

RESEARCH ARTICLE

Inhibitory Potential of Chitosan Derivatives against Severe Acute Respiratory Syndrome Coronavirus 2: An *In Silico* Prospective

Supplementary Files

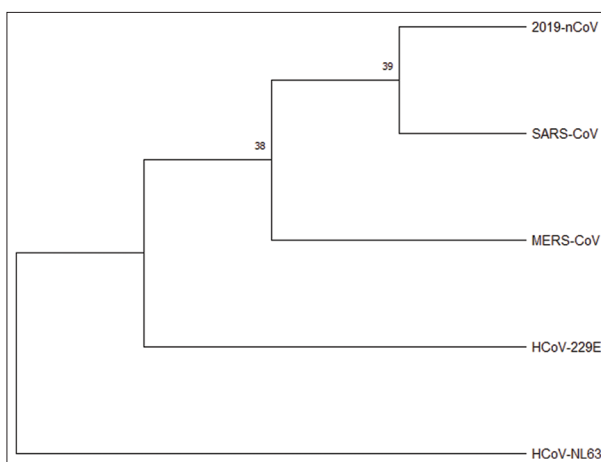


Figure S1. Phylogenetic analysis of severe acute respiratory syndrome coronavirus 2.

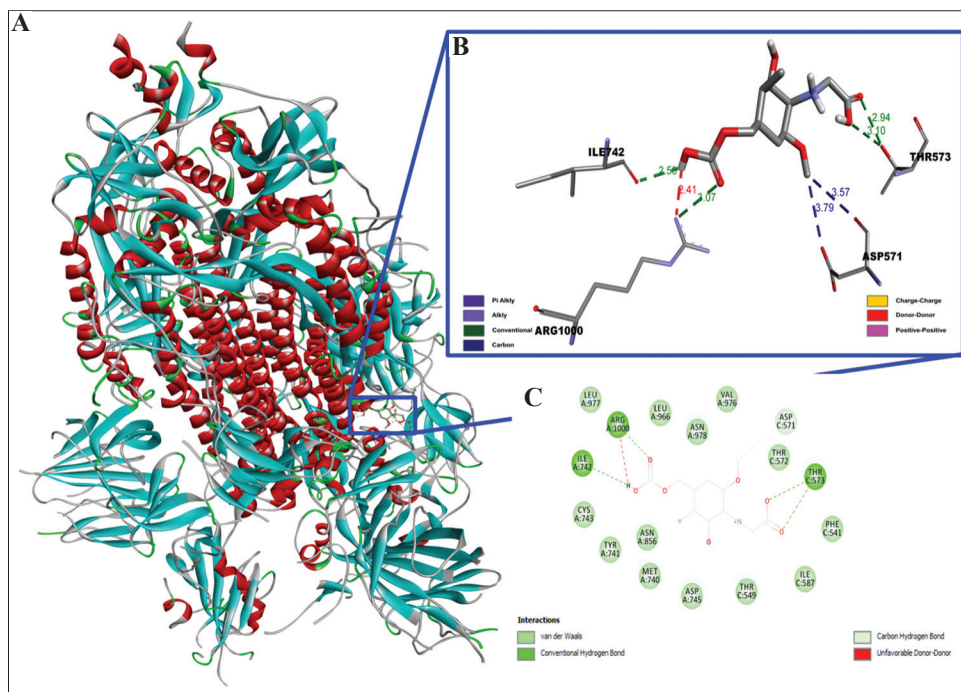


Figure S2. (A) Molecular docking of N,O-carboxy methyl chitosan with the S protein. (B) Brief 3D representation of N,O-carboxy methyl chitosan and S protein interaction, including bond types and length. (C) 2D representation of N,O-carboxy methyl chitosan and S protein with participating amino acids and types of bonds.

