## LETTER TO THE EDITOR

### **Open Access**



# Longitudinal assessment of systemic steroid therapy on hyperinflammatory endothelial biomarker profiles and serology responses of COVID-19 patients

Jonathan T. Sims<sup>1</sup>, Ching-Yun Chang<sup>1</sup>, Josh Poorbaugh<sup>1</sup>, Montanea Daniels<sup>1</sup>, Stephanie L. Beasley<sup>1</sup>, Lin Zhang<sup>1</sup>, George H. Rodgers<sup>1</sup>, Fabio Lena<sup>2</sup>, Leonardo G. Lacerenza<sup>2</sup>, Bruno Sposato<sup>2</sup>, Annabelle Dupont<sup>3</sup>, Sophie Susen<sup>3</sup>, Giacomo Casalini<sup>4</sup>, Mario Corbellino<sup>5</sup>, Justin Stebbing<sup>6</sup> and Venkatesh Krishnan<sup>1\*</sup>

Keywords: COVID-19, CXCL9, Cytokine, Dexamethasone, Inflammation, Serology

This first of its kind study provides objective context to the potential mechanism of action of corticosteroid use by connecting inflammatory biomarkers to IgG levels for the SARS-CoV-2 spike protein antigens and neutralization of ACE2 binding within patients across 3 institutions from Italy and France who received corticosteroids (dexame has one, n=5, or prednisolone, n=1) or usual standard of care (SOC, n=22) therapy. The median Ordinal Scale (WHO) upon admission was OS-5 and no difference in days from symptom onset or Ordinal scale at study entry was observed between these 2 groups (Additional file 1: Table S1, Fig. S1). Utilizing Olink multiplex technology and IL-19 ELISA, we assessed 185 analytes in the circulation of COVID-19 patients along with Luminex-based measurement of 10 immunoglobulins, including neutralization assessment (Additional file 2).

We observed prominent dysregulation of IL-8, CCL7/ MCP-3, S100A12/ENRAGE, and IL-6 in the circulation of these patient cohorts relative to 12 age/sex-matched healthy controls (HC) (Fig. 1A). IL-6 correlated modestly with baseline Ordinal scale and the broader inflammatory biomarker profile was characterized by strong

\*Correspondence: krishnan\_gary@lilly.com

correlations to circulating neutrophils and serum creatinine (Fig. 1A). Levels of TNFRSF10A, IL-10RA, CXCL9, TRAILR2, IL-18, and TNFa negatively correlated with SpO2 percentage and positively correlated with Ordinal Scale at admission emphasizing the potentially important role of these molecular pathways within hospitalized patients (Fig. 1A) [1].

Focusing on markers differentially regulated in COVID-19 patients, we observed a unique biomarker profile in patients depending on their therapeutic journey. Analytes elevated in baseline COVID-19, FGF21, CCL3, CCL4, CEACAM8, CCL13/MCP4, IL-19, IL15RA, ANGPT1, MMP7, and CXCL9, all decreased more in corticosteroid-treated patients relative to SOC (Fig. 1A).

We observed serology titers for all SARS-CoV-2 antigens with RBD mutant proteins increasing or remaining relatively unchanged within steroid-treated patients relative to the SOC-treated patients over the course of 7 days in this small study, mirrored by an increase in neutralization of the RBD-ACE2 interaction in the same steroid-treated patients (Fig. 1B). Further research into the mechanism of action of corticosteroid-reduced IgG levels will need to occur, but in this study, there was no detrimental impact on antibody responses to COVID-19 infection. Within these patients, repeated measured correlations demonstrated inverse correlation of



© The Author(s) 2022. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by/4.0/. The Creative Commons Public Domain Dedication waiver (http://creativecommons.org/publicdomain/zero/1.0/) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

<sup>&</sup>lt;sup>1</sup> Eli Lilly and Company, Lilly Corporate Center, Indianapolis, IN 46285, USA Full list of author information is available at the end of the article



ACE2 neutralization IC50 not only with ACE2, but also TNFSF11/TRANCE protein in the serum (Fig. 1B). This data argues that circulating antibodies with more potent neutralization (lower IC50) are found in patients with higher serum levels of ACE2 and TNFSF11/TRANCE, the former known to be shed by interactions with viral spike protein and the latter to decrease when viral pathogenesis is most robust [2, 3]. Related to the broad suppressive nature of corticosteroids, certain analytes reduced in circulation of COVID-19 patients, such as SCF and GDF2, were inversely correlated with serological IgG response markers (Fig. 1B). Intriguingly, a unique inflammatory biomarker profile was observed to correlate with the seasonal coronavirus protein control (NL63) relative to the profile linked to SARS-CoV-2 antigens and related RBD mutant proteins (Fig. 1B). Positive correlations were observed for IL-27, MERTK, IL-17A, CXCL9, CCL3, GAL9, TNF, CCL4, FGF21, VSIG2, DCN, CX3CL1, IL-19, CASP8, IL-10, AGRP, FABP2, CXCL10, CD84, SCF, and IFNg with ACE2 neutralization IC50 values (Fig. 1B). Higher ACE2 neutralization IC50 values indicate less potency of the antibody response and hence the observed increase in many proinflammatory signals (e.g. IL-19, TNF, CXCL9, and IFNg). This tight regulatory phenomenon between inflammatory biomarkers and serology was clearly evident within the tested timeframe of 7 days post corticosteroid treatment.

The correlation of COVID-19 inflammatory markers, which we previously linked to severe disease (e.g., IFNg, TNF, CXCL9, and IL-10), with higher IC50 neutralization values may demonstrate poor levels of ACE2/RBD interference in patients experiencing the hyperinflammatory endothelial-linked cytokine storm of COVID-19 [4].

