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Background

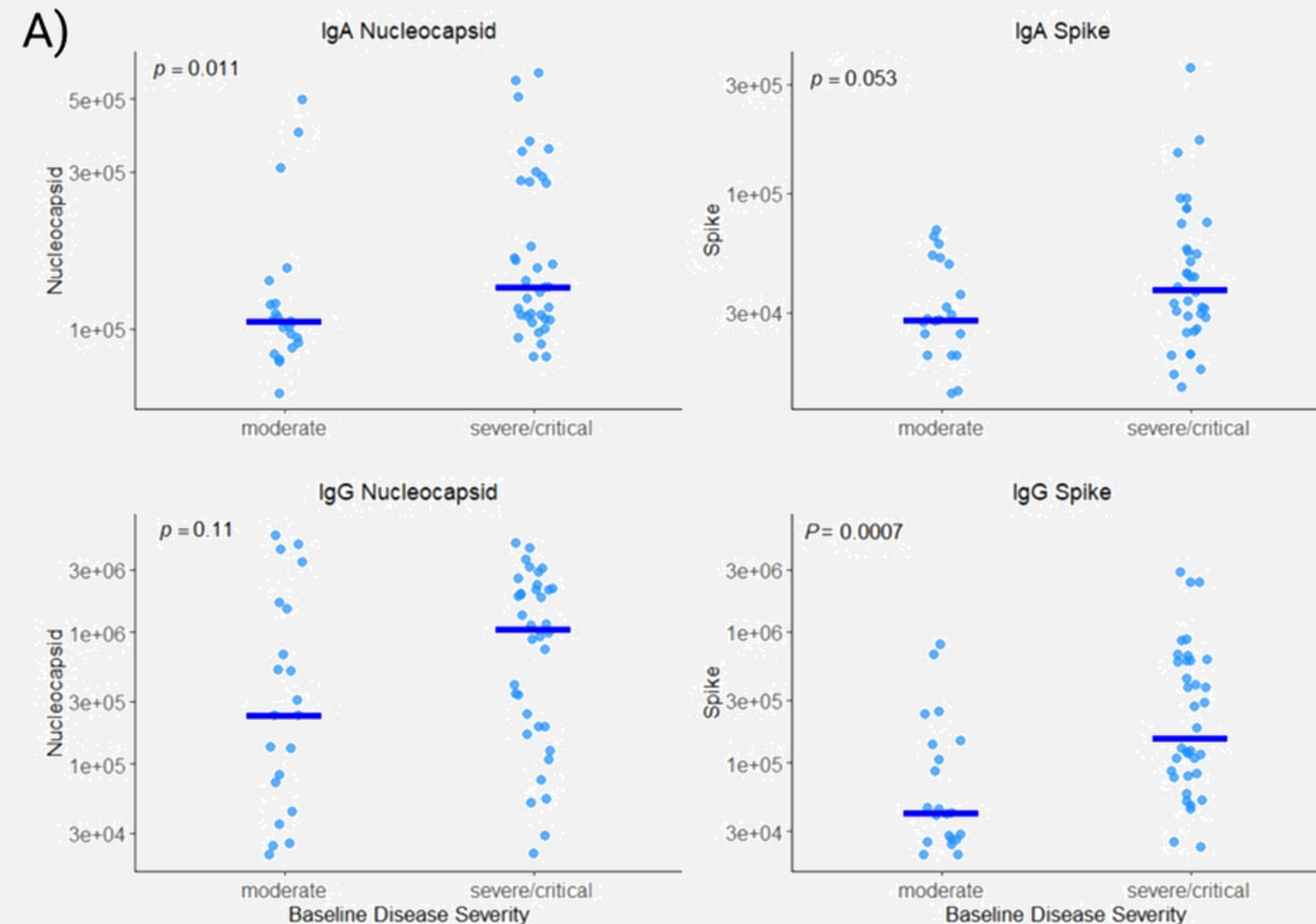
- In Coronavirus Disease 2019 (COVID-19) neutrophilia and a hyperactive neutrophil response has been associated with worse outcomes as activated neutrophils provoke endothelial injury, coagulation, and progression to ARDS^{1,2,3}.
- COVID-19 induces an altered and exuberant antibody response, with titers and subclasses of antibodies correlating with disease severity⁴.
- Neutrophils respond to IgG and IgA antibody complexes, leading to a variety of effector functions such as phagocytosis, degranulation, and NETosis^{5,6}
- Here we query associations between the endogenous SARS-CoV-2 antibody response and neutrophil activation in COVID-19.

