics | Aim: To evaluate whether early initiation of prophylactic anticoagulation compared with no anticoagulation was associated with decreased risk of death among COVID-19 patients admitted to hospital in USA.  
Methods: Observational cohort study including 4297 patients admitted to hospital from 1 March to 31 July 2020  
> Main outcome: 30 day mortality  
> Secondary outcomes: inpatient mortality, initiating therapeutic anticoagulation (a proxy for clinical deterioration, including thromboembolic events), and bleeding that required transfusion.  
Findings:  
> From 4297 patients, 3627 (84.4%) received prophylactic anticoagulation within 24 hours of admission. More than 99% (n=3600) of treated patients received subcutaneous heparin or enoxaparin  
> 622 deaths occurred within 30 days of hospital admission, 513 among those who received prophylactic anticoagulation.  
> The cumulative incidence of mortality at 30 days was 14.3% among those who received prophylactic anticoagulation and 18.7% among those who did not.  
> Compared with patients who did not receive prophylactic anticoagulation, those who did had a 27% decreased risk for 30 day mortality (hazard ratio 0.73).  
> Receipt of prophylactic anticoagulation was not associated with increased risk of bleeding that required transfusion.  
Early initiation of prophylactic anticoagulation compared with no anticoagulation among COVID-19 patients admitted to hospital was associated with a decreased risk of 30 day mortality.  
> Weeks after the resolution of SARS-CoV-2 infection, children who develop MIS-C maintained highly inflammatory monocyte-activating SARS-CoV-2 IgG antibodies, distinguishable from acute disease in children but with antibody levels similar to those in convalescent adults.  
These data provide insights into the potential mechanisms of IgG and IgA that might underlie differential disease severity in children infected with SARS-CoV-2 |
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| Euro Surveill., 11FEB2021 | Impact of age, ethnicity, sex and prior infection status on immunogenicity following a single dose of the BNT162b2 mRNA COVID-19 vaccine: real-world evidence from healthcare workers, Israel, December 2020 to January 2021 | Jabal KA., et al. Israel [gotopaper](#) | Vaccine | Description of one dose immunogenicity of BNT162b2 vaccine in various age and ethnic groups. **Background:**  
- As at 25 January 2021, Israel had vaccinated 29.2% of its population with a single dose of vaccine (almost exclusively BNT162b2 mRNA from Pfizer/BioNTech).  
- Ziv Medical Center (ZMC), located in Safed, Israel, is a 350-bed hospital, staffed by a multi-ethnic workforce of ca 1,500 persons including Jews, Arabs and Druze among others. ZMC has offered the BNT162b2 mRNA-based vaccine to all its staff, including administrative and support staff, with no specific exclusion for pregnant women. As at 21 January 2021, one-dose uptake was ca 90%. **Findings:**  
- 519 participants to the study (19-77 years of age). IgGs levels measured at 21d  
- 475 (92%) had detectable anti-SARS-CoV-2 IgG. Among these, GMC was 68.6 AU/mL (95% CI: 64–73.6). No differences between ethnicity or sex. Titres decreasing with age.  
- 39 non-respondant: median age older than respondent (57 vs 45) and more likely to be Jewish (31/38 non-responders of known ethnicity, 82% vs 291/459 responders of known ethnicity; 63%)  
- IgGs level postvaccination were higher among those with previous evidence of infection (at least one order of magnitude regardless the titre before vaccination) (GMC 573 vs 61.5) **Conclusion:** age and ethnicity (but not sex) may be associated with the likelihood of non-response (findings based on 39 observations). |
| PNAS 09FEB2021 | Exhaled aerosol increases with COVID-19 infection, age, and obesity | Edwards D., et al. USA [gotopaper](#) | Public Health / Epidemiology | > Respiratory droplet generation and exhalation in human and nonhuman primate subjects with and without COVID-19 infection to explore whether SARS-CoV-2 infection, and other changes in physiological state, translate into observable evolution of numbers and sizes of exhaled respiratory droplets in healthy and diseased subject  
**Method**  
- Observational cohort study of the exhaled breath particles of 194 healthy human subjects  
- Experimental infection study of 8 nonhuman primates infected, by aerosol, with SARS-CoV-2  
**Findings**  
- Exhaled aerosol particles vary between subjects by three orders of magnitude, with exhaled respiratory droplet number increasing with degree of COVID-19 infection and elevated BMI-years  
- 18% of human subjects (35) accounted for 80% of the exhaled bioaerosol of the group (194), reflecting a superspreader distribution of bioaerosol analogous to a classical 20:80 superspreader of infection distribution  
- The capacity of airway lining mucus to resist breakup on breathing varies significantly between individuals with a trend to increasing with the advance of COVID-19 infection and body mass index multiplied by age (i.e., BMI-years). **Conclusion**  
- Our studies of exhaled aerosol suggest that a critical factor in these and other transmission events is the propensity of certain individuals to exhale large numbers of small respiratory droplets.  
- Understanding the source and variance of respiratory droplet generation, and controlling it via the stabilization of airway lining mucus surfaces, may lead to effective approaches to reducing COVID-19 infection and transmission  
- These findings suggest that quantitative assessment and control of exhaled aerosol may be critical to slowing the airborne spread of COVID-19 in the absence of an effective and widely disseminated vaccine. |
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| Nature 10FEB2021 | mRNA vaccine-elicited antibodies to SARS-CoV-2 and circulating variants | Wang Z., et al. USA gotopaper | Vaccines | Antibody and memory B cell responses in volunteers who received either the Moderna (mRNA-1273) or Pfizer-BioNTech (BNT162b2) vaccines  
**Findings:**  
> Eight weeks after the second vaccine injection volunteers showed high levels of IgM, and IgG anti-S and anti-RBD  
> Plasma neutralizing activity, and the relative numbers of RBD-specific memory B cells were equivalent to individuals who recovered from natural infection  
> Vaccine-elicited monoclonal antibodies potently neutralize SARS-CoV-2, targeting a number of different RBD epitopes in common with mAbs isolated from infected donors  
> However, neutralization by 14 of the 17 most potent mAbs tested was reduced or abolished by either K417N, or E484K, or N501Y mutations.  
> Activity against SARS-CoV-2 variants encoding E484K or N501Y or the K417N:E484K:N501Y combination was reduced by a small but significant margin.  
> The same mutations were selected when recombinant vesicular stomatitis virus (rVSV)/SARS-CoV-2 S was cultured in the presence of the vaccine elicited mAbs.  
**Conclusion:**  
This results suggest that the monoclonal antibodies in clinical use should be tested against newly arising variants, and that mRNA vaccines may need to be updated periodically to avoid potential loss of clinical efficacy. |
| Nature 09FEB2021 | Lasting antibody and T cell responses to SARS-CoV-2 in COVID-19 patients three months after infection | Jiang X.L., et al. China gotopaper | Immunology | Aim : Longitudinal assessment of 25 SARS-CoV-2-infected patients up to 3–4 months post-infection and analysis of the specific antibody and memory T cell responses over time.  
**Findings:**  
> All patients seroconvert for IgG against N, S, or RBD, as well as IgM against RBD, and produce neutralising antibodies (NAb) by 14 days post symptoms onset (PSO) with the peak levels attained by 15–30 days PSO.  
> Anti-SARS-CoV-2 IgG and NAb remain detectable and relatively stable 3–4 months PSO, whereas IgM antibody rapidly decay.  
> 65% of patients have detectable SARS-CoV-2-specific CD4+ or CD8+ T cell responses 3–4 months PSO.  
> T cell responses maintain in most recovered patients for at least 3–4 months after infection.  
**Assessment of the duration and resiliency of the SARS-CoV-2 antibody and T cell responses in a large cohort study would be desirable for validation of the results.** |
Follow up of 26 HCW with mild COVID-19 three weeks (D21), two months (M2) and three months (M3) after the onset of symptoms.  
**Findings:**  
> All the HCW had anti-receptor binding domain (RBD) IgA at D21, decreasing to 38.5% at M3 (p < 0.0001).  
> Concomitantly a significant decrease in NAb titers was observed between D21 and M2 (p > 0.03) and between D21 and M3 (p < 0.0001).  
> SARS-CoV-2 can elicit a NAb response correlated with anti-RBD antibody levels, however neutralizing activity declines, and may even be lost, in association with a decrease in systemic IgA antibody levels, from two months after disease onset.  
**Conclusions:**  
This short-lasting humoral protection supports strong recommendations to maintain infection prevention and control measures in HCW, and suggests that periodic boosts of SARS-CoV-2 vaccination may be required. |
Lancet Child Adolesc Health 08FEB2021

SARS-CoV-2 transmission among children and staff in daycare centres during a nationwide lockdown in France: a cross-sectional, multicentre, seroprevalence study

Lachassinne E., et al.

Public Health / Epidemiology

**Aim:** to estimate the seroprevalence of antibodies against SARS-CoV-2 in daycare centres that remained open for key workers’ children during a nationwide lockdown in France (March 15 – May 09, 2020).

> 327 children enrolled (mean age 1·9 years), 197 daycare centre staff (40 yrs), and 164 adults in the comparator group (42 yrs).

> Positive serological tests were observed for 14 children (raw seroprevalence 4·3%) and 14 daycare centre staff (7·7%). After accounting for imperfect sensitivity and specificity of the assay, we estimated that 3·7% of the children and 6·8% of daycare centre staff had SARS-CoV-2 infection.

> The comparator group fared similarly to the daycare centre staff; 9 participants had a positive serological test (raw seroprevalence 5·5%), leading to a seroprevalence of 5·0% after adjusting.

> An exploratory analysis suggested that seropositive children were more likely than seronegative children to have been exposed to an adult household member with laboratory-confirmed COVID-19 (6/14 [43%] vs 19/327 [6%], relative risk 7.1).

The proportion of young children in this sample with SARS-CoV-2 infection was low. Intrafamily transmission seemed more plausible than transmission within daycare centres.

Nature Med. 08FEB2021

Neutralization of SARS-CoV-2 spike 69/70 deletion, E484K and N501Y variants by BNT162b2 vaccine-elicited sera

Xie X., et al.

Immunology

> Examine the effect of several key spike mutations from the UK and SA strains on BNT162b2 vaccine-elicited neutralization.

**Methods**

> Engineered three SARS-CoV-2 containing key spike mutations from the newly emerged United Kingdom (UK) and South African (SA) variants

- Mutant N501Y virus contains the N501Y mutation that is shared by both the UK and SA variants

- Mutant Δ69/70 + N501Y +D614G virus contains two additional changes present in the UK variants: amino acid 69 and 70 deletion (Δ69/70) and D614G substitution (D614G mutation is dominant in circulating strains around the world)

- Mutant E484K + N501Y +D614G virus addition-ally contains the E484K substitution, which is also located in the viral RBD

> Neutralization assays with the same 20 sera samples

**Findings**

> All sera showed equivalent neutralization titers between the WT and mutant viruses, with differences of four-fold or less.

