

Higher infectivity of the SARS-CoV-2 new variants is N501Y mutants: An insight from structural data

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Bioinformatic and MD Analysis of N501Y SARS-CoV-2 (UK) Variant. IFIP Advances in Information and Communication Technology, 2021, , 1-13.	0.5	1
2	The SARS-CoV-2 B.1.618 variant slightly alters the spike RBD's ACE2 binding affinity and is an antibody escaping variant: a computational structural perspective. RSC Advances, 2021, 11, 30132-30147.	1.7	57
4	Higher infectivity of the SARS-CoV-2 new variants is associated with K417N/T, E484K, and N501Y mutants: An insight from structural data. Journal of Cellular Physiology, 2021, 236, 7045-7057.	2.0	283
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