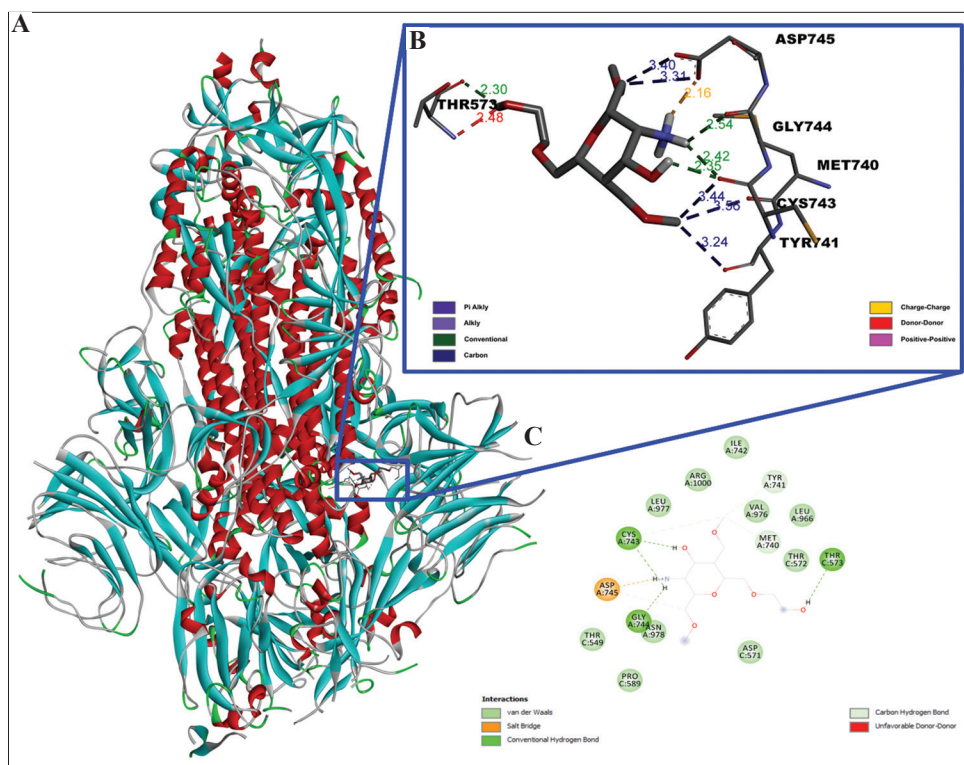


Figure S3. (A) Molecular docking of glycerol chitosan with the S protein. (B) Brief 3D representation of glycerol chitosan and S protein interaction, including bond types and length. (C) 2D representation of glycerol chitosan and S protein with participating amino acids and types of bonds.

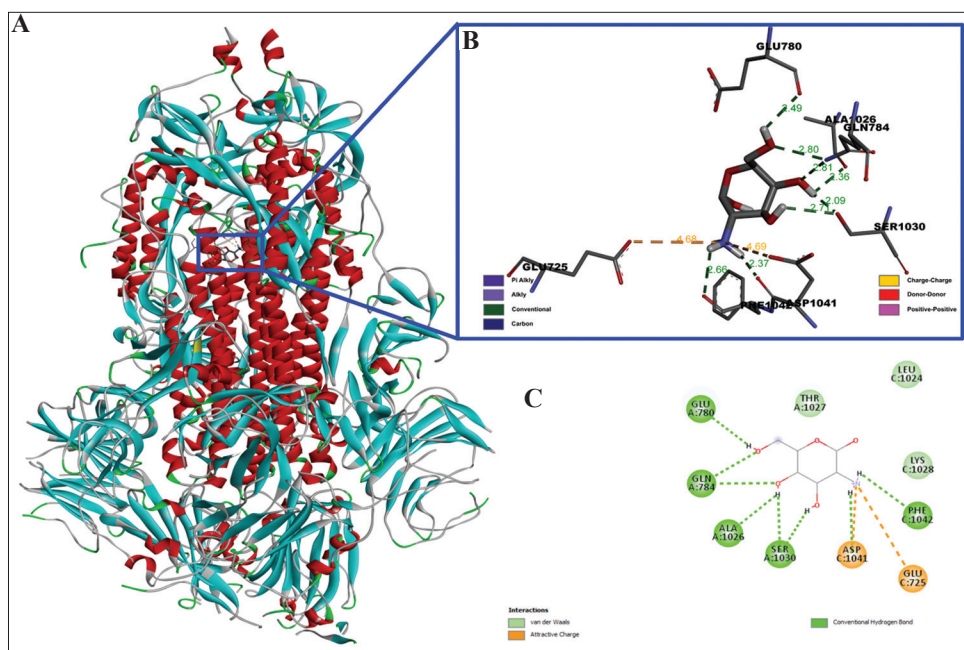


Figure S4. (A) Molecular docking of monomeric chitosan with the S protein. (B) Brief 3D representation of monomeric chitosan and S protein interaction, including bond types and length. (C) 2D representation of monomeric chitosan and S protein with participating amino acids and types of bonds.

