Summary of Major Literature Related to COVID-19 (Week of May 4–10)

Led by Loren Lipworth and Qi Dai, with contribution from XO Shu, D Yu, M Shrubsole, S Sudenga, A Beeghly-Fadiel, Q Cai, P Pradhan, M Pike (Epidemiology) and A Ahonkhai, H Algood (Infectious Diseases) Department of Medicine

*This is informational and not intended to create variance from VUMC policies/guidance.



STATISTICS - Tennessee and Nashville

EPIDEMIOLOGY

- <u>Assessing Differential Impacts of COVID-19 on Black Communities</u>. Millett et al. amfAR preprint. May 2. See also: <u>COVID-19 Racial Disparities in U.S. Counties</u>.
- Reporting by state health departments of cumulative COVID-19 data by race/ethnicity is incomplete
- Using publicly available county-level data, COVID-19 cases and deaths were compared between disproportionately (>13%) black and all other (<13% black) counties in the US
 - o 91% (616/677) of disproportionately black counties are located in the southern US
 - disproportionately black counties had higher comorbidity rate, higher proportion uninsured or unemployed, more crowded living conditions, and higher levels of air pollution
- 49% and 55% of COVID-19 cases and deaths, respectively, occurred in the 22% of counties that are disproportionately black
- Counties with higher proportions of black residents had more COVID-19 diagnoses (RR 1.24, 95% CI 1.17-1.33) and deaths (RR 1.18, 95% CI 1.00-1.40), after adjusting for county-level characteristics such as age, poverty, comorbidities, and epidemic duration

- COVID-19 deaths were higher in disproportionally black rural and small metro counties
- <u>Limitations</u>: ecologic analysis based on county-level data; potential for residual confounding by comorbidities, as well as socioeconomic and behavioral factors including social distancing practices
- Obesity could shift severe COVID-19 disease to younger ages. Kass et al. Lancet. May 4.
 Obesity a Risk Factor for Severe COVID-19 Infection: Multiple Potential Mechanisms. Sattar et al. Circulation. April 22
- Early reports suggested obesity may be an underappreciated risk factor for COVID-19 in the US
- Study examined the correlation between BMI and age among 265 COVID-19 patients (58% males, age 15-90 years) admitted to ICUs in the US
 - $\circ~$ Median BMI was 29 kg/m², with 25% exceeding a BMI of 35 kg/m²
- Younger individuals admitted to ICU were more likely to be obese
- In an earlier study among individuals with COVID-19 aged less than 60 years in New York City, those with a BMI between 30-34 and >35 Kg/m were 1.8 times and 3.6 times more likely to be admitted to critical care, respectively, than individuals with a BMI <30
- Obesity can restrict ventilation and reduce forced expiratory volume, impair immune responses, increase basal inflammatory status, and induce oxidant stress
- Implication: Obesity shifts severe COVID-19 to younger ages
- <u>Covid-19 in Immune-Mediated Inflammatory Diseases Case Series from New York</u>. Haberman et al. NEJM. April 29.
- Case series of 86 confirmed or highly suspected symptomatic COVID-19 patients at NYU Langone Health with known immune-mediated inflammatory disease and receiving anticytokine biologics or other immune-modulatory therapies
 - 62 (72%) were receiving biologics or JAK inhibitors
 - Incidence of hospitalization was 16%
- Compared to non-hospitalized patients, hospitalized patients:
 - were older (50 vs 46 years)
 - had higher percentage of hypertension (36 vs 8%), diabetes (14 vs 4%), or COPD (7 vs 4%)
 - had 30-40% higher odds of use of oral glucocorticoids, hydroxychloroquine or methotrexate, but not biologics or JAK inhibitors, after adjusting for comorbidities
- Implications:
 - incidence of hospitalization among patients with immune-mediated inflammatory disease is consistent with general population in NYC
 - baseline use of biologics is not associated with worse COVID-19 outcomes in this study

