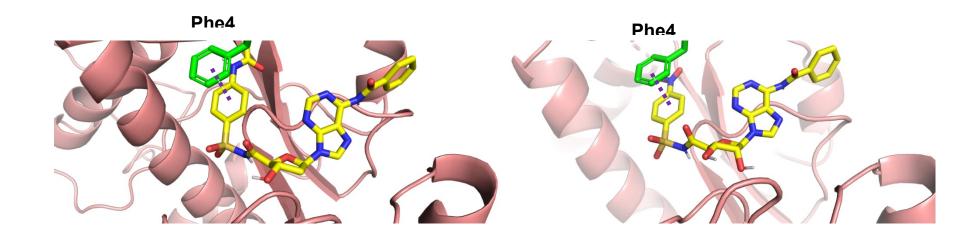
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Figure S1. Modeling results in the cap-binding pocket of SARS-CoV nsp14 with 7d (left) and 8a (right). Phe426 is shown in green. π - π stacking interactions (purple) are shown.



Modeling results in the cap-binding pocket of SARS-CoV nsp14 with 7d (left) and 8a (right). Phe426 is shown in green. π - π stacking interactions (purple) are shown.