

Mina Kelleni¹

¹Affiliation not available

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Abstract

In the first version, I have made a mistake SARS COV-2 Vaccines have not shared in development of the original SARS CoV-2 B.1.617 variants in India (yet they might have shared in the evolution of the more virulent delta plus variant). I have updated the manuscript, thanks to Authorea, and in the updated versions Remdesivir, Favipiravir and Dexamethasone are suggested as potential crucial causes that led to B.1.617 variants in India and elsewhere. Moreover, SARS CoV-2 mass vaccination programs and the unfortunately anticipated Molnupiravir are suggested to share in evolution of potentially more virulent variants e.g. delta plus and the catastrophic Indian surge of mortality is also anticipated to be repeated elsewhere unless some prompt interventions are considered.

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