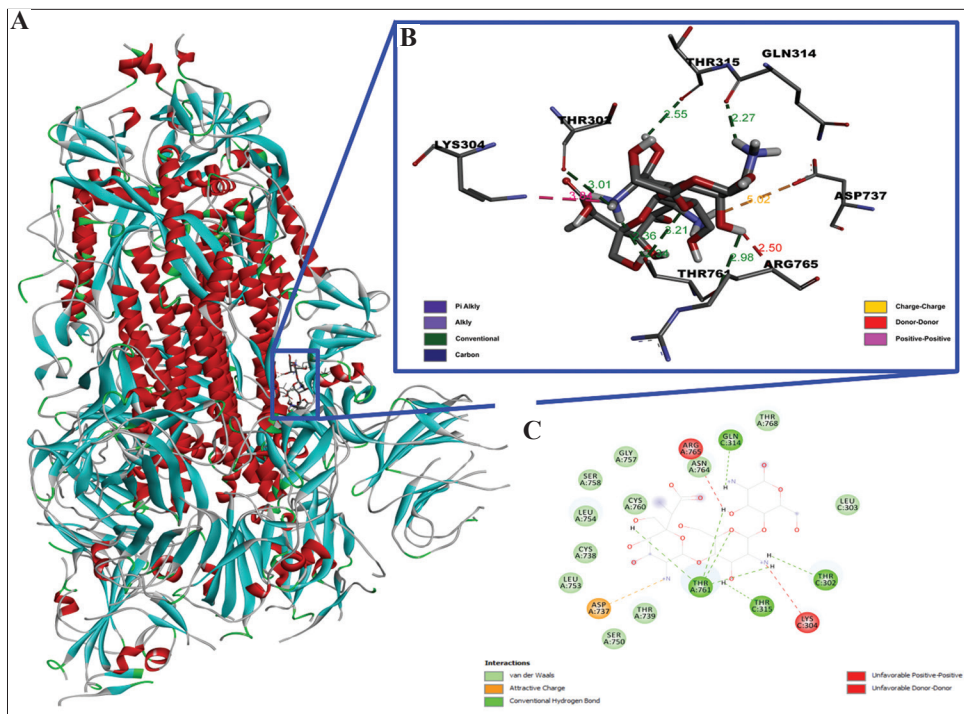


Figure S5. (A) Molecular docking of carboxy ethyl chitosan with the S protein. (B) Brief 3D representation of carboxy ethyl chitosan and S protein interaction, including bond types and length. (C) 2D representation of carboxy ethyl chitosan and S protein with participating amino acids and types of bonds.

