

Table I. Shared identical epitope between Ankyrin 1 and SARS-CoV-2 surface glycoprotein¹

Protein	Accession number	Epitope amino acids	Identity percentage, %
SARS-CoV-2 surface glycoprotein	NCBI ID: YP_009724390.1	752-LLLQY-756	100
Ankyrin 1	UniProt ID: P16157	323-LLLQY-327	

ID, identifier; NCBI, National Center for Biotechnology Information.

¹We used for comparative analyses BlastP (available at: <https://blast.ncbi.nlm.nih.gov/Blast.cgi?PAGE=Proteins>) and the whole virus proteome (available at: <https://www.ncbi.nlm.nih.gov/nuccore/MN908947>).


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Keywords: ankirin 1, autoantibodies, autoimmunity, COVID-19, molecular mimicry, severe acute respiratory syndrome coronavirus 2

First published online 8 June 2020

doi: 10.1111/bjh.16883

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COVID-19 and ABO blood group: another viewpoint

Li *et al.*¹ have recently published 'Association between ABO blood groups and risk of SARS-CoV-2 pneumonia', an observation already reported a few weeks ago as a MedRxiv preprint by Zhao *et al.*² and which had a certain impact in the press.

In both studies, the ABO blood groups distribution of patients with coronavirus disease 2019 (COVID-19) were compared to that of controls from the local populations that showed that blood group A was associated with an increased risk of infection, whereas group O was associated with a decreased risk. Considering this information rather as a working hypothesis, some scientists have called for caution.³

However, as already strongly suggested by others,⁴ this variable susceptibility to severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) infection could be linked to circulating anti-A antibodies, which could interfere or even inhibit the virus–cell adhesion process.

We had the idea to analyse these important available data series from the anti-A or -B antibodies viewpoint instead of ABO blood group antigens as the authors did.

In fact, considering the largest series of patients with COVID-19 ($N = 1888$) analysed by Zhao *et al.*,² we compared the proportion of those possessing anti-A in their serum (i.e. those of B and O blood groups) and those who

Table I. Comparison of subjects with/without anti-A antibodies in their serum.

	RBC blood group	Control, n (%)	COVID-19, n (%)	χ^2	P	OR (95% CI)
With anti-A	B and O	2170 (58.7)	927 (52.2)			
	A and AB	1524 (41.3)	848 (47.8)	20.74	<0.001	1.30 (1.16–1.46)
Without anti-A	A	1188 (32.2)	670 (37.7)	19.97	<0.001	1.32 (1.17–1.49)
	AB	336 (9.1)	178 (10.0)	4.58	0.0323	1.24 (1.02–1.51)

Table II. Comparison of anti-A from O and from B subjects.

	RBC blood group	Control, n (%)	COVID-19, n (%)	χ^2	P	OR (95% CI)
Anti-A from O	O	1250 (57.6)	458 (49.4)			
Anti-A from B	B	920 (42.4)	469 (50.6)	17.64	<0.001	1.39 (1.19–1.62)

did not (i.e. those of A and AB blood groups) to the control cohort ($N = 3694$; Table I).

The results (Table I) indicate that subjects with anti-A in serum (i.e. B and O blood groups) are significantly less represented in the COVID-19 group than those lacking anti-A whatever the group: A and AB ($P < 0.001$), A ($P < 0.001$) or AB ($P = 0.0323$), whereas there was no significant difference versus circulating anti-B (data not shown).

We then wondered if there was a difference between anti-A from O and anti-A from B, and then we compared the supposed protective effect of anti-A from O and from B (Table II).


Whereas both blood group O and B patients possess circulating seric anti-A, it appears and it is statistically highly significant ($P < 0.001$) that O group patients are underrepresented (49.4 % vs. 57.6%), whereas B group patients are, on the contrary, overrepresented (50.6% vs. 42.4%), meaning that anti-A from O is more protective than anti-A from B.


This latter observation is probably related to the fact that the immunoglobulin predominant isotype of anti-B/anti-A is IgM in serum from group A and B individuals, but IgG in O group serum, an already known notion,⁵ which has been well documented by flow cytometry.⁶

In conclusion, this way of analysing the data strongly suggests that the presence of anti-A antibodies in serum and more specifically IgG anti-A, should be considered as a factor more significant than the blood group itself, as far as the relationship between COVID-19 susceptibility and ABO blood groups is concerned.

Far from intending to corroborate the authors' conclusions as such, we wanted to show that the resources of immuno-haematology allow several approaches that could perhaps be useful for the disease follow-up.

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Keywords: antibodies, blood group serology, COVID-19

First published online 8 June 2020

doi: 10.1111/bjh.16884

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