Supplementary Material

Table 1. Average geometrical values (Å) over the last 70 ns of 100-ns-long MD simulations.

System	RMSD	RG
SARS-CoV-2_{TA}	1.9 ± 0.2	2.58 ± 0.01
SARS-CoV-2_{TF3}	1.8 ± 0.2	26.0 ± 0.07
SARS-CoV-2_{TF2B}	2.0 ± 0.1	25.8 ± 0.07
SARS-CoV_{TA}	2.5 ± 0.2	26.1 ± 0.06
SARS-CoV_{TF3}	2.7 ± 0.2	25.9 ± 0.06
SARS-CoV_{TF2B}	2.8 ± 0.2	25.8 ± 0.07

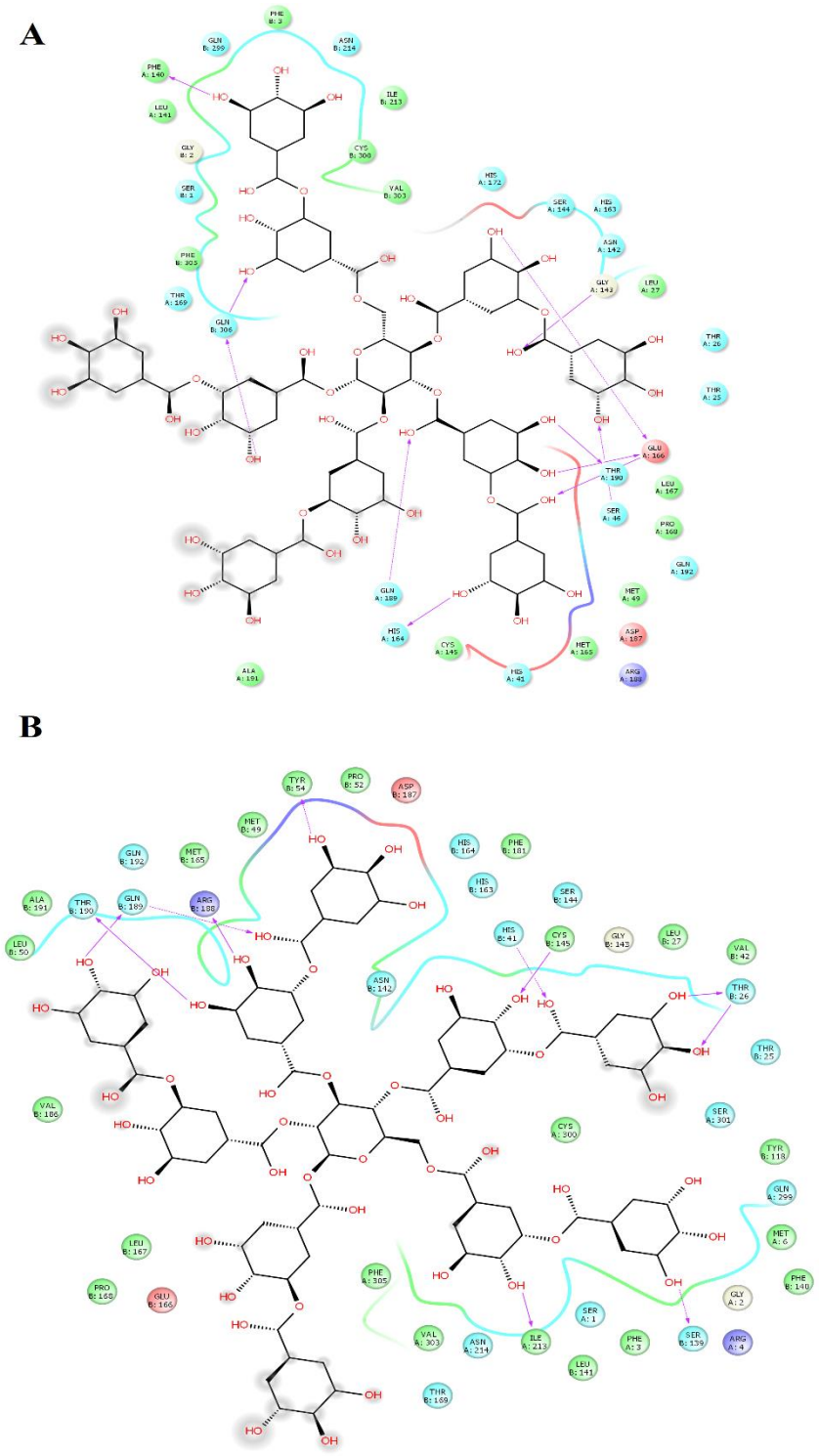

Figure 1. Interactions of the docked complexes between tannic acid and 3CL^{pro} of SARS-CoV-2. Map of interaction of tannic acid at subunit 1 (A) and subunit 2 (B).