Methods

- Baseline plasma samples from 57 patients hospitalized on oxygen with COVID-19 treated in a phase II study evaluating the safety and efficacy of fostamatinib, a spleen tyrosine kinase inhibitor, were obtained. We subsequently performed:
 - Quantitative measurements of SARS-CoV-2 specific antibodies using a luciferase-based immunoprecipitation system assay
 - Quantitative measurements of neutrophil specific biomarkers using Luminex technology
 - Neutrophil extracellular traps (NETs) as measured by myeloperoxidase-DNA (MPO-DNA) complexes by ELISA.
- Absolute neutrophil count (ANC) and immature granulocyte count (IGC) were measured from complete blood counts (CBC).

Acknowledgements

This work was supported by the Division of Intramural Research (DIR) of the National Heart, Lung and Blood Institute, National Institute of Allergy and Infectious Disease and the Clinical Center of the US National Institutes of Health



B)

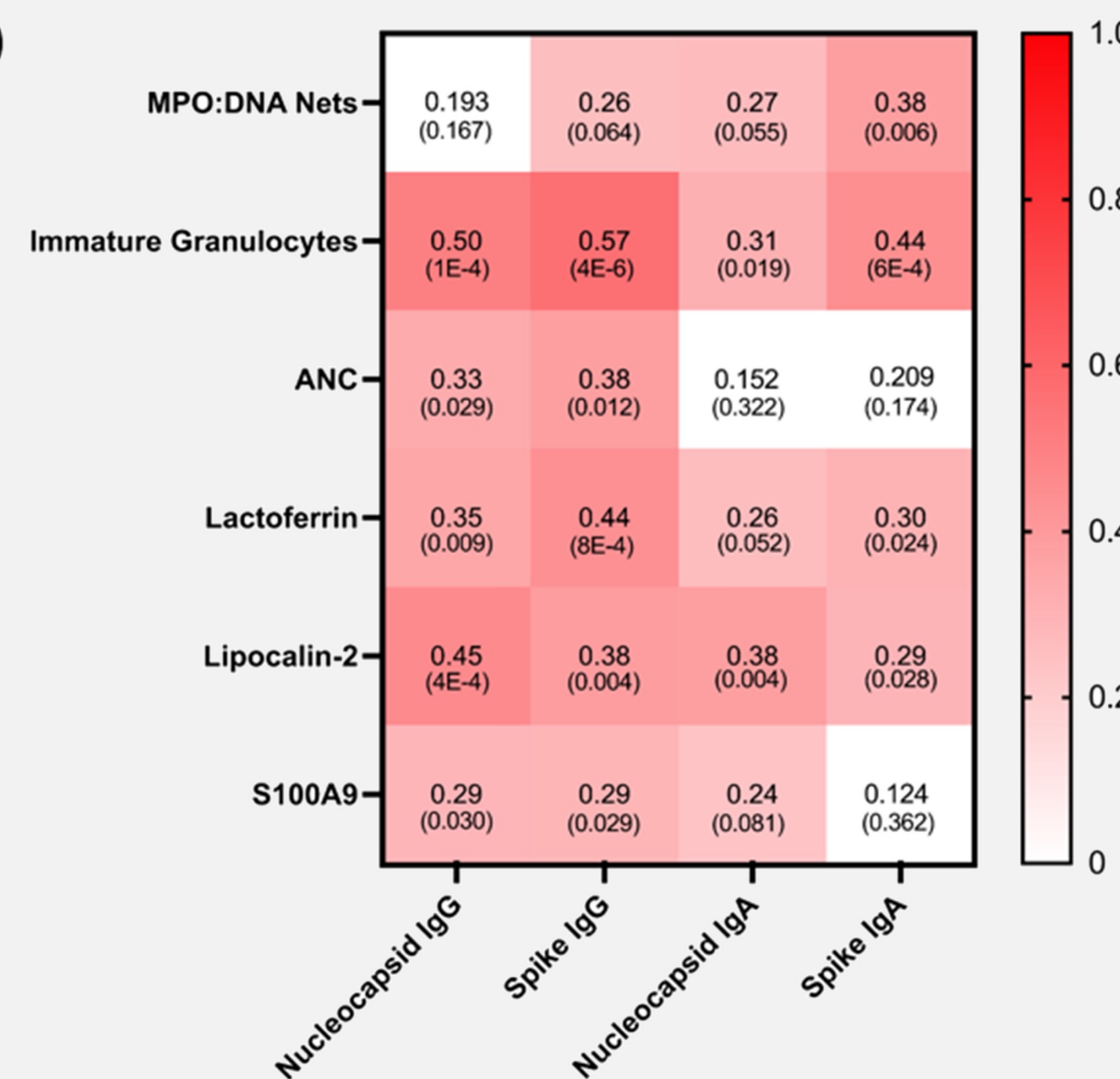


Figure 1A) Levels of anti-Spike and anti-Nucleocapsid IgA and IgG levels measured in the serum of 57 unvaccinated hospitalized COVID-19 patients. Moderate illness represents ordinal scale 5 requiring low flow oxygen, while severe/critical patients represent ordinal scale 6 and 7, requiring high flow oxygen, non-invasive or mechanical ventilation, respectively. P values are compared by a Wilcoxon ranked sum test.

Figure 1B) Heatmap showing Spearman correlations between levels of anti-Spike and anti-Nucleocapsid IgA and IgG and markers of neutrophil activation. P values for individual correlations are represented in parentheses. MPO (myeloperoxidase), ANC (absolute neutrophil count), S100A9 (S100 calcium binding protein A9).

Results

- Severe COVID-19 was associated with higher levels of nucleocapsid-IgA as well as spike-IgG compared to moderate disease
- Levels of IgG-spike and IgG-nucleocapsid both had significant correlations with Absolute Neutrophil Count