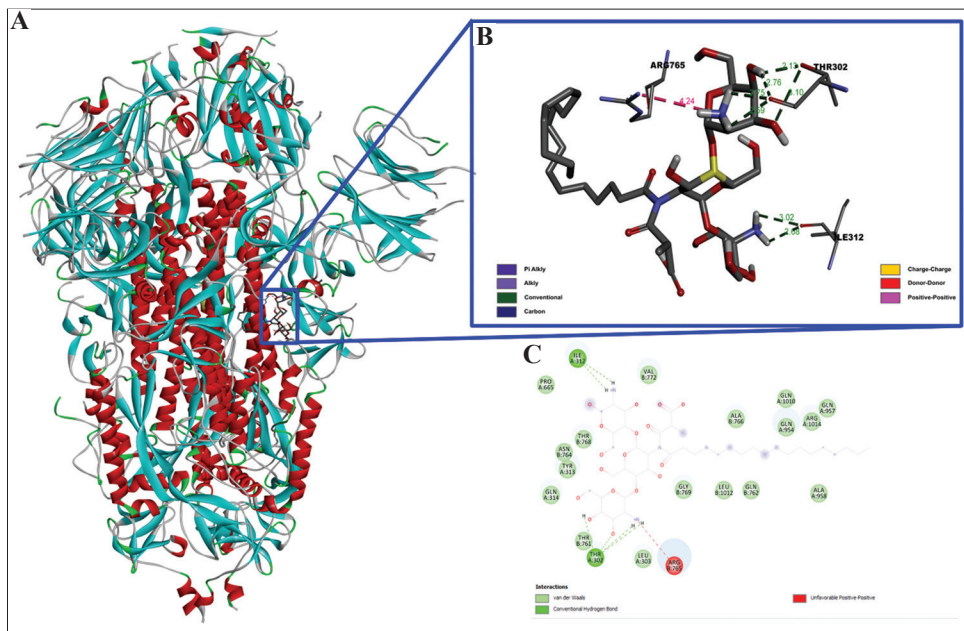


Figure S6. (A) Molecular docking of N-octadecanoyl-N-3-carboxy propionyl chitosan with the S protein. (B) Brief 3D representation of N-octadecanoyl-N-3-carboxy propionyl chitosan and S protein interaction, including bond types and length. (C) 2D representation of N-octadecanoyl-N-3-carboxy propionyl chitosan and S protein with participating amino acids and types of bonds.

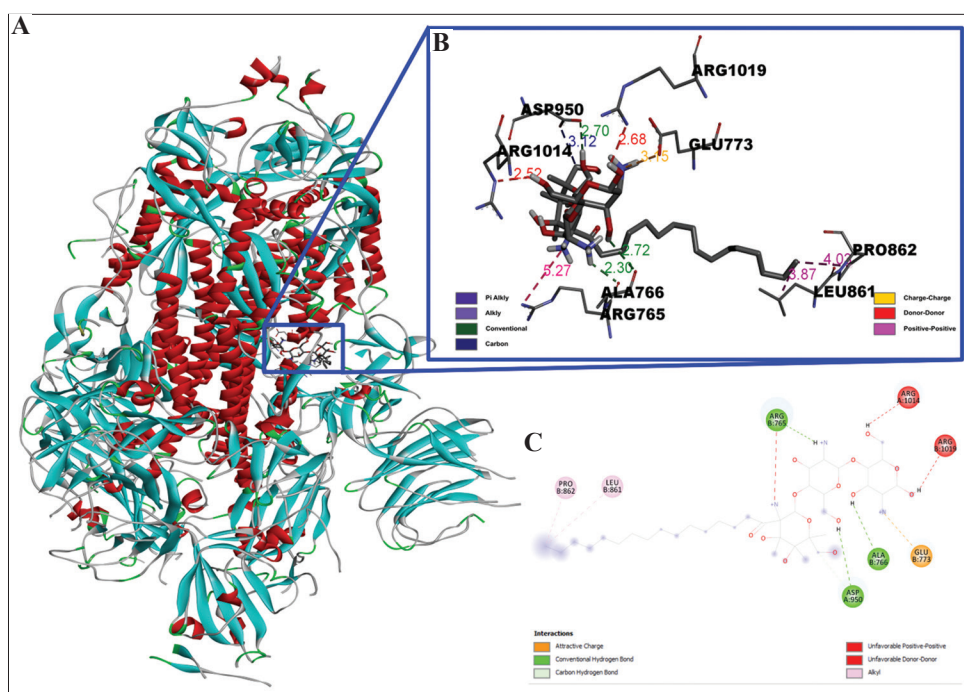


Figure S7. (A) Molecular docking of palmitoyl-trimethyl-chitosan with the S protein. (B) Brief 3D representation of palmitoyl-trimethyl-chitosan and S protein interaction, including bond types and length. (C) 2D representation of palmitoyl-trimethyl-chitosan and S protein with participating amino acids and types of bond.



Figure S8. Protein-protein interaction depicting five lowest binding energy for S Protein-ACE2 complex in the presence mHTCC. Green- lowest binding energy 1, Yellow- lowest binding energy 2, Blue- lowest binding energy 3, Cyans- lowest binding energy 4, Magentas- lowest binding energy 5, and Red- S Protein.