Contents lists available at ScienceDirect

EClinicalMedicine



journal homepage: https://www.journals.elsevier.com/eclinicalmedicine

Complement inhibition in severe COVID-19 – Blocking C5a seems to be key

Endry H.T. Lim^a, Alexander P.J. Vlaar^{a,*}, Sanne de-Bruin^a, Matthijs C. Brouwer^b, Diederik van-de-Beek^b

^a Department of Intensive Care Medicine, Amsterdam University Medical Center, University of Amsterdam, Amsterdam UMC, 1100DD Amsterdam, the Netherlands ^b Department of Neurology, Amsterdam Neuroscience, Amsterdam University Medical Center, University of Amsterdam, Amsterdam, the Netherlands

ARTICLE INFO

Article History: Received 12 December 2020 Accepted 6 January 2021 Available online xxx

With great interest we read the report by Annane and colleagues describing the effect of blocking complement factor C5 with the antibody eculizumab in patients with severe COVID-19 [1]. Results of this non-controlled study show an important proof of principle of complement inhibition therapy in patients with severe COVID-19. Increasing evidence point towards a critical role of the proinflammatory anaphylatoxin C5a in the pathogenesis of severe COVID-19 [2–4]. A previous study showed that controlling the anaphylatoxin C5a in disease requires a specifically targeted inhibition [5].

The authors mention that frequency and dosage of eculizumab had to be increased during the study to achieve complete and sustained complement inhibition [1]. Concentrations of C5a during the study of patients treated with or without high or higher dose eculizumab may provide important information about the potential of complement inhibition in COVID-19. Alternatively, selective approaches blocking C5a could be preferred. We recently published results of a phase 2 trial, showing that selective C5a inhibition with vilobelimab is safe in patients with severe COVID-19, with secondary outcome results in favour of vilobelimab [4]. Because blockade of an upstream component in the complement pathways will inevitably affect the formation of the membrane attack complex, such upstream intervention might put patients with COVID-19 at risk of secondary bacterial infections. The authors could provide more insight in this issue by presenting C5a concentrations of patients treated with eculizumab, with a breakdown for initial and high dose of eculizumab.

Declaration of Competing Interest

Dr. Vlaar reports personal fees from InflaRx, outside the submitted work. All other authors declare no competing interests.

References

- Annane D, Heming N, Grimaldi-Bensouda L, Frémeaux-Bacchi V, Vigan M, Roux AL, et al. Eculizumab as an emergency treatment for adult patients with severe COVID-19 in the intensive care unit: a proof-of-concept study. EClinicalMedicine 2020. doi: 10.1016/j.eclinm.2020.100590.
- [2] Skendros P, Mitsios A, Chrysanthopoulou A, Mastellos DC, Metallidis S, Rafailidis P, et al. Complement and tissue factor–enriched neutrophil extracellular traps are key drivers in COVID-19 immunothrombosis. J Clin Investig 2020. doi: 10.1172/ JCI141374.
- [3] Carvelli J, Demaria O, Vély F, Batista L, Benmansour NC, Fares J, et al. Association of COVID-19 inflammation with activation of the C5a–C5aR1 axis. Nature 2020. doi: 10.1038/s41586-020-2600-6.
- [4] Vlaar APJ, de Bruin S, Busch M, Timmermans SAMEG, van Zeggeren IE, Koning R, et al. Anti-C5a antibody IFX-1 (vilobelimab) treatment versus best supportive care for patients with severe COVID-19 (PANAMO): an exploratory, open-label, phase 2 randomised controlled trial. Lancet Rheumatol 2020. doi: 10.1016/S2665-9913(20) 30341-6.
- [5] Riedemann NC, Habel M, Ziereisen J, Hermann M, Schneider C, Wehling C, et al. Controlling the anaphylatoxin C5a in diseases requires a specifically targeted inhibition. Clin Immunol 2017. doi: 10.1016/j.clim.2017.03.012.

https://doi.org/10.1016/j.eclinm.2021.100722

2589-5370/© 2021 The Author(s). Published by Elsevier Ltd. This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0/)



Letter

DOI of original article: http://dx.doi.org/10.1016/j.eclinm.2020.100590.

^{*} Corresponding author.

E-mail address: a.p.vlaar@amsterdamumc.nl (A.P.J. Vlaar).