- All four antibody titers showed strong correlations with Immature Granulocyte count, evidence of emergency granulopoiesis and altered neutrophil populations
- S100A9, a component of calprotectin, lactoferrin and lipocalin-2, components of primary and secondary granules correlated strongly with antibody titers
- Circulating NETs levels demonstrated a significant correlation with spike IgA levels, and its correlations with IgG-spike and IgA-nucleocapsid additionally approached significance

Conclusion

- Higher anti-spike and anti-nucleocapsid IgG and IgA levels associated with more severe COVID-19 illness.
- Endogenous SARS-CoV-2 specific antibody levels associate with markers of emergency granulopoiesis and neutrophil hyperactivation
- Circulating NETs levels, which have been previously been associated with disease severity and thrombotic markers, correlated with antibody levels⁷
- Targeting antibody mediated neutrophil activation may prove beneficial to treatment of SARS-CoV-2⁸

References

- Bonaventura, A., et al., Endothelial dysfunction and immunothrombosis as key pathogenic mechanisms in COVID-19. *Nat Rev Immunol*, 2021. 21(5): p. 319-329
- Reusch, N., et al., Neutrophils in COVID-19. *Front Immunol*, 2021. 12: p. 652470.
- Meizlish ML, et al. A neutrophil activation signature predicts critical illness and mortality in COVID-19. *Blood Adv*. 2021 Mar 9;5(5):1164-1177.
- Garcia-Beltran, WF., COVID-19 neutralizing antibodies predict disease severity and survival. *Cell*. 2021 184(2):476-488
- Stacey HD, et al., IgA potentiates NETosis in response to viral infection. *Proc Natl Acad Sci U S A*. 2021 Jul 6;118(27):e2101497118.
- Wang Y, et al., Expression, Role, and Regulation of Neutrophil Fcγ Receptors. *Front Immunol*. 2019 Aug 27;10:1958.
- Zuo Y, et al., Neutrophil extracellular traps in COVID-19. *JCI Insight*. 2020 Jun 4;5(11):e138999.
- Strich, JR et al., Fostamatinib for the treatment of hospitalized adults with COVID: A randomized clinical trial, *Clin Infect Dis*, 2021;ciab732