> Notably, 10 out of the 20 sera had neutralization titers against mutant Δ69/70 + N501Y + D614G virus that were twice their titers against the WT virus, whereas 6 out of the 20 sera had neutralization titers against mutant E484K + N501Y + D614G virus that were half their titers against the WT virus.

> The ratios of the neutralization GMTs of the sera against the N501Y, Δ69/70 + N501Y + D614G and E484K + N501Y + D614G viruses to their GMTs against the USA-WA1/2020 virus were 1.46, 1.41 and 0.81, respectively.

> Neutralization geometric mean titers (GMTs) of 20 BTN162b2 vaccine-elicited human sera against the three mutant viruses were 0.81- to 1.46-fold of the GMTs against parental virus, indicating small effects of these mutations on neutralization by sera elicited by two BTN162b2 doses.

> Clinical data are needed for firm conclusions about vaccine effectiveness against variant viruses.
Eur J Epidemiol. 06FEB2021

Evidence of early circulation of SARS-CoV-2 in France: findings from the population-based “CONSTANCES” cohort

Carrat F., et al. France gotopaper

Public Health / Epidemiology

Analysis of serological status for SARS-CoV-2 antibodies on serum samples routinely collected in 9144 adults from a French general population-based cohort (CONSTANCES).

- 353 participants with a positive anti-SARS-CoV-2 IgG test were identified, among whom 13 were sampled between November 2019 and January 2020.
- Evidence was confirmed by neutralizing antibodies testing.
- Investigations in 11 of these participants revealed evidence of symptoms possibly related to a SARS-CoV-2 infection or situations at risk of potential SARS-CoV-2 exposure.

These results suggest early circulation of SARS-CoV-2 in Europe.

Nature 05FEB2021

SARS-CoV-2 evolution during treatment of chronic infection

Kemp S.A., et al. UK gotopaper

Virology

Aim: to report chronic SARS-CoV-2 with reduced sensitivity to neutralising antibodies in an immune suppressed individual treated with convalescent plasma (whole genome ultradeep sequences over 23 time points spanning 101 days).

- Little change was observed in the overall viral population structure following two courses of remdesivir over the first 57 days.
- Following convalescent plasma therapy, large, dynamic virus population shifts were observed, with the emergence of a dominant viral strain bearing D796H in S2 and ΔH69/ΔV70 in the S1 N-terminal domain NTD of the Spike protein.
- As passively transferred serum antibodies diminished, viruses with the escape genotype diminished in frequency, before returning during a final, unsuccessful course of convalescent plasma.
- In vitro, the Spike escape double mutant bearing ΔH69/ΔV70 and D796H conferred modestly decreased sensitivity to convalescent plasma, whilst maintaining infectivity similar to wild type. D796H appeared to be the main contributor to decreased susceptibility but incurred an infectivity defect. The ΔH69/ΔV70 single mutant had two-fold higher infectivity compared to wild type, possibly compensating for the reduced infectivity of D796H.

These data reveal strong selection on SARS-CoV-2 during convalescent plasma therapy associated with emergence of viral variants with evidence of reduced susceptibility to neutralising antibodies.

Nature 05FEB2021

Efficacy and tolerability of bevacizumab in patients with severe Covid-19

Pang J., et al. China gotopaper

Therapeutics

Aim: to evaluate the efficacy of the anti-vascular endothelial growth factor (VEGF) drug bevacizumab for treatment of Covid-19 patients.

Single-arm trial (NCT04275414) including 26 patients with severe Covid-19 followed up for 28 days, from 2-centers (China and Italy).

Patients received a single dose of bevacizumab

Findings:
- PaO2/FiO2 values markedly increased at days 1 and 7 after bevacizumab administration compared to the baseline values.
- 24 of 26 patients (92%) showed improvement and 2 patients (8%) showed no change in oxygen-support within 28-day follow-up, 17 (65%) patients are discharged, and none show worsen oxygen-support status nor die.
- Significant reduction of lesion areas/ratios are shown in chest computed tomography (CT) or X-ray within 7 days.
- Of 14 patients with fever, body temperature normalizes within 72 h in 13 (93%) patients.
- Relative to comparable controls, bevacizumab shows clinical efficacy by improving oxygenation and shortening oxygen-support duration. Bevacizumab plus standard care is highly beneficial for patients with severe Covid-19.
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<th>Title</th>
<th>Authors and link</th>
<th>Field of expertise</th>
<th>Key facts</th>
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<td>Lancet Resp Med. 05FEB2021</td>
<td>Peginterferon lambda for the treatment of outpatients with COVID-19: a phase 2, placebo-controlled randomised trial</td>
<td>Feld J.J., et al. Canada <a href="#">gotopaper</a></td>
<td>Therapeutics</td>
<td>Aim – Test therapeutic effects of Peginterferon lambda (PGL), a type III interferon. Double-blind, placebo-controlled trial, on 60 outpatients with laboratory-confirmed COVID-19 receiving PGL (single subcutaneous injection, 180 μg) or placebo within 7 days of symptoms onset or first positive swab. Primary endpoint: proportion of patients who were negative for SARS-CoV-2 RNA on day 7 after the injection. &gt; The decline in SARS-CoV-2 RNA was greater in patients treated with PGL than placebo from day 3 onwards, with a difference of 2-42 log copies per mL at day 7 (p=0.0041). &gt; By day 7, 24 (80%) participants in the PGL group had an undetectable viral load, compared with 19 (63%) in the placebo group (p=0.15). &gt; After controlling for baseline viral load, patients in the PGL group were more likely to have undetectable virus by day 7 than were those in the placebo group (odds ratio [OR] 4.12). &gt; Of those with baseline viral load above 106 copies per mL, 15/19 (79%) in the PGL group had undetectable virus on day 7, compared with 6/16 (38%) in the placebo group (OR 6.25). &gt; PGL was well tolerated, and adverse events were similar between groups (mild and transient aminotransferase, concentration increases more frequently observed in the PGL group). Peginterferon lambda accelerated viral decline in outpatients with COVID-19, increasing the proportion of patients with viral clearance by day 7, particularly in those with high baseline viral load.</td>
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<td>Lancet Public Health 05FEB2021</td>
<td>COVID-19 vaccine hesitancy in a representative working-age population in France: a survey experiment based on vaccine characteristics</td>
<td>Schwarzinger M., et al. France <a href="#">gotopaper</a></td>
<td>Public Health / Epidemiology</td>
<td>Aim: to assess the effects of vaccine characteristics, information on herd immunity, and general practitioner recommendation on vaccine hesitancy in a representative working-age population in France. Findings: Online survey in July 2020, adults aged 18–64 years residing in France, with no history of SARS-CoV-2 infection. Responses were analysed with a two-part model to disentangle outright vaccine refusal from vaccine hesitancy. &gt;Outright vaccine refusal and vaccine hesitancy were both significantly associated with female gender, age, lower educational level, poor compliance with recommended vaccinations in the past, and no report of specified chronic conditions. &gt;Outright vaccine refusal was associated with a lower perceived severity of COVID-19. &gt;Vaccine hesitancy was lower when herd immunity benefits were communicated and in working versus non-working individuals, and those with experience of COVID-19 (Symptoms or close contact). &gt;For a mass vaccination campaign involving mass vaccination centres and communication of herd immunity benefits, the model predicted outright vaccine refusal in 29.4% of the French working-age population. &gt;Predicted hesitancy was highest for vaccines manufactured in China (vaccine acceptance 27.4%), and lowest for a vaccine manufactured in the EU (vaccine acceptance 61.3%).</td>
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<td><strong>Lancet 05FEB2021</strong></td>
<td>Factors associated with the spatial heterogeneity of the first wave of COVID-19 in France: a nationwide geo-epidemiological study</td>
<td>Gaudart J., et al. France gotopaper</td>
<td>Public Health / Epidemiology</td>
<td>&gt; better understand the factors associated with the heterogeneity of in-hospital COVID-19 morbidity and mortality across France</td>
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**Methods**

> Geo-epidemiological analysis was based on data publicly available for the 96 administrative departments of metropolitan France between March 19 and May 11, 2020, Assessment:

> Multidimensional variables (spatiotemporal spread of the epidemic, national lockdown, demographic population structure, baseline intensive care capacities, ...)

> in-hospital COVID-19 incidence, mortality, and case fatality rates

**Findings**

> clear spatial heterogeneity of in-hospital COVID-19 incidence and mortality rates, following the spread of the epidemic

> Delay between the first COVID-19-associated death and the onset of the national lockdown was positively associated with in-hospital incidence, mortality, and case fatality rates

> Mortality and case fatality rates were higher in departments with older populations (adjusted standardised ratio for populations with a high proportion older than aged >85 years 2·17 [95% CI 1·20–3·90] for mortality and 1·43 [1·08–1·88] for case fatality rate)

> Mortality rate was also associated with incidence rate (1·0004, 1·0002–1·001), but mortality and case fatality rates did not appear to be associated with baseline intensive care capacities

> No association between climate and in-hospital COVID-19 incidence, or between economic indicators and in-hospital COVID-19 incidence or mortality rates

This ecological study highlights the impact of the epidemic spread, national lockdown, and reactive adaptation of intensive care capacities on the spatial distribution of COVID-19 morbidity and mortality

| **Lancet Infect Dis. 03FEB2021** | Safety, tolerability, and immunogenicity of an inactivated SARS-CoV-2 vaccine (CoronaVac) in healthy adults aged 60 years and older: a randomised, double-blind, placebo-controlled, phase 1/2 clinical trial | Wu Z., et al. China gotopaper | Vaccines | > Randomised, double-blind, placebo-controlled, phase 1/2 clinical trial of CoronaVac in healthy adults aged 60 years and older (NCT04383574).

> Vaccine or placebo by IM injection (in two doses, days 0 and 28).

> Phase 1: dose-escalation study. 72 participants (24 per intervention group and 24 in the placebo group; mean age 65·9 years [SD 4·9])

- Block 1: 3 μg inactivated virus in 0·5 mL of aluminium hydroxide

- Block 2 (6 μg per injection).

> Phase 2: 1·5 μg, 3 μg, or 6 μg per dose, or placebo. 350 participants were enrolled in phase 2 (100 in each intervention group and 50 in the placebo group; mean age 66·6 years [SD 4·7] in 349 participants)

**Primary safety endpoint**: adverse reactions within 28 days after each injection in all participants who received at least one dose.

**Primary immunogenicity endpoint**: seroconversion rate at 28 days after the second injection (NCT04383573).

**Findings**

> Safety: any adverse reaction within 28 days after injection occurred in 20 (20%) of 100 participants in the 1·5 μg group, 25 (20%) of 125 in the 3 μg group, 27 (22%) of 123 in the 6 μg group, and 15 (21%) of 73 in the placebo group.

> All adverse reactions were mild or moderate in severity and injection site pain (39 [9%] of 421 participants) was the most frequently reported event.

> Eight serious adverse events, considered unrelated to vaccination, have been reported by seven (2%) participants.