Children

- 4. Hyperinflammatory shock in children during COVID-19 pandemic. Riphagen et al. Lancet. May 7.
- Case cluster of 8 children with hyperinflammatory shock, showing features similar to atypical Kawasaki disease, in pediatric intensive care unit in London, UK
 - Four (50%) children had known family exposure to COVID-19
 - All children tested SARS-CoV-2 negative while inpatient
 - No pathological organisms were identified in 7 of the children
 - $\circ~$ All but one were in the 75% for weight
- Clinical presentation was similar with unrelenting fever, variable rash, conjunctivitis, peripheral edema, and generalized extremity pain with significant gastrointestinal symptoms
- Since discharge, two children tested positive for SARS-CoV-2 (one postmortem). All children are receiving ongoing surveillance for coronary abnormalities

- Authors suggest this clinical picture represents a new phenomenon that children with asymptomatic SARS-CoV-2 infection may manifest as a hyperinflammatory syndrome with multiorgan involvement similar to Kawasaki syndrome
- Update states authors have now treated more than 20 children with similar clinical presentation, 10 of these are antibody positive for SARS-CoV-2.
- See also: <u>NYC Department of Health Advisory Statement.</u> Similar hyperinflammatory syndrome among 64 children and teens in NYC has been reported
- 5. Atypical presentation of COVID-19 in young infants. Nathan et al. Lancet. April 27.
- Trousseau Hospital in Paris observed an increase in number of young (< 3 months) infants presenting with SARS-CoV-2 infections one week after quarantine began
- Admitted 14 infants with 5 (all boys) testing positive for SARS-CoV-2. All 14 infants presented with fever and/or respiratory symptoms
- 4 of the COVID-19 positive boys showed neurological symptoms at admission. Their clinical course was rapidly favorable, which allowed for discharge after 1-3 days.
- All infants' parents showed mild signs of viral infection that could be related to undiagnosed COVID-19 with no mention of testing
- Implication: infants younger than 3 months with isolated fever should be tested for SARS-CoV-2

TRANSMISSION/PPE

- 6. <u>Universal Masking is Urgent in the COVID-19 Pandemic: SEIR and Agent Based Models, Empirical</u> <u>Validation, Policy Recommendations</u>. Kai et al. arXiv preprint. April 21.
- Two modeling approaches for predicting the impact of universal face mask wearing on the spread of SARS-CoV-2
- Significant impact when at least 80% of the population wears masks
- Minimal impact when only 50% or less of the population wears masks
- Significant impact when universal masking is adopted early (by Day 50 of a regional outbreak) compared to late
- Effects hold for homemade masks
- Empirical data showed strong correlation between early universal masking and successful suppression of daily case growth rates and/or reduction from peak daily growth rates, consistent with model predictions
- <u>Implication</u>: **Universal masking is a key non-pharmaceutical intervention** for containing or slowing the spread of the COVID-19 pandemic as governments exit society lockdowns
 - combined with social distancing and mass contact tracing, this is more sustainable from economic, social and mental health perspectives than full lockdown
 - o depends on strong public health messaging and the acceptance of masking in the population
 - proper masking techniques and norms need to be taught
- <u>Changes in contact patterns shape the dynamics of the COVID-19 outbreak in China</u>. Zhang et al. Science. April 29.
 - Mathematical models were used to address age differences in susceptibility and how altered social patterns affect transmission of COVID-19
 - Using a mean time between successive generations of cases of 5.1 days, strict social distancing measures were found to significantly reduce R₀ to below an epidemic threshold
 - Findings were robust across a variety of sensitivity analyses
 - School closures also impacted disease dynamics. However, models indicated that this approach alone would not be sufficient to prevent an outbreak

- <u>Limitations</u>:
 - contact testing and rapid quarantine of suspected and confirmed positive cases was also in place, which could influence model results
 - o differences in population structures and age-mixing behavior patterns limit generalizability