Importantly, the elevation of markers indicative of innate immune activation (CCL13/MCP4, CCL3, CCL4, CXCL9) are reduced in steroid-treated patients. We observed that CXCL9 is an important COVID-19 biomarker describing immunological responses in corticosteroid-treated patients based on its correlation to Ordinal scale at baseline, decreased levels over time in corticosteroid-treated patients, and correlation with ACE2 neutralization IC50 values; therefore, warranting further evaluation and clinical monitoring within patients treated with corticosteroids. However, key systemic inflammatory cytokines linked to adaptive immune responses elevated in COVID-19 vs. HC are not differentially regulated in steroid vs SOC-treated patients, demonstrating that early effective targeted therapies against these will be most successful in rapidly reducing the inflammatory burden that severe patients experience.

#### **Supplementary Information**

The online version contains supplementary material available at https://doi.org/10.1186/s12967-022-03583-5.

Additional file 1: Table S1. Baseline characteristics of COVID-19 patients. Table S2. Recombinant proteins used in serology measurements. Figure S1. A Patient classification based on therapeutic decision prior to collection of first sample as either steroids or SOC (non-steroid). B, C Graphical box plot representations of the time (days) between a patient's symptom onset and collection of first sample (B) or Ordinal Scale classification at admission (C) between steroid and SOC patient groups.

Additional file 2. Additional Materials and Methods.

#### Acknowledgements

The authors would like to thank the continued dedication of healthcare workers across the world for their response to COVID-19. We are indebted to Julien Poissy, Carlotta Pucci, Michele Bindi, Cesira Nencioni, Leonardo Croci, and Maja Rossi for collaboration and assistance with this study. We thank Jeffrey Boyles and Denisa Foster for generating recombinant proteins for the serology assay. We thank Autum Auxier and Tomasz Izbicki for sample management and Robert Benschop for scientific input and support.

#### Author contributions

JTS, VK, JS, MC, FL, and SS conceived the study. JTS, JP, MD, SLB, and GHR conducted the laboratory experiments. JTS, C-YC, VK, JP and LZ designed and conducted the main analyses and interpreted the results. JP, GHR, FL, LGL, BS, AD, SS, GC, MC and JS also contributed to the discussion of analyses. JTS and VK wrote the first draft of the manuscript. JTS, JP, JS, and VK wrote the manuscript, with contributions and review by all other authors. All authors read and approved the final manuscript.

#### Funding

The analytical measures using Olink within the study were funded by Eli Lilly and Company. The physicians and the COVID-19 patients did not receive any funding or reimbursement from Eli Lilly and Company.

#### Availability of data and materials

The data from the current study are available from the corresponding author on reasonable request.

#### Declarations

#### Ethics approval and consent to participate

All consent forms and protocols were approved by appropriate ethical review boards prior to study initiation.

#### **Consent for publication**

Not applicable.

#### Competing interests

Jonathan T. Sims, Venkatesh Krishnan, Ching-Yun Chang, Josh Poorbaugh, Montanea Daniels, Stephanie L. Beasley, Lin Zhang, & George H. Rodgers are employees and may be shareholders of Eli Lilly and Company. Fabio Lena, Leonardo G. Lacerenza, Bruno Sposato, Annabelle Dupont, Sophie Susen, Giacomo Casalini, and Mario Corbellino report no conflicts of interest. Justin Stebbing is editor-in-chief of Oncogene, sat on several scientific advisory boards, including Benevolent AI, and consults with Lansdowne partners, Vitruvian, and Eli Lilly and Company, and sits on the Board of Directors for BB Biotech Healthcare Trust.

#### Author details

<sup>1</sup>Eli Lilly and Company, Lilly Corporate Center, Indianapolis, IN 46285, USA. <sup>2</sup>Department of Pharmaceutical Medicine, Misericordia Hospital, Grosseto, Italy. <sup>3</sup>Inserm, CHU Lille, Institut Pasteur de Lille, U1011-EGID, University of Lille, 59000 Lille, France. <sup>4</sup>Luigi Sacco Department of Clinical and Biomedical Sciences, University of Milan, Milan, Italy. <sup>5</sup>Division of Infectious Diseases, ASST Fatebenefratelli Sacco, Milan, Italy. <sup>6</sup>Department of Surgery and Cancer, Imperial College, London, UK.

# Received: 3 January 2022 Accepted: 23 February 2022 Published online: 08 September 2022

#### References

- Ye Q, Wang B, Mao J. The pathogenesis and treatment of the 'Cytokine Storm' in COVID-19. J Infect. 2020;80(6):607–13.
- Jia H, Neptune E, Cui H. Targeting ACE2 for COVID-19 therapy: opportunities and challenges. Am J Respir Cell Mol Biol. 2021;64(4):416–25.
- Haljasmägi L, et al. Longitudinal proteomic profiling reveals increased early inflammation and sustained apoptosis proteins in severe COVID-19. Sci Rep. 2020;10(1):20533.
- Sims JT, et al. Characterization of the cytokine storm reflects hyperinflammatory endothelial dysfunction in COVID-19. J Allergy Clin Immunol. 2021;147(1):107–11.

#### **Publisher's Note**

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

#### Ready to submit your research? Choose BMC and benefit from:

- fast, convenient online submission
- thorough peer review by experienced researchers in your field
- rapid publication on acceptance
- support for research data, including large and complex data types
- gold Open Access which fosters wider collaboration and increased citations
- maximum visibility for your research: over 100M website views per year

#### At BMC, research is always in progress.

Learn more biomedcentral.com/submissions

