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Correspondence

SARS-CoV-2 spike E484K mutation reduces antibody neutralisation

It is concerning that emerging variants of SARS-CoV-2 can evade neutralising antibodies induced by previous infection or vaccination through mutations in the spike protein, including the receptor-binding domain (RBD). The asparagine (N) to tyrosine (Y) substitution at position 501 (N501Y), present in variants of concern belonging to the B.1.1.7, B.1.351, and P.1 lineages, does not seem to affect in-vitro neutralisation of human convalescent or post-vaccination sera. However, additional substitutions, such as E484K present in B.1.351 and P.1 lineages, might allow evasion from neutralising antibodies.¹⁻⁴ We did in-vitro microneutralisation assays with the USA-WA1/2020 virus and a recombinant (r)SARS-CoV-2 virus, which is identical to USA-WA1/2020 except for the E484K mutation introduced in the spike RBD. A total of 34 sera were selected from study participants on the basis of their SARS-CoV-2 S ELISA antibody titre (negative [N=4] vs weak [N=8], moderate [N=11], or strong positive [N=11]; appendix p 1). Additionally, we included sera from five individuals who received two doses of the Pfizer-BioNTech SARS-CoV-2 vaccine BNT162b2 (V1-V5). The same sera have been tested for neutralisation studies with a N501Y SARS-CoV-2 variant in our recent report.5 Serum neutralisation efficiency was lower with E484K rSARS-CoV-2 versus USA-WA1/2020 for vaccinated (3.4-fold) and convalescent samples (2.4 fold for low IgG, 4.2 fold for moderate IgG, and 2.6 fold for high IgG, based on geometric means; figure), and data were significantly different for the convalescent sera. These data suggest that the single E484K mutation in the RBD affects the binding of serum polyclonal neutralising antibodies. The decrease in neutralisation efficiency for sera with low or moderate IgG against SARS-CoV-2 spike protein could result in loss of neutralisation of the E484K recombinant virus. These data show the influence of a single mutation (E484K) present in SARS-CoV-2 B.1.351 and P.1 lineages on the neutralising activity of convalescent (infected with previous SARS-CoV-2 strains) and post-vaccination polyclonal sera. A limitation of our study is the small number of postvaccination samples tested. However, sera with high neutralisation titres against the USA-WA1/2020 strain (convalescent and post-vaccination) were still able to neutralise E484K rSARS-CoV-2. Therefore. vaccinations should induce the highest neutralisation titres possible, to maximise protection against antigenically drifted SARS-CoV-2 strains. Most SARS-CoV-2 vaccines are recommended as a prime-boost regimen. Due to vaccine shortage, some public health authorities have recommended to postpone the booster vaccination to be able to provide more individuals with a primer vaccination. This approach will result in lower neutralising antibody titres. Our data suggest that lower neutralising antibody titres might be problematic in the context of newly emerging SARS-CoV-2 variants, considering that this approach could leave some vaccinees unprotected. Which neutralisation titre correlates with (full) protection, and to what extent immune mechanisms beyond direct virus neutralisation, such as antibody-dependent cellular cytotoxicity by non-neutralising antibodies or T-cell mediated immunity, contribute to protection is unclear. The worldwide vaccination effort should aim at fully vaccinating as many people as possible by use of vaccination strategies that result in high neutralising antibody titres.



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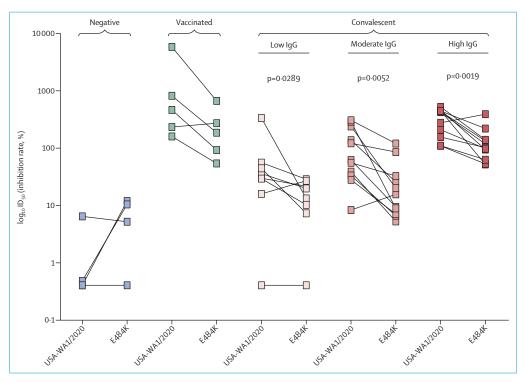


Figure: Human convalescent and post-vaccination sera neutralise E484K recombinant SARS-CoV-2 less efficiently than USA-WA1/2020 in an in-vitro microneutralisation assay

Convalescent sera are subdivided in low, moderate, and high IgG classes on the basis of anti-spike ELISA titres. Two-sided Mann Whitney-U tests were used to calculate statistical differences.

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Sonia Jangra, Chengjin Ye, Raveen Rathnasinghe, Daniel Stadlbauer, Personalized Virology Initiative study group, Florian Krammer, Viviana Simon, Luis Martinez-Sobrido, *Adolfo García-Sastre, *Michael Schotsaert adolfo.garcia-sastre@mssm.edu or michael.schotsaert@mssm.edu

Department of Microbiology (SJ, RR, DS, FK, VS, AG-S, MS), Department of Medicine, Division of Infectious Diseases (VS, AG-S), Global Health and

Emerging Pathogens Institute (SJ, RR, VS, AG-S, MS), Graduate School of Biomedical Sciences (RR), and The Tisch Cancer Institute (AG-S), Icahn School of Medicine at Mount Sinai New York, NY 10029, USA; Texas Biomedical Research Institute, San Antonio, TX, USA (CY, LM-S)

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