> In phase 1, seroconversion after the second dose was observed in 24 of 24 participants (100·0% [95% CI 85·8–100·0]) in the 3 μg group and 22 of 23 (95·7%) [78·1–99·9]) in the 6 μg group.

> In phase 2, seroconversion was seen in 88 of 97 participants in the 1·5 μg group (90·7% [83·1–95·7]), 96 of 98 in the 3 μg group (98·0% [92·8–99·8]), and 97 of 98 (99·0% [94·5–100·0]) in the 6 μg group.

**Conclusion**

CoronaVac is safe and well tolerated in older adults. Neutralising antibody titres by the 3 μg dose were similar to those of the 6 μg dose, and higher than those of the 1·5 μg dose.
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<th>Authors and link</th>
<th>Field of expertise</th>
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> The model predicted a median peak viral load that coincided with symptom onset.  
> Patients with age ≥65y had a smaller loss rate of infected cells, leading to a delayed median time to viral clearance occurring 16d after symptom onset as compared to 13 d in younger patients.  
> In multivariate analysis, the risk factors associated with mortality were age ≥65y, male gender, and presence of chronic pulmonary disease (hazard ratio [HR] > 2.0). Using a joint model, viral dynamics after hospital admission was an independent predictor of mortality (HR = 1.31, P < 10−3).  
> Simulation of effectiveness of pharmacological interventions: a treatment able to reduce viral production by 90% upon hospital admission would shorten the time to viral clearance by 2.0 and 2.9d in patients of age <65 y and ≥65y, respectively. Assuming a similar association between viral dynamics and mortality in patients of age ≥65y with risk factors, this could translate into a reduction of mortality from 19 to 14%.  
Viral dynamics is associated with mortality in hospitalized patients. |
| The Lancet 02FEB2021 | Safety and efficacy of an rAd26 and rAd5 vector-based heterologous prime-boost COVID-19 vaccine: an interim analysis of a randomised controlled phase 3 trial in Russia | Logunov D.Y., et al. Russia | Vaccines | Background  
> Sputnik V: heterologous recombinant adenovirus (rAd)-based vaccine.  
> Good safety profile and strong humoral and cellular immune responses (phase 1/2 clinical trials).  
Preliminary results on the efficacy and safety of this vaccine from the interim analysis of this phase 3 trial. (NCT04530396).  
Methods  
> Randomised, double-blind, placebo-controlled, phase 3 trial (25 hospitals and polyclinics in Moscow, Russia).  
> Participants aged at least 18 years, with negative SARS-CoV-2 PCR and IgG and IgM tests, no infectious diseases in the 14 days before enrolment, and no other vaccinations in the 30 days before enrolment.  
> Randomly assigned (3:1) to receive vaccine or placebo (0·5 mL/dose) IM: prime-boost regimen at 21-day interval  
> First dose (rAd26) and the second dose (rAd5), both vectors carrying the gene for the full-length SARS-CoV-2 glycoprotein S.  
Primary outcome: proportion of participants with PCR-confirmed COVID-19 from day 21 after receiving the first dose.  
SAE: assessed in all participants who had received at least one dose at the time of database lock  
Findings  
> 21 977 adults randomly assigned to the vaccine group (n=16 501) or the placebo group (n=5476).  
> 19 866 received two doses of vaccine or placebo and were included in the primary outcome analysis.  
> From 21 days after the first dose of vaccine (the day of dose 2):  
> 16 (0·1%) of 14 964 participants in the vaccine group and 62 (1·3%) of 4902 in the placebo group were confirmed to have COVID-19: vaccine efficacy was 91·6% (95% CI 85·6–95·2).  
> Most reported AEs were grade 1 (7485 [94·0%] of 7966 total events).  
> SAE: 45 (0·3%) of 16 427 participants in the vaccine group and 23 (0·4%) of 5435 participants in the placebo group. None were considered associated with vaccination, with confirmation from the independent data monitoring committee.  
> Four deaths were reported during the study (three [<0·1%] of 16 427 participants in the vaccine group and one [<0·1%] of 5435 participants in the placebo group), none of which were considered related to the vaccine.  
Conclusion: This interim analysis of the phase 3 trial of Gam-COVID-Vac showed 91·6% efficacy against COVID-19 and was well tolerated in a large cohort. |
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<th>Field of expertise</th>
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| Science 02FEB2021 | Age groups that sustain resurging COVID-19 epidemics in the United States | Monod M., et al. UK [gotopaper](#) | Public Health / Epidemiology | > Understanding the age demographics driving transmission and how these affect the loosening of interventions is crucial  

**Methods**  
> Analyze aggregated, age-specific mobility trends from more than 10 million individuals in the US and link these mechanistically to age-specific COVID-19 mortality data  

**Findings**  
> Estimation: as of October 2020, individuals aged 20-49 are the only age groups sustaining resurgent SARS-CoV-2 transmission with reproduction numbers well above one, and that at least 65 of 100 COVID-19 infections originate from individuals aged 20-49 in the US  

Targeting interventions – including transmission-blocking vaccines – to adults aged 20-49 is an important consideration in halting resurgent epidemics and preventing COVID-19-attributable deaths. |

| Cell 02FEB2021 | Maturation and persistence of the anti-SARS-CoV-2 memory B cell response | Sokal A., et al. France [gotopaper](#) | Immunology | Analysis of the longevity and functionality of the anti-SARS-CoV-2 memory B cell response  

**Methods**  
> longitudinal deep profiling of the anti-SARS-CoV-2 memory B cell response in two parallel cohorts of patients with severe and mild COVID-19 (39 total patients)  

> They combined single cell transcriptomics, single cell culture and IgH VDJ sequencing to track and characterize the cellular and molecular phenotype and clonal evolution of spike-specific MBCs clones from early time points after SARS-CoV-2 infection up to 6 months after the initial symptoms  

**Findings**  
> Distinct SARS-CoV-2 spike-specific activated B cell clones fueled an early antibody-secreting cell burst as well as a durable synchronous germinal center response  

> While highly mutated memory B cells, including pre-existing cross-reactive seasonal Betacoronavirus-specific clones, were recruited early in the response, neutralizing SARS-CoV-2 RBD-specific clones accumulated with time and largely contributed to the late remarkably stable memory B-cell pool.  

> Seasonal coronavirus-specific memory B cells contribute an early anti-SARS-CoV2 response  

> Spike-specific memory B cells with a resting phenotype increase up to 6 months  

> Highlighting germinal center maturation, these cells displayed clear accumulation of somatic mutations in their variable region genes over time  

> Longitudinal study reveals a temporal switch to RBD-specific neutralizing memory B cells  

These findings demonstrate that an antigen-driven activation persisted and matured up to 6 months after SARS-CoV-2 infection and may provide long-term protection. |
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| Lancet 02FEB2021 | Azithromycin in patients admitted to hospital with COVID-19 (RECOVERY): a randomised, controlled, open-label, platform trial | RECOVERY Collaborative Group UK gotopaper | Therapeutics | **Aim:** to evaluate the safety and efficacy of azithromycin (500 mg once per day by mouth or intravenously for 10 days or until discharge) in patients admitted to hospital with COVID-19.  
**Primary outcome:** 28-day all-cause mortality  
**Results**  
> Between April 7 and Nov 27, 2020, 7763 were included in the assessment of azithromycin. Mean age was 65·3 years, approx. a third were women. 2582 patients were randomly allocated to receive azithromycin and 5181 to usual care alone.  
> Overall, 561 (22%) patients allocated to azithromycin and 1162 (22%) patients allocated to usual care died within 28 days (rate ratio 0·97).  
> No significant difference was seen in duration of hospital stay (median 10 days vs 11 days) or the proportion of patients discharged from hospital alive within 28 days (rate ratio 1·04).  
> Among those not on invasive mechanical ventilation at baseline, no significant difference was seen in the proportion meeting the composite endpoint of invasive mechanical ventilation or death (risk ratio 0·95).  
In patients admitted to hospital with COVID-19, azithromycin did not improve survival or other prespecified clinical outcomes. |
> We identified 314 patients with COVID-19, with 282 (90%) having at least one contact (753 contacts in total), resulting in 282 clusters.  
> 90 (32%) of 282 clusters had at least one transmission event. The secondary attack rate was 17% (125/753 contacts), with a variation from 12% when the index case had a viral load lower than $1 \times 10^6$ copies per mL to 24% when the index case had a viral load of $1 \times 10^{10}$ copies per mL or higher (adjusted odds ratio per log10 increase in viral load 1·3).  
> Increased risk of transmission was also associated with household contact (3·0) and age of the contact (per year: 1·02, 1·01–1·04).  
> 449 contacts had a positive PCR result at baseline. 28 (6%) of 449 contacts had symptoms at the first visit.  
> Of 421 contacts who were asymptomatic at the first visit, 181 (43%) developed symptomatic COVID-19, with a variation from approx. 38% in contacts with an initial viral load lower than $1 \times 10^7$ copies per mL to >66% for those with an initial viral load of $1 \times 10^{10}$ copies per mL or higher (hazard ratio per log10 increase in viral load 1·12).  
> **Time to onset of symptomatic disease decreased** from a median of 7 days (IQR 5–10) for individuals with an initial viral load lower than $1 \times 10^7$ copies per mL to 6 days (4–8) for those with an initial viral load between $1 \times 10^7$ and $1 \times 10^9$ copies per mL, and 5 days (3–8) for those with an initial viral load higher than $1 \times 10^9$ copies per mL.  
The viral load of index cases was a leading driver of SARS-CoV-2 transmission. The risk of symptomatic COVID-19 was strongly associated with the viral load of contacts at baseline and shortened the incubation time of COVID-19 in a dose-dependent manner. |
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<th>Field of expertise</th>
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| Nature 01FEB2021 | Immunogenic BNT162b vaccines protect rhesus macaques from SARS-CoV-2 | Vogel A.B., et al. Germany [gotopaper](#) | Immunology / Preclinical model | > Preclinical development of two BNT162b vaccine candidates: lipid-nanoparticle (LNP) formulated nucleoside-modified mRNA encoding SARS-CoV-2 spike glycoprotein-derived immunogens  
> BNT162b1 encodes a soluble, secreted, trimerised receptor-binding domain (RBD-foldon)  
> BNT162b2 encodes the full-length transmembrane spike glycoprotein, locked in its prefusion conformation (PS2)  
> flexibly tethered RBDs of the RBD-foldon bind ACE2 with high avidity  
> Approximately 20% of the P25 trimers are in the two-RBD ‘down,’ one-RBD ‘up’ state  

Findings  
> In mice, one intramuscular dose of either candidate elicits a dose-dependent antibody response with high virus-entry inhibition titres and strong TH1 CD4+ and IFNγ+ CD8+ T-cell responses  
> Prime/boost vaccination of rhesus macaques with BNT162b candidates elicits SARS-CoV-2 neutralising geometric mean titres 8.2 to 18.2 times that of a SARS-CoV-2 convalescent human serum panel  
> Vaccine candidates protect macaques from SARS-CoV-2 challenge, with BNT162b2 protecting the lower respiratory tract from the presence of viral RNA and with no evidence of disease enhancement |
> Hospitalized patients with laboratory-confirmed COVID-19 from 2 Italian tertiary referral centres (derivation cohort, n = 187 patients; validation cohort, n = 62 patients).  
> Three-day angiopoietin-2 increase of at least twofold from baseline was significantly associated with in-hospital mortality by multivariate analysis (hazard ratio [HR], 6.69) with Area under the receiver operating characteristic curve (AUROC) = 0.845.  
> Ten-day angiopoietin-2 of at least twofold from baseline was instead significantly associated with nonresolving pulmonary condition by multivariate analysis (HR, 5.33) with AUROC = 0.969.  
> Patients with persistent elevation of 10-day angiopoietin-2 levels showed severe reticular interstitial thickening and fibrous changes on follow-up computed tomography scans. Angiopoietin-2 and Tie2 were diffusely colocalized in small-vessel endothelia and alveolar new vessels and macrophages.  

