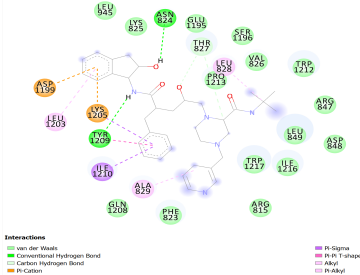
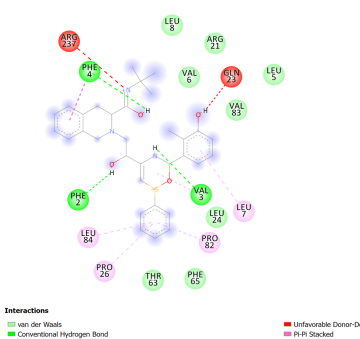
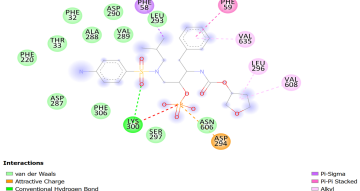


Supplementary Tables

Table S1: List of the ten top hits of antiviral drugs against SARS-CoV-2 Spike protein by order of lowest to highest binding energy value. Indinavir seems to exhibit the lowest binding energy (-9.8 kcal/Mol) when docked with Spike protein, whereas Cobicistat exhibits the highest binding energy (-6.37 kcal/Mol).

	Antivirals	Binding energy (kcal/Mol)	2D interaction	Interface amino acids	Type of interactions
1	Indinavir (C ₃₆ H ₄₇ N ₅ O ₄), 5362440	-9.8		Ala, 2Arg, Asn, 2Asp, Gln, Glu, 2Ile, 4Leu, 2Lys, Phe, Pro, Ser, Thr, 2Trp, Tyr	1Alk, 1CHB, 2HB, 1Pi-An, 3Pi-Alk, 1Pi-Cat, 2Pi-Sig, 1Pi-Pi-T, 14VW
2	Nelfinavir (C ₃₂ H ₄₅ N ₃ O ₄ S), 64143	-9		2Arg, Gln, 5Leu, 3Phe, 2Pro, Thr, 3Val	3HB, 5Pi-Alk, Pi-Pi-Stk, 7VW, 1UDD, 1UPP
3	Fosamprenavir (C ₂₅ H ₃₆ N ₃ O ₉ PS), 131536	-8.2		Ala, Asn, 3Asp, 5Phe, 2Leu, Lys, Ser, 3Val	1AC, 2Alk, 1HB, 1Pi-Alk, 1Pi-Pi-Stk, 1Pi-Sig, 1UPP, 11VW
4	Rintatolimod (C ₂₈ H ₄₀ N ₉ O ₂₅ P ₃), 135537060	-7.6		Arg, 3Asn, 2Asp, 2Gln, 2Glu, Gly, 2Ile, 2Lys, Pro, 2Ser, 1Val	1AC, 1CHB, 5HB, 14VW

			Interactions - van der Waals - Conventional Hydrogen Bond - Attractive Charge - Carbon Hydrogen Bond		
5	Loviride (C ₁₇ H ₁₆ Cl ₂ N ₂ O ₂), 3963	-7.3	Interactions - van der Waals - Conventional Hydrogen Bond - Carbon Hydrogen Bond - Pi-Sigma - Pi-Pi Stacked - Alkyl	Ala, Asn, 3Asp, Leu, Lys, 4Phe, Ser, Thr, 2Val	2Alk, 1CHB, 3HB, 1Pi-Pi- Stk, 1Pi-Sig, 9VW
6	Nevirapine (C ₁₅ H ₁₄ N ₄ O), 4463	-7.3	Interactions - van der Waals - Carbon Hydrogen Bond	Ala, Asn, 2Ile, 2Leu, Phe, Pro, Thr, 2Trp, Tyr, Val	1CHB, 12VW
7	Nitazoxanide (C ₁₂ H ₉ N ₃ O ₅ S), 41684	-7.1	Interactions - van der Waals - Conventional Hydrogen Bond - Pi-Sigma - Pi-Alkyl	2Arg, Asn, 2Gln, Phe, Pro, 4Leu, 3Val	4HB, 12Pi- Alk, 1Pi-Sig, 9VW
8	Imiquimod (C ₁₄ H ₁₆ N ₄), 57469	-6.8		Asn, Ile, 2Leu, Lys, Phe, Pro, Thr, Trp, Tyr, Val	2Alk, 2HB, 1Pi-Alk, 5VW

