

CORRECTION

Correction: Breadth and function of antibody response to acute SARS-CoV-2 infection in humans

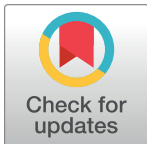
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There is an error in the caption for [S2 Fig](#). The cells in this experiment were transduced to express the unmutated full length wild-type spike protein and did not contain the stabilizing mutations as stated in the figure legend. Please see the full [S2 Fig](#) legend below.

There is also an error in the last sentence of the second paragraph of the Results subsection “Genetic and phenotypic characteristics of anti-spike glycoprotein antibodies”. The correct sentence is: All anti S2 MAbs bound to the MDCK-SIAT1 cells transduced to express the full-length spike antigen.

Supporting information

S2 Fig. The binding activity of anti-SARS-CoV-2 S2 antibodies with spike-expressed MDCK cells in flow cytometry. We produced MDCK-Spike by stably transducing parental MDCK-SIAT1 cells with cDNA expressing full-length unmutated SARS-CoV-2 spike glycoprotein. MDCK-H3 cells were stained in the control experiment. Anti-influenza H3 MAb BS-1A was included as an antibody control. Each experiment was repeated twice (n = 2). The binding percentage was presented as mean ± standard error of the mean. (TIF)



Reference

1. Huang K-YA, Tan TK, Chen T-H, Huang C-G, Harvey R, Hussain S, et al. (2021) Breadth and function of antibody response to acute SARS-CoV-2 infection in humans. *PLoS Pathog* 17(2): e1009352. <https://doi.org/10.1371/journal.ppat.1009352> PMID: 33635919

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