Angiopoietin-2 course is strongly associated with COVID-19 in-hospital mortality and nonresolving pulmonary condition, and may be an early and useful predictor of COVID-19 clinical course.  

Description of disease phenotypes of SARS-CoV-2 exposure occurring around the time of vaccine administration  
- Disease phenotypes of a one-dose regimen given 3 days prior (D-3), 1 (D1) or 2 (D2) days after, or on the day (D0) of virus challenge in golden Syrian hamster  
- Monitoring of serial clinical severity, tissue histopathology, virus burden, and antibody response of the vaccinated hamsters.  

Findings:  
> One-dose vaccinated hamsters had significantly lower clinical disease severity score, body weight loss, lung histology score, nucleocapsid protein expression in lung, infectious virus titres in the lung and nasal turbinate, inflammatory changes in intestines and a higher serum neutralizing antibody or IgG titre against the spike receptor-binding domain or nucleocapsid protein when compared to unvaccinated controls.  
> Improvements particularly noticeable in D-3, but also in D0, D1 and even D2 vaccinated hamsters to varying degrees.  
> No increased eosinophilic infiltration was found in the nasal turbinate, lung, and intestine after virus challenge.  
> Significantly higher serum titre of fluorescent foci microneutralization inhibition antibody was detected in D1 and D2 vaccinated hamsters at day 4 post-challenge compared to controls despite undetectable neutralizing antibody titre.  

Vaccination just before or soon after exposure to SARS-CoV-2 does not worsen disease phenotypes and may even ameliorate infection. |
| Clin Infect Dis 30JAN21 | Absence of vaccine-enhanced disease with unexpected positive protection against SARS-CoV-2 by inactivated vaccine given within three days of virus challenge in Syrian hamster model | Li C., et al. China [gotopaper](#) | Vaccines (viral mutants) |  

- Disease phenotypes of a one-dose regimen given 3 days prior (D-3), 1 (D1) or 2 (D2) days after, or on the day (D0) of virus challenge in golden Syrian hamster  
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| Science 29JAN2021 | Neutralization of SARS-CoV-2 lineage B.1.1.7 pseudovirus by BNT162b2 vaccine–elicited human sera | Muik M., et al. Germany/USA [gotopaper](#) | Vaccines (viral mutants) | **Background:**
> The new SARS-CoV-2 lineage called B.1.1.7 emerged in the UK and is reported to spread more efficiently and faster than other strains.
> This variant contains 10 amino acid changes in the spike protein: ΔH69/V70, ΔY144, N501Y, A570D, D614G, P681H, T716I, S982A, D1118H.
> N501Y mutation is located in the receptor binding site. The spike with this mutation binds more tightly to its cellular receptor ACE-2.

**Is this virus strain recognized by neutralizing antibodies induced after vaccination?**

**Methods:**
> VSV SARS-CoV-2-S pseudoviruses bearing the Wuhan reference strain or the B.1.1.7 lineage spike protein tested with sera of 40 participants given the BNT162b2 vaccine from Pfizer (phase I/II, DE)
> The 50% neutralization geometric mean titer (GMT) of sera against the SARS-CoV-2 lineage B.1.1.7 spike pseudotyped VSV for the younger adult group and the full analysis set were slightly, statistically significantly reduced compared to the GMTs against the Wuhan reference spike pseudotyped VSV.
> GMTs were not significantly different for the older adult group (0.78 [0.68;0.89] for the younger and 0.83 [0.65;1.1] for the older adults (0.80 [0.71;0.89] CI 95%).

**Conclusions:**
> Based on experience from antibody correlates of disease protection for influenza virus vaccines, a 20% reduced titer does not indicate a biologically significant change in neutralization activity.
> The largely preserved neutralization of pseudoviruses bearing the B.1.1.7 spike by BNT162b2-immune sera makes it unlikely that the UK variant virus will escape BNT162b2-mediated protection. |
> Understanding the dynamics of maternal antibody (Ab) responses to SARS-CoV-2 infection during pregnancy and transplacental Ab transfer.
> Assessing association between maternal and neonatal SARS-CoV-2-specific Ab concentrations.

**Background:**
> Maternal and cord blood sera were available for Ab measurement for 1471 mother/newborn dyads (09Apr-08Aug 2020).
> IgG and IgM to the receptor-binding domain of the SARS-CoV-2 spike protein were measured by enzyme-linked immunosorbent assay.
> Ab concentrations and transplacental transfer ratios were analyzed in combination with demographic and clinical data.

**Findings:**
> SARS-CoV-2 IgG Ab were transferred across the placenta in 72 of 83 pregnant women who were seropositive.
> Cord blood IgG concentrations were directly associated with maternal Ab concentrations.
> IgM antibodies were not detected in any cord blood sera.
> Transfer ratios were associated with time elapsed from maternal infection to delivery and not associated with severity of maternal infection.

**Efficient transplacental transfer of SARS-CoV-2 IgG Ab supports potential maternal Ab neonate protection from SARS-CoV-2 infection.** |
| Cell 28JAN2021 | Circulating SARS-CoV-2 spike N439K variants maintain fitness while evading antibody-mediated immunity | Thomson E.C., et al. UK/USA [gotopaper](#) | Virology | **Background:**
> The immunodominant SARS-CoV-2 spike (S) receptor binding motif (RBM) is a highly variable region of S, and provide epidemiological, clinical, and molecular characterization of a prevalent, sentinel RBM mutation, N439K.
> N439K S protein has enhanced binding affinity to the hACE2 receptor, and N439K viruses have similar in vitro replication fitness and cause infections with similar clinical outcomes to wild-type.
> The N439K mutation confers resistance against several neutralizing monoclonal antibodies, including one authorized for emergency use by the FDA, and reduces the activity of some polyclonal sera from persons recovered from infection.

Immune evasion mutations that maintain virulence and fitness such as N439K can emerge within SARS-CoV-2 S, highlighting the need for ongoing molecular surveillance. |
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<th>Authors and link</th>
<th>Field of expertise</th>
<th>Key facts</th>
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| Nature Commun. 27JAN2021 | Integrating deep learning CT-scan model, biological and clinical variables to predict severity of COVID-19 patients | Lassau N., et al. France gotopaper | Diagnostics       | Identifying predictors of disease severity is a priority  
> Collect 58 clinical and biological variables, and chest CT scan data, from 1003 coronavirus-infected patients from two French hospitals.  
> Train a deep learning model based on CT scans to predict severity  
> Construct the multimodal AI-severity score that includes 5 clinical and biological variables (age, sex, oxy-genation, urea, platelet) in addition to the deep learning model  

**Findings**  
Neural network analysis of CT-scans brings unique prognosis information, although it is correlated with other markers of severity (oxygenation, LDH, and CRP) explaining the measurable but limited 0.03 increase of AUC obtained when adding CT-scan information to clinical variables.  
When comparing AI-severity with 11 existing severity scores, we find significantly improved prognosis performance; AI-severity can therefore rapidly become a reference scoring approach. |
| JAMA Netw Open 27JAN2021 | Respiratory and Psychophysical Sequelae Among Patients With COVID-19 Four Months After Hospital Discharge | Bellan M., et al Italy gotopaper | Public Health / Epidemiology - Long Covid | **Aim:** Evaluate the prevalence of lung function anomalies, exercise function impairment, and psychological sequelae among patients hospitalized for COVID-19, 4 months after discharge  

**Methods**  
> Prospective cohort study at an academic hospital  
> Patients ≥18 years old (or their caregivers) hospitalized with SARS-CoV-2 infection (March 1-June 29, 2020)  
> Confirmed via RT-PCR testing, bronchial swab, serological testing, or suggestive computed tomography results  

To describe proportion of patients with:  
> Diffusing lung capacity for carbon monoxide (DLCO) <80% of expected value  
> Severe lung function impairment (DLCO <60% expected value)  
> Posttraumatic stress symptoms (measured using the Impact of Event Scale–Revised total score)  
> Functional impairment (assessed using the Short Physical Performance Battery [SPPB] score and 2-minute walking test);  
> Identification of factors associated with DLCO reduction and psychological functional sequelae  

**Findings**  
> 238/767 patients (31.0%) (median age, 61 [50-71] years; 142 [59.7%] men; median comorbidities, 2 [1-3]) had sequelae.  
> 219 patients were able to complete both pulmonary function tests and DLCO measurement. DLCO was reduced to <80% of the estimated value in 113 patients (51.6%) and <60% in 34 patients (15.5%)  
> The SPPB score was suggested limited mobility (score <11) in 53 patients (22.3%).  

Patients with normal SPPB scores underwent a 2-minute walk test, which was outside reference ranges of expected performance for age and sex in 75 patients (40.5%) → 128 patients (53.8%) had functional impairment. Posttraumatic stress symptoms were reported in a total of 41 patients (17.2%)  

4 months after discharge, respiratory, physical, and psychological sequelae were common among patients who had been hospitalized for COVID-19. |
| Cell 26JAN2021 | Two-component spike nanoparticle vaccine protects macaques from SARS-CoV-2 infection | Brouwer P.J.M., et al. The Netherlands gotopaper | Vaccines | **> Two-component protein nanoparticles display multiple copies of the SARS-CoV-2 Spike protein potentially protecting from infection**  

**Immunization studies :**  
> Vaccination induces potent neutralizing antibody responses in mice, rabbits and cynomolgus macaques  
> Spike protein nanoparticles enhance cognate B cell activation in vitro  
> Vaccination protects macaques against a high-dose SARS-CoV-2 challenge, resulting in strongly reduced viral infection and replication in upper and lower airways.  