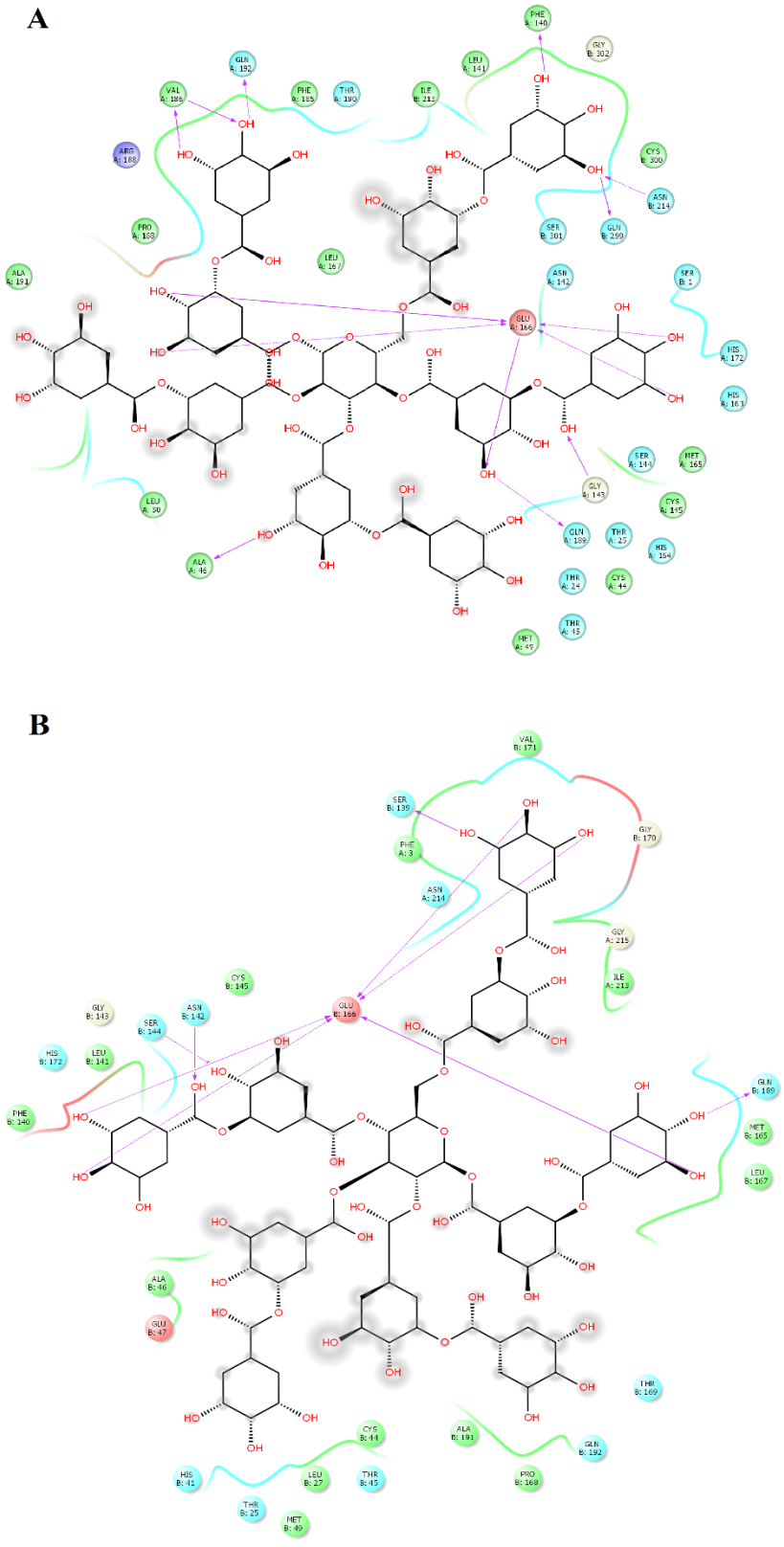

Figure 2. Interactions of the docked complexes between tannic acid and 3CL^{pro} of SARS-CoV. Map of interaction of tannic acid at subunit 1 (A) and subunit 2 (B).

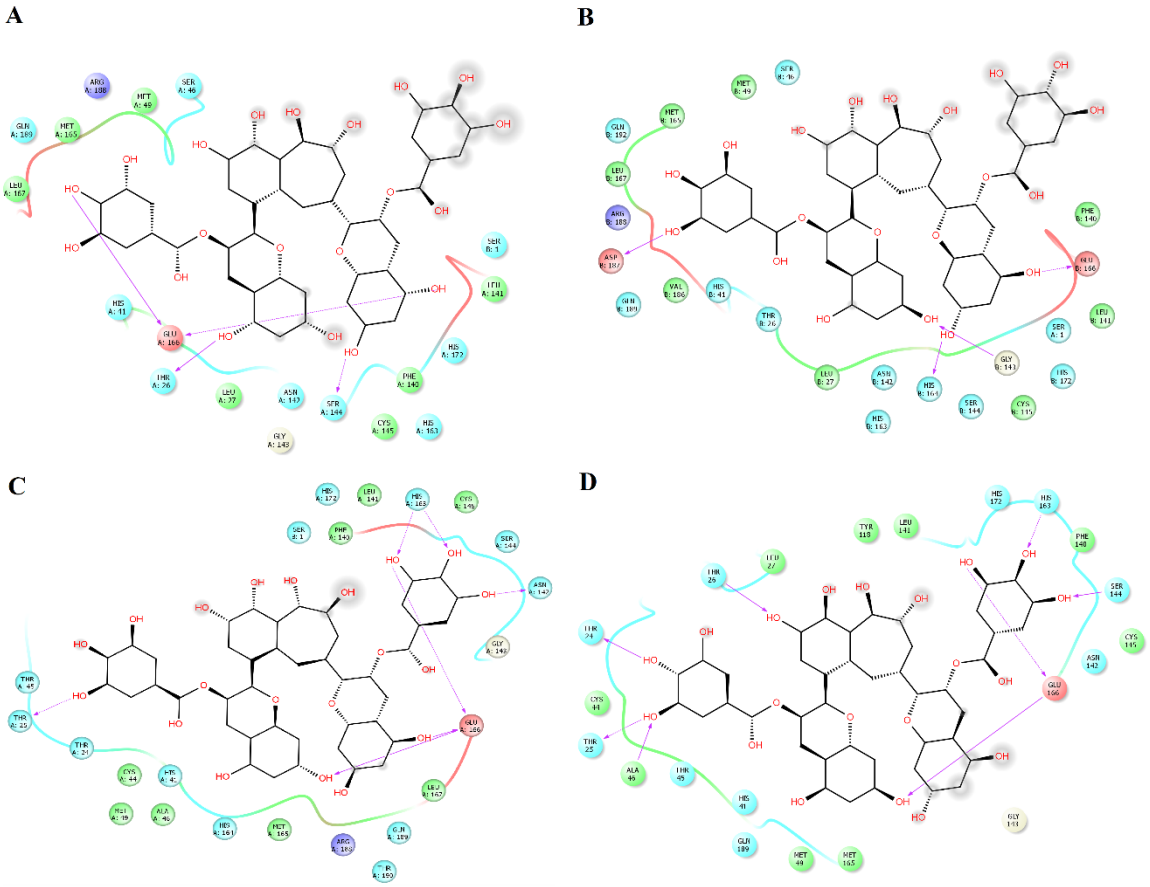

Figure 3. Interactions of the docked complex between TF3 and 3CL^{pro} of SARS-CoV-2 and SARS-CoV. Map of interaction of TF3 at subunit 1 (A) and subunit 2 (B) of SARS-CoV-2. TF3 at subunit 1(C) and 2 (D) of SARS-CoV.