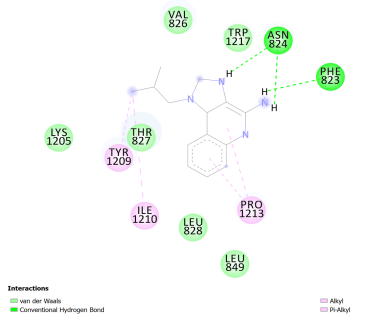
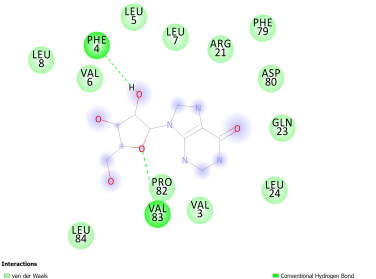
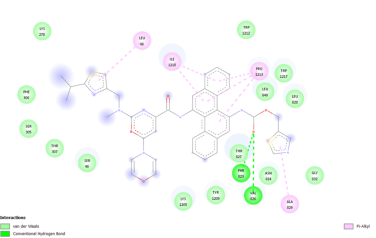
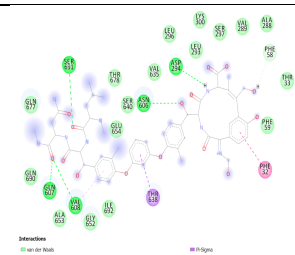
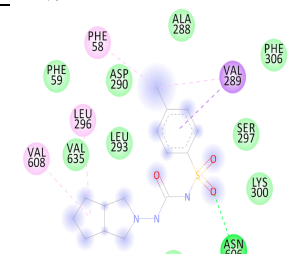
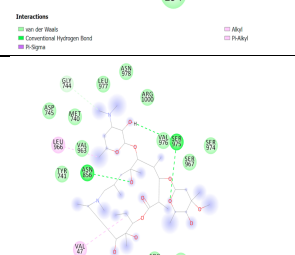
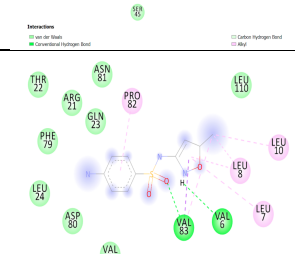
					
9	Inosine (C ₁₀ H ₁₂ N ₄ O ₅), 135398641	-6.5		Arg, Asp, 5Leu, 2Phe, Pro, 3Val	2HB, 12VW
10	Cobicistat (C ₄₀ H ₅₃ N ₇ O ₅ S ₂), 25151504	-6.37		1Ala, 1Asn, 1Gly, 1Ile, 3Leu, 2Lys, 2Phe, 1Pro, 2Ser, 2Thr, 2Trp, 1Tyr, 1Val	2HB, 7Pi- Alk, 14VW

Table S2: List of the ten top hits of antibiotics drugs against SARS-CoV-2 Spike protein by order of lowest to highest binding energy value. Vancomycin seems to exhibit the lowest binding energy (-10.2 kcal/Mol) when docked with Spike protein, whereas Levofloxacin exhibits the highest binding energy (-5.11 kcal/Mol).

	Antibiotics	Binding energy (kcal/Mol)	2D interaction	Interface amino acids	Type of interactions
1	Vancomycin (C ₆₆ H ₇₅ Cl ₂ N ₉ O ₂₄), 14969	-10.2		2Ala, Asn, Asp, 3Gln, Glu, Gly, Ile, 2Leu, Lys, 3Phe, 3Ser, 3Thr, 2Val	5HB, 1Pi-Alk, 1Pi-DHB, 1Pi-Pi-T, 1Pi-Sig, 17VW
2	Gliclazide (C ₁₅ H ₂₁ N ₃ O ₃ S), 3475	-8.6		Ala, Asn, 2Asp, 2Leu, Lys, 3Phe, Ser, 2Val	3Alk, 1HB, 1Pi-Alk, 1Pi-Sig, 9VW
3	Azithromycin (C ₃₈ H ₇₂ N ₂ O ₁₂), 447043	-7.9		2Arg, 2Asn, Asp, Gly, 2Leu, Met, Phe, 4Ser, Tyr, 3Val	2Alk, 1CHB, 3HB, 13VW
4	Sulfamethoxazole (C ₁₀ H ₁₁ N ₃ O ₃ S), 5329	-6.6		Arg, Asn, Asp, Gln, 5Leu, Phe, Pro, Thr, 3Val	3Alk, 2HB, 3Pi-Alk, 1Pi-Sig, 9VW