These nanoparticles are a promising vaccine candidate. |
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**Methods:** whole-blood preserving single-cell analysis to integrate contributions from all major cell types including neutrophils, monocytes, platelets, lymphocytes and the contents of serum.  
**Findings:** > Patients with mild COVID-19 disease display a coordinated pattern of interferon-stimulated gene (ISG) expression across every cell population and these cells are systemically absent in patients with severe disease  
> Severe COVID-19 patients paradoxically produce very high anti-SARS-CoV-2 antibody titers and have lower viral load as compared to mild disease eve two weeks beyond symptom onset.  
> Examination of the serum from severe patients demonstrates that they uniquely produce Abs that functionally block the production of the mild disease-associated ISG-expressing cells, by engaging conserved signaling circuits that dampen cellular responses to interferons  
Global targeting of ISG archetypes might be addressable with drugs such as rituximab to reduce B cell responses, perhaps in the presence of convalescent serum, through introduction of IVIG to compete with serum antibodies for FcR engagement, or with rapid development of antibodies that clinically block FCyRIIb. |
| Science 25JAN2021 | Prospective mapping of viral mutations that escape antibodies used to treat COVID-19 | Starr T.N., et al. USA gotopaper | Immunology | **Aim:** mapping how all mutations to SARS-CoV-2’s receptor-binding domain (RBD) affect binding by the antibodies in the REGN-COV2 cocktail and the antibody LY-CoV016.  
**Methods:** To validate the antigenic effects of key mutations, neutralization assays using spike-pseudotyped lentiviral particles were made.  
**Findings:** > Regarding REGN-COV2 antibodies: a mutation at site 486 escaped neutralization only by REGN10933, whereas mutations at sites 439 and 444 escaped neutralization only by REGN10987  
> One mutation (E406W) strongly escapes the cocktail of both antibodies  
> E406W is not accessible by a single-nucleotide change, which may explain why it was not identified by the Regeneron cocktail  
> Mutations at RBD residues that contact antibody do not always mediate escape, and several prominent escape mutations occur at residues not in contact with antibody.  
> The maps reveal that mutations escaping the individual antibodies are already present in circulating SARS-CoV-2 strains. |
| Science 25JAN21 | Plitidepsin has potent preclinical efficacy against SARS-CoV-2 by targeting the host protein eEF1A | White K.M., et al. International gotopaper | Therapeutics | Previous author’s work on SARS-CoV-2 highlighted 332 host proteins that are likely to play a role in the viral life cycle of SARS-CoV-2. Drugs modulating these host proteins were tested and those that targeted the eukaryotic translation machinery (eIF4H interacts with SARS-CoV-2 Nsp9) demonstrated particularly potent antiviral activities.  
In this study, the eEF1A inhibitor plitidepsin was tested.  
Plitidepsin has been clinically developed for the treatment of multiple myeloma with a well-established safety profile and pharmacokinetics.  
**Findings:** > Antiviral activity (IC90 = 0.88 nM) 27.5-fold more potent than remdesivir against SARS-CoV-2 in vitro, limiting toxicity  
> The dynamics between the antiviral effects of plitidepsin and remdesivir when used together in vitro suggests that plitidepsin has an additive effect with remdesivir  
> The antiviral activity of plitidepsin against SARS-CoV-2 is mediated through inhibition of the known target eEF1A.  
> In vivo studies in mouse models of SARS-CoV-2 infection showed a reduction of viral replication in the lungs by two orders of magnitude when using Plitidepsin in prophylactic treatment.  
**Conclusions:** This study establishes plitidepsin as a host-targeted anti-SARS-CoV-2 agent with in vivo efficacy. *Phase II/III study to come* |
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<td>&gt; Among the 222 patients included, 174 (78%) were on microbiotics.</td>
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<td>&gt; Univariate analysis: patients with antibiotics were significantly older, frailer and with a more severe presentation at admission.</td>
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<td>&gt; An unfavorable outcome was more frequent in patients with antibiotic therapy (HR = 2.94).</td>
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<td>&gt; In multivariate analysis and propensity score, antibiotic therapy was not significantly associated with outcome (HR = 1.612).</td>
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<td>Antibiotics were frequently prescribed in our study and associated with a more severe presentation at admission. However, receiving antibiotics was not associated with outcome.</td>
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<td>- Usual care + anakinra (200 mg twice a day on days 1–3, 100 mg twice on day 4, 100 mg once on day 5) vs usual care only.</td>
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<td>Two coprimary outcomes: proportion of patients who had died or needed non-invasive or mechanical ventilation by day 4 (ie, a score of &gt;5 on the WHO-CPS) and survival without need for mechanical or non-invasive ventilation (including high-flow oxygen) at day 14.</td>
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<td>Results</td>
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<td>&gt; 116 patients recruited: 59 in the anakinra group, and 57 in the usual care group (2 withdrew). Median age was 66 years, 70% were men.</td>
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<td>&gt; In the anakinra group, 21/59 (36%) patients had a WHO-CPS score &gt;5 at day 4 versus 21/55 (38%) in the usual care group (median posterior absolute risk difference [ARD] −2.5%), with a posterior probability of ARD of less than 0 (ie, anakinra better than usual care) of 61.2%.</td>
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<td>&gt; At day 14, 28 (47%) patients in the anakinra group and 28 (51%) in the usual care group needed ventilation or died, with a posterior probability of any efficacy of anakinra (hazard ratio [HR] &lt;1) of 54.5% (median posterior HR 0.97).</td>
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<td>&gt; At day 90, 16 (27%) patients in the anakinra group and 15 (27%) in the usual care group had died. Serious adverse events occurred in 27 (46%) patients in the anakinra group and 21 (38%) in the usual care group (p=0.45).</td>
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<td>JAMA</td>
<td>Effect of Bamlanivimab as Monotherapy or in Combination With Etesevimab on Viral Load in Patients With Mild to Moderate COVID-19</td>
<td>Gottlieb R.L., et al. USA gotopaper</td>
<td>Theraupetics</td>
<td>Aim: to determine the effect of bamlanivimab monotherapy and combination therapy with bamlanivimab and etesevimab on SARS-CoV-2 viral load in mild to moderate COVID-19 (BLAZE-1 study).</td>
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<td>- Bamlanivimab: a single infusion of 700 mg (n = 101), 2800 mg (n = 107), or 7000 mg (n = 101)</td>
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<td>- Combination treatment: 2800 mg of bamlanivimab and 2800 mg of etesevimab [n = 112]</td>
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<td>- Placebo (n = 156). Primary end point: change in SARS-CoV-2 log viral load at D11 (±4 dys).</td>
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<td>&gt; Among the 577 randomized (mean age, 44.7 years; 54.6% women), 533 (92.4%) completed the efficacy evaluation period (day 29).</td>
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<td>&gt; Change in log viral load from baseline at D11 was −3.72 for 700 mg, −4.08 for 2800 mg, −3.49 for 7000 mg, −4.37 for combination treatment, and −3.80 for placebo. Compared with placebo, differences in the change in log viral load at D11 were 0.09 for 700 mg, −0.27 for 2800 mg, 0.31 for 7000 mg, and −0.57 for combination treatment.</td>
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<td>&gt; Among the secondary outcome measures, differences between each treatment group vs the placebo group were statistically significant for 10 of 84 end points. The proportion of patients with COVID-19–related hospitalizations or ED visits was 5.8% (9 events) for placebo, 1.0% (1 event) for 700 mg, 1.9% (2 events) for 2800 mg, 2.0% (2 events) for 7000 mg, and 0.9% (1 event) for combination treatment.</td>
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<td>&gt; Immediate hypersensitivity reactions were reported in 9 patients (6 bamlanivimab, 2 combination treatment, and 1 placebo).</td>
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<td>&gt; No deaths occurred during the study treatment.</td>
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<td>In nonhospitalized patients with mild to moderate COVID-19, bamlanivimab and etesevimab treatment, compared with placebo, was associated with a reduction in SARS-CoV-2 viral load at day 11.</td>
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### Lancet Infect Dis 21JAN21

**Safety and immunogenicity of an inactivated SARS-CoV-2 vaccine, BBV152: a double-blind, randomised, phase 1 trial**

**Authors and link**

Ella R., et al. India

gotopaper

**Field of expertise**

Vaccines

**Key facts**

**Background**

BBV152: whole-virion inactivated SARS-CoV-2 vaccine formulated with a toll-like receptor 7/8 agonist molecule adsorbed to alum (Algel-IMDG) or alum (Algel).

**Methods**

- Double-blind, multicentre, randomised, controlled phase 1 trial to assess the safety and immunogenicity of BBV152 at 11 hospitals across India. (NCT04471519).
- Healthy adults aged 18–55 years Individuals with positive SARS-CoV-2 nucleic acid and/or serology tests excluded.
- Participants randomly assigned to receive either one of three vaccine formulations: 3 μg with Algel-IMDG / 6 μg with Algel-IMDG / 6 μg with Algel / Algel only
- Two IM doses at d0 et d14
- Primary outcomes: solicited local and systemic reactogenicity events at 2 h and 7 days after vaccination
- Secondary outcome: seroconversion
- Cell-mediated responses were evaluated by intracellular staining and ELISPOT.

**Findings**

- 375 participants enrolled: 100 each were randomly assigned to the three vaccine groups, and 75 were randomly assigned to the control group (Algel only).
- Solicited local and systemic adverse reactions after 2 doses: 17 (17%; 95% CI 10.5–26.1) participants in the 3 μg with Algel-IMDG group, 21 (21%; 13.8–30.5) in the 6 μg with Algel-IMDG group, 14 (14%; 8.1–22.7) in the 6 μg with Algel group, and ten (10%; 6.9–23.6) in the Algel-only group.
- Most common solicited adverse events: injection site pain [17 (5%) of 375 participants], headache [13 (3%)], fatigue [11 (3%)], fever [nine (2%)], and nausea or vomiting (seven [2%]). All solicited adverse events were mild or moderate, and more frequent after the first dose.
- One SAE (viral pneumonitis) reported in the 6 μg with Algel group, unrelated to the vaccine.
- Seroconversion rates (%) of 87.9, 91.9, and 82.8 in the 3 μg, 6 μg with Algel-IMDG, and 6 μg with Algel groups, respectively.
- CD4+ and CD8+ T-cell responses were detected in a subset of 16 participants from both Algel-IMDG groups.

**BBV152 led to tolerable safety outcomes and enhanced immune responses. Both Algel-IMDG formulations were selected for phase 2 immunogenicity trials.**

### Science Immunol. 21JAN2021

**Severely ill COVID-19 patients display impaired exhaustion features in SARS-CoV-2-reactive CD8+ T cells**

**Authors and link**

Kusnadi A., et al. USA

gotopaper

**Field of expertise**

Immunology

**Key facts**

**Aim:** Understand anti-viral immune responses. Report from data generated by single-cell RNA sequencing of virus-reactive CD8+ T cells from COVID-19 patients with different clinical severity.

**Methods:** Single-cell transcriptomes of >80,000 virus-reactive CD8+ T cells, obtained using a modified Antigen-Reactant T cell Enrichment (ARTE) assay, from 39 COVID-19 patients and 10 healthy subjects.
- Recent reports from COVID-19 patients have suggested the presence of exhaustion-related markers in global CD8+ T cell populations.
- COVID-19 patients were segregated into two groups based on whether the dominant CD8+ T cell response to SARS-CoV-2 was "exhausted" or not.