TREATMENT/EMERGING DRUG TARGETS

- Triple combination of interferon beta-1b, lopinavir-ritonavir, and ribavirin in the treatment of patients admitted to hospital with COVID-19: an open-label, randomised, phase 2 trial. Hung et al. Lancet. May 8.
- Multicenter, prospective, open-label, randomised, phase 2 trial in adults with COVID-19 admitted to six hospitals in Hong Kong (<u>NCT04276688</u>)
- Patients were randomly assigned (2:1) to a 14-day triple combination of lopinavir 400 mg and ritonavir 100 mg every 12 h, ribavirin 400 mg every 12 h, and three doses of 8 million IUs of interferon beta-1b on alternate days (combination group, N=86) or to 14 days of lopinavir 400 mg and ritonavir 100 mg every 12 h (control group, N=41)
- Primary endpoint was time to providing a nasopharyngeal swab negative for SARS-CoV-2 RT-PCR
- Median number of days from symptom onset to start of study treatment was 5 days (IQR 3–7)
- The combination group had a significantly shorter median time from start of study treatment to negative nasopharyngeal swab (7 days [IQR 5–11]) than the control group (12 days [8–15]; hazard ratio 4.37 [95% CI 1.86–10.24], p=0.001)
- Combination group also showed improvements in secondary outcomes, including shorter time to complete resolution of symptoms and shorter hospital stay
- Adverse events included self-limited nausea and diarrhea with no significant difference between the two treatment groups. One patient in the control group discontinued treatment because of biochemical hepatitis
- Limitations:
 - Open label, no placebo group
 - Subgroup (n=34) within combination group who started treatment 7 days or more after symptom onset omitted interferon beta-1b because of concerns about proinflammatory sideeffects
- <u>Implications</u>: Early treatment with the triple combination of antiviral therapy with interferon beta-1b, lopinavir—ritonavir, and ribavirin is safe and highly effective in shortening duration of virus shedding, alleviating symptoms, and facilitating the discharge of patients with mild to moderate COVID-19
- 9. Interleukin-1 blockade with high-dose anakinra in patients with COVID-19, acute respiratory distress syndrome, and hyperinflammation: a retrospective cohort study. Cavalli et al. Lancet Rheum. May 7.
- A small retrospective study conducted in Italy to evaluate clinical outcomes of patients treated with anakinra in addition to standard treatment (hydroxychloroquine + lopinavir + ritonavir)
- At 21 days, 29 patients who received anakinra (5 mg/kg twice a day intravenously) showed reductions in serum CRP; 21 (72%) of 29 patients showed improvements in respiratory function vs. 8 (50%) of 16 patients who received standard treatment only
- Survival was 90% in anakinra group vs. 56% in standard treatment group (p=0.009); ventilation-free survival was 72% in anakinra group vs. 50% in standard treatment group (p=0.15)
- Discontinuation of anakinra was not followed by inflammatory relapses
- A phase 2 trial of intravenous anakinra in COVID-19 is ongoing (<u>NCT04324021</u>).
- <u>Implication</u>: Treatment with anakinra in addition to standard treatment was safe and associated with better clinical outcomes
- <u>Limitations</u>: Small and non-randomized study

VIROLOGY

- 10. <u>The coronavirus proofreading exoribonuclease mediates extensive viral recombination</u>. Gribble et al. bioRxiv preprint. April 25.
- Extensive RNA recombination is performed by SARS-CoV-2, MERS-CoV, and murine hepatitis virus (MHV) in culture determined by combining both short- and long-read RNA sequencing
- These divergent β-CoVs generate similar patterns of recombination junctions and diverse populations of defective viral genomes (DVG) and subgenomic mRNAs (sgmRNA)
- Inactivating the proofreading nonstructural protein (nsp14) 3'-to-5' exoribonuclease (nsp14-ExoN) in MHV decreased recombination events and patterns of recombination suggesting nsp14-ExoN is required not just for high fidelity replication and also recombination
- <u>Limitation</u>: no proofreading-deficient nsp14-ExoN catalytic mutant has been rescued in MERS-CoV or SARS-CoV-2 limiting the genetic inactivation study to MHV at this time.
- Implications:
 - nsp14-ExoN represents a highly-conserved and vulnerable target for virus inhibition and attenuation
 - treatment of SARS-CoV-2 infection with small molecules that target the replicase proteins including nsp14-ExoN may result in an altered recombination landscape and perturb viral fitness