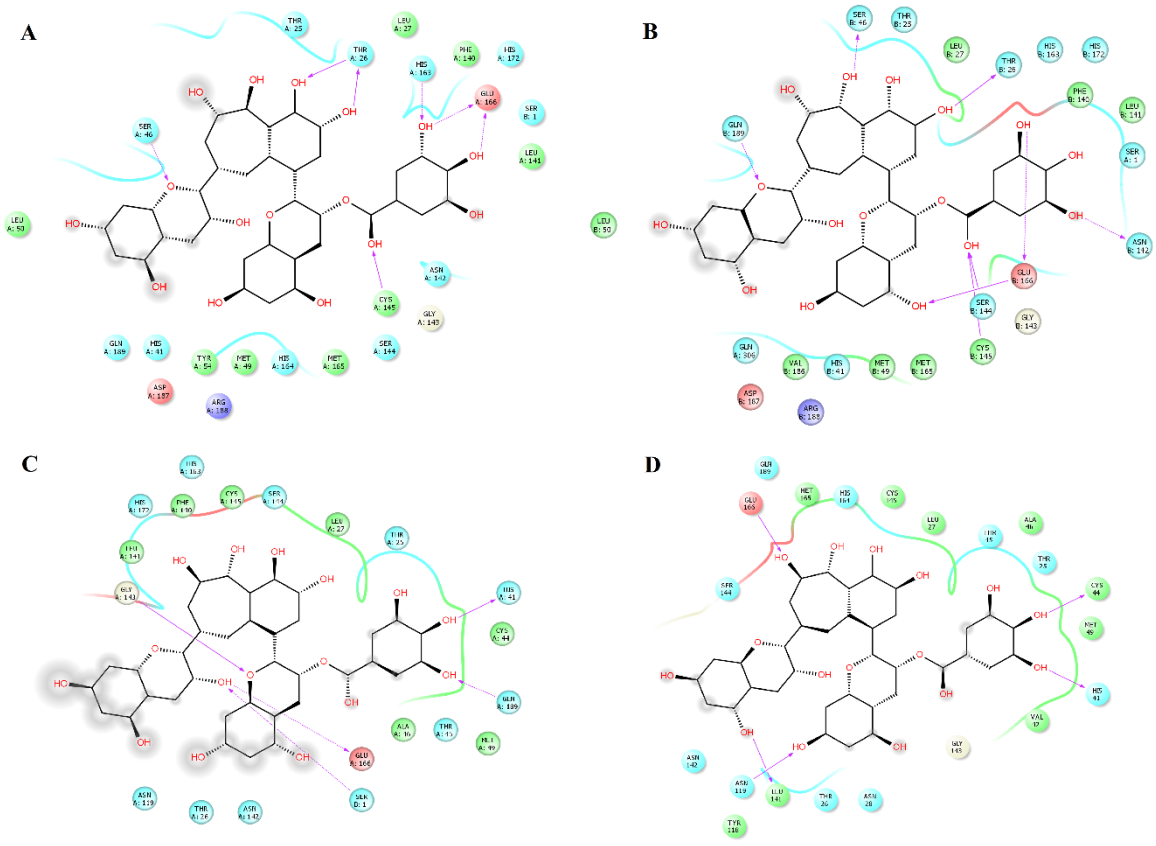

Figure 4. Interactions of the docked complex between TF2B and 3CL^{pro} of SARS-CoV-2 and SARS-CoV. Map of interaction of TF2B at subunit 1 (A) and subunit 2 (B) of SARS-CoV-2. TF2B at subunit 1(C) and 2 (D) of SARS-CoV.