**Findings:**

- SARS-CoV-2-reactive cells in the exhausted subset were increased in frequency and displayed lesser cytotoxicity and inflammatory features in COVID-19 patients with mild compared to severe illness.
- SARS-CoV-2-reactive cells in the dominant non-exhausted subset from patients with severe disease showed enrichment of transcripts linked to co-stimulation, pro-survival NF-κB signaling, and anti-apoptotic pathways, suggesting the generation of robust CD8+ T cell memory responses in patients with severe COVID-19 illness.

**Overall, the single-cell analysis revealed substantial diversity in the nature of CD8+ T cells responding to SARS-CoV-2.**
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| Lancet Public Health 20JAN2021 | Quarantine and testing strategies in contact tracing for SARS-CoV-2: a modelling study | Quilty B.J., et al. UK gotopaper | Public Health / Epidemiology | **Aim:** to assess the merit of testing contacts to avert onward transmission and to replace or reduce the length of quarantine for uninfected contacts.  
> Assuming moderate levels of adherence to quarantine and self-isolation, self-isolation on symptom onset alone can prevent 37% of onward transmission potential from secondary cases.  
> 14 days of post-exposure quarantine reduces transmission by 59%.  
> Quarantine with release after a negative PCR test 7 days after exposure might avert a similar proportion (54%; risk ratio [RR] 0.94), as would quarantine with a negative lateral flow antigen test 7 days after exposure (50%; RR 0.88) or daily testing without quarantine for 5 days after tracing (50%; RR 0.88) if all tests are returned negative.  
**Testing might allow for a substantial reduction in the length of, or replacement of, quarantine with a small excess in transmission risk.** Decreasing test and trace delays and increasing adherence will further increase the effectiveness of these strategies. |
| BMJ 20JAN21 | Effect of tocilizumab on clinical outcomes at 15 days in patients with severe or critical coronavirus disease 2019: randomised controlled trial | V.C. Veiga, et al. Brazil gotopaper | Therapeutics | Does tocilizumab improves clinical outcomes for patients with severe or COVID-19?.  
**Methods:**  
> Randomised, open label trial (NCT04403685)  
> Nine hospitals in Brazil, 8 May to 17 July 2020.  
> Adults with confirmed Covid-19 who were receiving supplemental oxygen or mechanical ventilation and had abnormal levels of at least two serum biomarkers (C reactive protein, D dimer, lactate dehydrogenase, or ferritin).  
> Interventions Tocilizumab (single intravenous infusion of 8 mg/kg) plus standard care (n=65) versus standard care alone (n=64).  
> Main outcome: clinical status measured at 15 days, analysed as a composite of death or mechanical ventilation (assumption of odds proportionality was not met).  
> *The data monitoring committee recommended stopping the trial early, after 129 patients had been enrolled, because of an increased number of deaths at 15 days in the tocilizumab group.*  
**Findings:**  
> 129 patients enrolled (mean age 57 years; 68% men) and all completed follow-up.  
> All patients in the tocilizumab group and two in the standard care group received tocilizumab.  
> 18 of 65 (28%) patients in the tocilizumab group and 13 of 64 (20%) in the standard care group were receiving mechanical ventilation or died at day 15 (odds ratio 1.54).  
> Death at 15 days occurred in 11 (17%) patients in the tocilizumab group compared with 2 (3%) in the standard care group (odds ratio 6.42).  
> Adverse events were reported in 29 of 67 (43%) patients who received tocilizumab and 21 of 62 (34%) who did not receive tocilizumab.  
In patients with severe or critical Covid-19, tocilizumab plus standard care was not superior to standard care alone in improving clinical outcomes at 15 days, and it might increase mortality. |
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| Lancet Microbe 19JAN2021 | Insight into the practical performance of RT-PCR testing for SARS-CoV-2 using serological data: a cohort study | Zhang Z., et al. China [gotopaper](#) | Diagnostics | **Aim:** Assess the practical performance of RT-PCR-based surveillance protocols and determine the extent of undetected SARS-CoV-2 infection in Shenzhen, China.  
**Methods:** cohort study in Shenzhen, China. All RT-PCR(-) close contacts (defined as those who lived in the same residence as, or shared a meal, travelled, or socially interacted with, an index case within 2 days before symptom onset) of all RT-PCR(+) cases of SARS-CoV-2 detected since January, 2020.  
**Findings:**  
> Serological samples from 2345 of 4422 RT-PCR (-) close contacts of cases of RT-PCR-confirmed SARS-CoV-2.  
> 80 of 880 RT-PCR (-) close contacts were positive on total antibody ELISA.  
> The seropositivity rate with total Ab ELISA among RT-PCR (-) close contacts, adjusted for assay performance, was 4.1%, which was significantly higher than among individuals residing in neighbourhoods with no reported cases  
> RT-PCR (+) individuals were 8.0 times more likely to report symptoms than those who were RT-PCR (-) but seropositive.  
> RT-PCR did not detect 48 of 134 infected close contacts, and false-negative rates appeared to be associated with stage of infection.  
Even rigorous RT-PCR testing protocols might miss a substantial proportion of SARS-CoV-2 infections, perhaps in part due to difficulties in determining the timing of testing in asymptomatic individuals for optimal sensitivity. |
**Primary outcome:** all-cause 28-day in-hospital mortality.  
**Secondary outcomes:** all-cause death at any time, receipt of mechanical ventilation (MV), readmissions.  
> Among 468 patients with COVID-19–related critical illness, 319 (68.2%) were treated with MV and 121 (25.9%) with vasopressors.  
> All-cause 28-day in-hospital mortality rate was 29.9%, median ICU stay was 8 days (IQR, 3–17), median hospital stay was 13 days (IQR, 7–25), and all-cause 30-day readmission rate (among nonhospice survivors) was 10.8%.  
> Mortality decreased over time, from 43.5% (CI, 31.3–53.8) to 19.2% (CI, 11.6–26.7) between the first and last 15-day periods in the core adjusted model, whereas patient acuity and other factors did not change.  
> Among patients with COVID-19–related critical illness admitted to ICUs, mortality seemed to decrease over time despite stable patient characteristics. |
| Nature 18JAN2021 | Evolution of antibody immunity to SARS-CoV-2 | Gaebler C., et al. USA [gotopaper](#) | Immunology | **Aim:** Assess the humoral memory response in a cohort of 87 individuals assessed at 1.3 and 6.2 months after infection.  
**Findings:**  
> IgM, and IgG anti-SARS-CoV-2 spike protein receptor binding domain (RBD) antibody titres decrease significantly with IgA being less affected.  
> The number of RBD-specific memory B cells is unchanged. Memory B cells display clonal turnover after 6.2 months, and the antibodies they express have greater somatic hypermutation, increased potency and resistance to RBD mutations.  
> Analysis of intestinal biopsies obtained from asymptomatic individuals, revealed persistence of SARS-CoV-2 nucleic acids and immunoreactivity in the small bowel (7/14 volunteers).  
> The memory B cell response to SARS-CoV-2 evolves between 1.3 and 6.2 months after infection in a manner that is consistent with antigen persistence.  
> Individuals who are infected with SARS-CoV-2 could mount a rapid and effective response to the virus upon re-exposure. |
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> 385 references, 16 unique studies (5922 unique patients). Significant variability in patient selection, study design, setting and stage of illness at which patients were enrolled.  

> In the primary analysis, the **saliva NAAT** pooled sensitivity was 83.2% (95% credible interval [CrI], 74.7%-91.4%) and the pooled specificity was 99.2% (95% CrI, 98.2%-99.8%).  

> The **nasopharyngeal swab** NAAT had a sensitivity of 84.8% (95% CrI, 76.8%-92.4%) and a specificity of 98.9% (95% CrI, 97.4%-99.8%).  

> Results were similar in secondary analyses (on peer-reviewed studies, and on ambulatory settings).  

**Saliva NAAT diagnostic accuracy is similar to that of nasopharyngeal swab NAAT,** especially in the ambulatory setting, supporting larger-scale research on the use of saliva NAAT as an alternative. |

Methods:  

> Multicenter, placebo-controlled, phase 1–2a trial, randomised  

> Healthy adults: between the ages of 18 and 55 years (cohort 1) and those 65 years of age or older (cohort 3) (=> 805 participants)  

> Cohorte 1& 3 : receive the Ad26.COV2.S vaccine at a dose of 5x1010viral particles (low dose) or 1x1011 viral particles (high dose) per milliliter or placebo in a single-dose or two-dose schedule  

> Cohorte 2: Longer-term data comparing a single-dose regimen with a two-dose regimen are being collected  

**Findings related to safety & reactogenicity**  

After first vaccine dose in cohorts 1 & 3 and after second dose in cohort 1:  

> Most frequent solicited adverse events (AE) were fatigue, headache, myalgia, and injection-site pain & most frequent systemic AE = fever  

> Systemic adverse events were less common in cohort 3 than in cohort 1 and in those who received the low vaccine dose than in those who received the high dose.  

> Reactogenicity was lower after the second dose.  

**Findings related to immunogenicity profiles**  

> Neutralizing-antibody titers against wild-type virus were detected in 90% or more of all participants on day 29 after the first vaccine dose, and reached 100% by day 57 with a further increase in titers in cohort 1a.  

> Titers remained stable until at least day 71. A second dose provided an increase in the titer by a factor of 2.6 to 2.9 (GMT, 827 to 1266). Spike-binding antibody responses were similar to neutralizing-antibody responses.  

> On day 14, CD4+ T-cell responses were detected in 76 to 83% of the participants in cohort 1 and in 60 to 67% of those in cohort 3, with a clear skewing toward type 1 helper T cells. CD8+ T-cell responses were robust overall but lower in cohort 3.  

