Reviews of "A rigorous framework for detecting SARS-CoV-2 spike protein mutational ensemble from genomic and structural features "

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Summary of Reviews: *In this study, the authors leverage computational approaches to track and structurally describe emerging SARS-CoV-2 mutations. Authors report the presence of three novel mutational clusters, namely A222, N439, and S477. Reviewers deem the study thorough and reliable.*

Reviewer 1 (Dalal Harkati) | ☐☐☐☐

Reviewer 2 (Basil Saleh) | ☐☐☐☐

RR:C19 Strength of [Evidence Scale Key](#)

☐ ☐☐☐ = Misleading

☐☐ ☐☐ = Not Informative

☐☐☐ ☐ = Potentially Informative

☐☐☐☐ = Reliable

☐☐☐☐☐ = Strong

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