**The safety and immunogenicity profiles of Ad26.COV2.S support further development of this vaccine candidate.** |
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<td>NEJM 13JAN2021</td>
<td>Convalescent Plasma Antibody Levels and the Risk of Death from Covid-19</td>
<td>Joyner M.J., et al. USA gotopaper</td>
<td>Therapeutics</td>
<td><strong>Aim:</strong> to assess whether convalescent plasma with high antibody levels rather than low antibody levels is associated with a lower risk of death. <strong>Primary outcome:</strong> death within 30 days after plasma transfusion.</td>
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<td>Science 12JAN2021</td>
<td>Mosaic nanoparticles elicit cross-reactive immune responses to zoonotic coronaviruses in mice</td>
<td>Cohen A.A., et al. USA gotopaper</td>
<td>Immunology</td>
<td>Construction of homotypic nanoparticles displaying the RBD of SARS-CoV-2 or co-displaying SARS-CoV-2 RBD along with RBDs from animal betacoronaviruses (mosaic nanoparticles; 4-8 distinct RBDs).</td>
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| Nature Commun. 11JAN2021 | Duration and key determinants of infectious virus shedding in hospitalized patients with coronavirus disease-2019 (COVID-19) | van Kampen J.I.A., et al. Netherlands gotopaper | Public Health / Epidemiology | **Aim:** assess the duration and key determinants of infectious SARS-CoV-2 shedding in patients with severe and critical COVID-19 **Methods:** Analysis of viral loads, neutralizing antibody titers (nAb), detection of the subgenomic RNAs from 129 hospitalized individuals diagnosed with COVID-19 by RT-PCR **Findings:** >Infectious virus shedding was detected by virus cultures in 23/129 patients (17.8%) hospitalized with COVID-19 >The median duration of shedding infectious virus is 8 days post onset of symptoms and drops below 5% after 15.2 days post onset of symptoms. >The probability of isolating infectious virus was less than 5% when the nAb titer was 1:80 or higher. >A serum nAb titre of at least 1:20 (OR of 0.01) is independently associated with non-infectious SARS-CoV-2. >Quantitative viral RNA load assays and serological assays could be used in test-based strategies to discontinue or de-escalate infection prevention and control precautions. >Detection of viral subgenomic RNA correlated poorly with shedding of infectious virus.
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| The Lancet 08JAN2021 | 6-month consequences of COVID-19 in patients discharged from hospital: a cohort study | Huang C.H., et al. China gotopaper | Public Health / Epidemiology - Long COVID | **Aim:** to describe the long-term health consequences of patients with COVID-19 who have been discharged from hospital and investigate the associated risk factors.  
> 1733 discharged patients with COVID-19 enrolled: median age of 57 years and 52% were men. The median follow-up time after symptom onset was 186·0 days.  
> **Fatigue or muscle weakness** (63%, 1038/1655) and **sleep difficulties** (26%, 437/1655) were the most common symptoms. **Anxiety or depression** was reported among 23% (367/1617) of patients.  
> The proportions of median 6-min walking distance less than the lower limit of the normal range were 24% for those at severity scale 3, 22% for severity scale 4, and 29% for severity scale 5–6.  
> The corresponding proportions of patients with **diffusion impairment** were 22% for severity scale 3, 29% for scale 4, and 56% for scale 5–6, and median CT scores were 3·0 for severity scale 3, 4·0 for scale 4, and 5·0 for scale 5–6.  
> After multivariable adjustment, patients showed an odds ratio (OR) 1·61 for scale 4 versus scale 3 and 4·60 for scale 5–6 versus scale 3 for diffusion impairment; OR 0·88 for scale 4 versus scale 3 and OR 1·77 for scale 5–6 versus scale 3 for anxiety or depression, and OR 0·74 for scale 4 versus scale 3 and 2·69 for scale 5–6 versus scale 3 for fatigue or muscle weakness.  
> Of 94 patients with **blood antibodies** tested at follow-up, the seropositivity (96·2% vs 58·5%) and median titres (19·0 vs 10·0) of the neutralising antibodies were significantly lower compared with at the acute phase.  
> 107 of 822 participants without acute kidney injury and with estimated glomerular filtration rate (eGFR) 90 mL/min per 1·73 m² or more at acute phase had eGFR less than 90 mL/min per 1·73 m² at follow-up.  
> At 6 months after acute infection, COVID-19 survivors were mainly troubled with fatigue or muscle weakness, sleep difficulties, and anxiety or depression. Patients who were more severely ill during their hospital stay had more severe impaired pulmonary diffusion capacities and abnormal chest imaging manifestations. |
| JAMA Netw. Open 07JAN2021 | SARS-CoV-2 Transmission From People Without COVID-19 Symptoms | Johansson M.A., et al. USA gotopaper | Public Health/Epidemiology | **Aim:** to assess the proportion of SARS-CoV-2 transmissions in the community that likely occur from persons without symptoms.  
Baseline assumptions for the model: incubation period at 5 days, infectious period of 10 days, peak infectiousness occurred at the median of symptom onset, 30% of individuals with infection never develop symptoms and are 75% as infectious as those who do develop symptoms. This implies that persons with infection who never develop symptoms may account for approximately 24% of all transmission.  
> In this base case, **59% of all transmission came from asymptomatic transmission**, comprising 35% from presymptomatic individuals and 24% from individuals who never develop symptoms.  
> Under a broad range of values for each assumption, at least **50% of new infections** was estimated to have originated from exposure to individuals with infection but without symptoms.  
> In this decision analytical model, transmission from asymptomatic individuals was **estimated to account for more than half of all transmissions**. Measures such as wearing masks, hand hygiene, social distancing, and strategic testing of people who are not ill will be foundational to slowing the spread of COVID-19. |
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| **Clin Infect Dis. 07JAN2021** | Association Between Chronic Use of Immunosuppressive Drugs and Clinical Outcomes From Coronavirus Disease 2019 (COVID-19) Hospitalization: A Retrospective Cohort Study in a Large US Health System | Anderson K.M., et al. USA gotopaper | Public Health / Epidemiology | Does chronic use of immunosuppressive drugs worsens or improves the severity of COVID-19?  
> Retrospective cohort study, adults with acute inpatient hospital admission (March -August 2020)  
> confirmed or suspected COVID-19  
> chronic immunosuppression was defined as prescriptions for immunosuppressive drugs current at the time of admission.  
> Outcomes: mechanical ventilation, in-hospital mortality, and length of stay.  

**Findings:**  
> 2121 patients admitted with laboratory-confirmed (1967; 93%) or suspected (154; 7%) COVID-19  
> median age of 55 years (40–67).  
> of these, 108 (5%) were classified as immunosuppressed before COVID-19, primarily with prednisone (>7.5 mg/day), tacrolimus, or mycophenolate mofetil.  
> Among the entire cohort, 311 (15%) received mechanical ventilation  
> The median (interquartile range) length of stay was 5.2 (2.5–10.6) days  
> 1927 (91%) survived to discharge  
> no significant differences in the risk of mechanical ventilation, in-hospital mortality or length of stay among individuals with immunosuppression and counterparts.  

Chronic use of immunosuppressive drugs was neither associated with worse nor better clinical outcomes among adults hospitalized with COVID-19 in this setting. |
Evaluation of a semiobjective olfactory test developed to assess patient-reported chemosensory dysfunction prior to testing for the presence SARS-CoV-2  
> Diagnostic study conducted in a COVID-19 screening center in France (March-April, 2020)  
> Participants: health care workers or outpatients with symptoms or with close contact with an index case.  
> Participants interviewed to ascertain their symptoms and then Clinical Olfactory Dysfunction Assessment (CODA) (ad hoc test developed for a simple and fast evaluation of olfactory function).  
Assessment followed a standardized procedure in which participants identified and rated the intensity of 3 scents (lavender, lemongrass, and mint) to achieve a summed score ranging from 0 to 6. The COVID-19 status was assessed using RT PCR.  

**Findings:**  
> 809 participants, female to male sex ratio: 2.8. Mean age: 41.8 years (18-94).  
> Asymptomatic or mild disease patients; 58 (7.2%) tested positive for SARS-CoV-2.  
> Chemosensory dysfunction was reported by 20 of 58 participants (34.5%) with confirmed COVID-19 vs 29 of 751 participants (3.9%) who tested negative for COVID-19  
> Olfactory dysfunction, either self-reported or clinically ascertained (CODA score ≤3), yielded similar sensitivity and specificity for COVID-19 diagnosis.  
> Concordance was high between reported and clinically tested olfactory dysfunction, with a Gwet AC1 of 0.95 (95% CI, 0.93-0.97).  
> Of 19 participants, 15 (78.9%) with both reported olfactory dysfunction and a CODA score of 3 or lower were confirmed to have COVID-19.  
> The CODA score also revealed 5 of 19 participants (26.3%) with confirmed COVID-19 who had previously unperceived olfactory dysfunction.  

Olfactory dysfunction was suggestive of COVID-19, particularly when clinical testing confirmed anamnesis. However, normal olfaction was most common among patients with COVID-19. |
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| NEJM 07JAN2021   | Antibody Status and Incidence of SARS-CoV-2 Infection in Health Care Workers | Lumley S.F., et al. UK [gotopaper](#) | Immunology | > Study relationship between the presence of antibodies to severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) and the risk of subsequent reinfection  
> Incidence of SARS-CoV-2 infection confirmed by polymerase chain reaction (PCR) in seropositive and seronegative HCW attending testing of asymptomatic and symptomatic staff at Oxford University Hospitals  
> Baseline antibody status was determined by anti-spike (primary analysis) and anti-nucleocapsid IgG assays  
> Followed for up to 31 weeks  
> 12,541 health care workers participated having anti-spike IgG measured  
**Findings:**  
> A total of 223 anti-spike–seronegative health care workers had a positive PCR test (1.09 per 10,000 days at risk), 100 during screening while they were asymptomatic and 123 while symptomatic, whereas 2 anti-spike–seropositive health care workers had a positive PCR test (0.13 per 10,000 days at risk), and both workers were asymptomatic when tested (adjusted incidence rate ratio, 0.11; 95% confidence interval, 0.03 to 0.44; P = 0.002)  
> The presence of anti-spike or anti-nucleocapsid IgG antibodies was associated with a substantially reduced risk of SARS-CoV-2 reinfection in the ensuing 6 months. |
| NEJM 06JAN21     | Early High-Titer Plasma Therapy to Prevent Severe Covid-19 in Older Adults | Libster R., et al. Argentina/USA [gotopaper](#) | Therapeutics | Convallescent plasma administration at early COVID19 patients  
> Randomized, double-blind, placebo-controlled trial of convalescent plasma with high IgG titers against SARS-CoV-2 in older adult patients within 72 hours after the onset of mild Covid-19 symptoms.  
> 160 patients randomized  
**Primary end point:** severe respiratory disease (respiratory rate of 30 breaths per minute or more, an oxygen saturation of less than 93% while the patient was breathing ambient air, or both)  
**Trial stopped early at 76% of projected sample size because a decrease in Covid-19.**  
**Findings:**  
> Severe respiratory disease developed in 13 of 80 patients (16%) who received convalescent plasma and 25 of 80 patients (31%) who received placebo (relative risk, 0.52; 95% confidence interval [CI], 0.29 to 0.94; P=0.03), with a relative risk reduction of 48%.  
> No solicited adverse events were observed.  
**Early administration of high-titer convalescent plasma against SARS-CoV-2 to mildly ill infected older adults reduced the progression of Covid-19.** |
| Nature 06JAN2021 | A longitudinal study of SARS-CoV-2-infected patients reveals a high correlation between neutralizing antibodies and COVID-19 severity | Legros V., et al. France [gotopaper](#) | Immunology | Cohort study of 140 SARS-CoV-2 qPCR-confirmed infections, including patients with mild symptoms and more severe forms (intensive care included).  
The neutralizing antibody (nAb) responses were assessed using either live SARS-CoV-2 particles or retroviruses pseudotyped with the SARS-CoV-2 S viral surface protein (Spike).  
**Findings:**  
> ICU patients displayed high nAb activity compared to other groups with milder disease symptoms. nAb titers correlated strongly with disease severity and with anti-spike IgG levels.  
> The anti-S IgG response can be used as a marker of neutralizing activity in individuals.  
> Serum from individuals diagnosed with OC43, 229E, NL63, and HKU1 coronavirus infections but not infected with SARS-CoV-2 failed to cross-neutralize SARS-CoV-2 suggesting the absence of cross-neutralization between SARS-CoV-2 and endemic coronaviruses.  
> The D614G mutation did not affect the nAb activity of the serum samples from our cohort indicating that this highly prevalent mutation is not associated with SARS-CoV-2 resistance to neutralization. |
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<th>Key facts</th>
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| Science 06JAN2021 | Immunological memory to SARS-CoV-2 assessed for up to 8 months after infection | Dan J.M., et al. USA [gotopaper](#) | Immunology | > Understanding immune memory to SARS-CoV-2 and for assessing the likely future course of the COVID-19 pandemic.  
> 2254 samples from 188 COVID-19 cases, including 43 samples at ≥ 6 months post-infection  
**Findings**  
> IgG to the Spike protein was relatively stable over 6+ months  
> Spike-specific memory B cells were more abundant at 6 months than at 1 month post symptom onset  
> SARS-CoV-2-specific CD4+ T cells and CD8+ T cells declined with a half-life of 3-5 months  
Each component of SARS-CoV-2 immune memory exhibited distinct kinetics |
| Clin Infect Dis. 06JAN2021 | The duration, dynamics and determinants of SARS-CoV-2 antibody responses in individual healthcare workers | Lumley S.F., et al. USA [gotopaper](#) | Immunology | > SARS-CoV-2 IgG antibody measurements used to estimate the proportion of a population exposed or infected and may be informative about the risk of future infection  
> 6 months of data from a longitudinal seroprevalence study of 3276 UK healthcare workers with measurements of SARS-CoV-2 anti-nucleocapsid and anti-spike IgG  
> Interval censored survival analysis was used to investigate the duration of detectable responses  
> Bayesian mixed linear models were used to investigate anti-nucleocapsid waning  
**Findings**  
> SARS-CoV-2 anti-nucleocapsid antibodies wane within months (Anti-nucleocapsid IgG levels rose to a peak at 24 (95% credibility interval, CrI 19-31) days post first PCR-positive test, before beginning to fall), and faster in younger adults and those without symptoms.  
> Higher maximum observed anti-nucleocapsid titres were associated with longer estimated antibody half-lives  
> Anti-spike IgG remains stably detected.  
> Ongoing longitudinal studies are required to track the long-term duration of antibody levels and their association with immunity to SARS-CoV-2 reinfection |
| JAMA Netw. 05JAN2021 | Estimation of US SARS-CoV-2 Infections, Symptomatic Infections, Hospitalizations, and Deaths Using Seroprevalence Surveys | Angulo F.J., et al. USA [gotopaper](#) | Public Health / Epidemiology | Cross-sectional study of respondents of all ages, data from 4 regional and 1 nationwide Centers for Disease Control and Prevention (CDC) seroprevalence surveys between April and August 2020 were used to estimate infection and symptomatic underreporting multipliers.  
**Main Outcomes:** SARS-CoV-2 infections, symptomatic infections, hospitalizations, and deaths.  
**Findings:**  
> 14.3% of the US population was infected with SARS-CoV-2 and 8.6% had a symptomatic infection, with an infection hospitalization ratio of 2.0% and symptomatic fatality ratio of 1.1% through Nov 15, 2020.  
The US population remains a long way from herd immunity. The number of estimated COVID-19 deaths is also remarkably more than the reported deaths in the US through Nov 15, 2020, supporting the conclusion that approximately 35% of COVID-19 deaths are not reported.  
**Limitations:** Estimate the COVID-19 disease burden in the US using underreporting multipliers derived from the 10 specific states may not be nationally representative. |
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<td>BMJ Thorax 05JAN2021</td>
<td>Current smoking and COVID-19 risk: results from a population symptom app in over 2.4 million people</td>
<td>Hopkinson N.S., et al. UK</td>
<td>Public Health / Epidemiology</td>
<td>Main study outcome: development of ‘classic’ symptoms of COVID-19 during the pandemic defined as fever, new persistent cough and breathlessness and their association with current smoking. &gt; UK users of the Zoe COVID-19 Symptom Study app provided baseline data including demographics, anthropometrics, smoking status and medical conditions, and were asked to log their condition daily. &gt; Participants who reported that they did not feel physically normal were then asked by the app to complete a series of questions, including 14 potential COVID-19 symptoms and about hospital attendance. &gt; The number of concurrent COVID-19 symptoms was used as a proxy for severity and the pattern of association between symptoms was also compared between smokers and non-smokers. <strong>Findings:</strong> Data on 2,401,982 participants, mean (SD) age 43.6 (15.1) years, 63.3% female, overall smoking prevalence 11.0%. &gt; 834,437 (35%) participants reported being unwell and entered one or more symptoms. &gt; Current smokers were more likely to report symptoms suggesting a diagnosis of COVID-19: classic symptoms adjusted OR (95% CI) 1.14 (1.10 to 1.18); &gt;5 symptoms 1.29 (1.26 to 1.31); &gt;10 symptoms 1.50 (1.42 to 1.58). &gt; The pattern of association between reported symptoms did not vary between smokers and non-smokers. <strong>Data are consistent with people who smoke being at an increased risk of developing symptomatic COVID-19.</strong></td>
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<td>JAMA Netw. 04JAN2021</td>
<td>Risk Factors Associated With All-Cause 30-Day Mortality in Nursing Home Residents With COVID-19</td>
<td>Panagiotou O.A., et al. USA</td>
<td>Public Health / Epidemiology</td>
<td>Cohort study conducted at 351 US nursing homes among 5256 nursing home residents with COVID-19–related symptoms who had SARS-CoV-2 infection confirmed by PCR testing between March 16 and September 15, 2020. <strong>Main Outcome:</strong> Death due to any cause within 30 days of the 1st positive SARS-CoV-2 test result. <strong>Findings:</strong> &gt; Compared with residents aged 75 to 79 years, the odds of death were 1.46 times higher for residents aged 80 to 84 years, 1.59 times higher for residents aged 85 to 89 years, and 2.14 times higher for residents aged 90 years or older. &gt; Women had lower risk for 30-day mortality than men (odds ratio 0.69). &gt; Comorbidities associated with 30-day mortality: diabetes (OR, 1.21) and chronic kidney disease (OR, 1.33). &gt; Fever (OR, 1.66), shortness of breath (OR, 2.52), tachycardia (OR, 1.31), and hypoxia (OR, 2.05). &gt; Compared with intact cognitively residents: the odds of death among residents with moderate cognitive impairment (CI) were 2.09 times higher, and 2.79 times higher for residents with severe CI. &gt; Compared with residents with no or limited impairment in physical function (IPF), the odds of death among residents with moderate IPF were 1.49 times higher, and 1.64 times higher among residents with severe IPF. Once infected, <strong>those with baseline functional limitations, cognitive impairment, and disease severity are at heightened risk for mortality.</strong></td>
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<td>Science 04JAN2021</td>
<td>Neutralizing antibody titres in SARS-CoV-2 infections</td>
<td>Lau E.H.Y., et al. USA gotopaper</td>
<td>Immunology</td>
<td>Characterization of neutralizing antibody persistence in infected patients. Testing of 293 sera from an observational cohort of 195 reverse transcription polymerase chain reaction (RT-PCR) confirmed SARS-CoV-2 infections collected from 0 to 209 days after onset of symptoms. Findings: &gt; Of 115 sera collected ≥61 days after onset of illness tested using plaque reduction neutralization (PRNT) assays, 99.1% remained seropositive for both 90% (PRNT90) and 50% (PRNT50) neutralization endpoints. &gt; PRNT50 titres dropping to the detection limit of a titre of 1:10 for severe, mild and asymptomatic patients takes at least 372, 416 and 133 days &gt; At day 90 after onset of symptoms (or initial RT-PCR detection in asymptomatic infections), it took 69, 87 and 31 days for PRNT50 antibody titres to decrease by half (T1/2) in severe, mild and asymptomatic infections, respectively. &gt; Patients with severe disease had higher peak PRNT90 and PRNT50 antibody titres than patients with mild or asymptomatic infections. &gt; Age did not appear to compromise antibody responses, even after accounting for severity. SARS-CoV-2 infection elicits robust neutralizing antibody titres in most individuals.</td>
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<td>Nature Commun. 04JAN2021</td>
<td>Dose-dependent response to infection with SARS-CoV-2 in the ferret model and evidence of protective immunity</td>
<td>Ryan K.A., et al. UK gotopaper</td>
<td>Immunology / Preclinical model</td>
<td>&gt; Understand if ferrets are a suitable species for a model of human SARS-CoV-2 infection &gt; Dose titration study of SARS-CoV-2 in the ferret model &gt; Animals are challenged intranasally with a range of titres of SARS-CoV-2 (5 × 10^2, 5 × 10^4 and 5 × 10^6 pfu) in 1 ml volume Findings &gt; After a high (5 × 10^6 pfu) and medium (5 × 10^4 pfu) dose of virus is delivered, intranasally, viral RNA shedding in the upper respiratory tract (URT) is observed in 6/6 animals &gt; Only 1/6 ferrets show similar signs after low dose (5 × 10^2 pfu) challenge &gt; Ferrets re-challenged, after virus shedding ceased, are fully protected from acute lung pathology &gt; The endpoints of URT viral RNA replication &amp; distinct lung pathology are observed most consistently in the high dose group &gt; This ferret model of SARS-CoV-2 infection presents a mild clinical disease</td>
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<td>Clin Infect Dis. 03JAN2021</td>
<td>Distinct disease severity between children and older adults with COVID-19: Impacts of ACE2 expression, distribution, and lung progenitor cells</td>
<td>Zhang Z., et al. China gotopaper</td>
<td>Epidemiology</td>
<td>&gt; Examine the expression pattern of angiotensin-converting enzyme 2 (ACE2), the cell-entry receptor for severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), and the role of lung progenitor cells in children and older patients. &gt; cohort of 299 patients with COVID-19 &gt; Expression and distribution of ACE2 and lung progenitor cells examinations: combination of public single-cell RNA-seq datasets, lung biopsies, and ex vivo infection of lung tissues with SARS-CoV-2 pseudovirus in children and older adults Findings &gt; Compared to children, ACE2 positive cells are generally decreased in older adults and mainly presented in the lower pulmonary tract (alveolar region) and rarely in airway regions in the older adults (p &lt; 0.01). &gt; The lung progenitor cells are also decreased. These risk factors may impact disease severity and recovery from pneumonia caused by SARS-CoV-2 infection in older patients.</td>
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