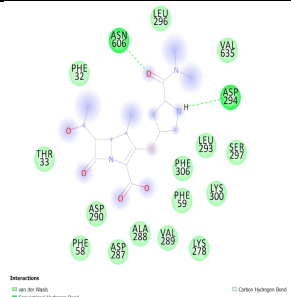
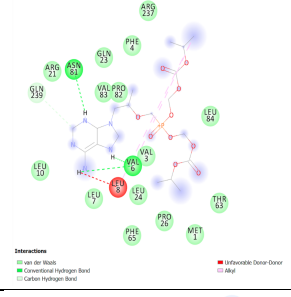
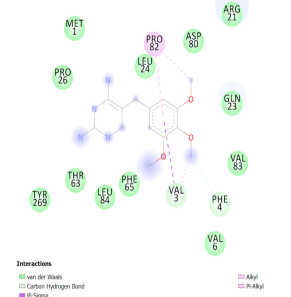
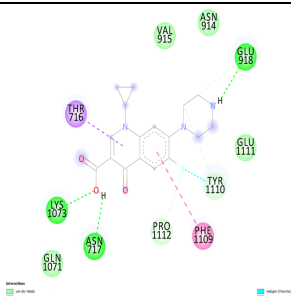
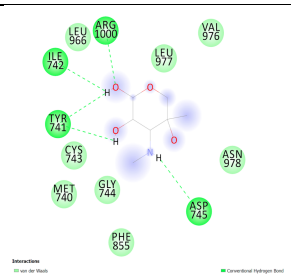
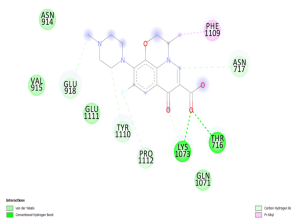
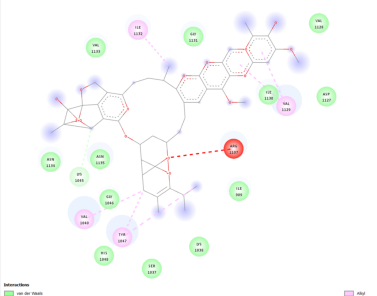
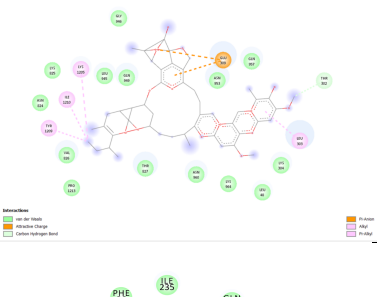
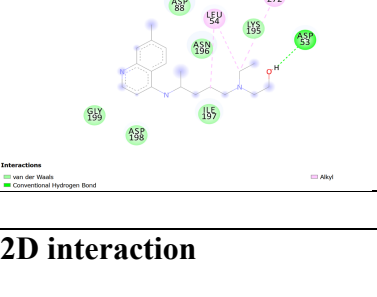
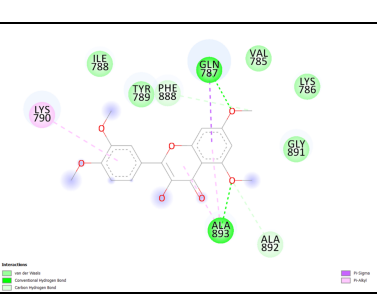
5	Meropenem (C ₁₇ H ₂₅ N ₃ O ₅ S), 441130	-6.4		Ala, Asn, 3Asp, 2Leu, 2Lys, 4Phe, Ser, Thr, 2Val	1CHB, 2HB, 15VW
6	Tenofovir Disoproxil (C ₁₉ H ₃₀ N ₅ O ₁₀ P), 5481350	-6.3		2Arg, Asn, 2Gln, 5Leu, Met, 2Phe, 2Pro, Thr, 3Val	2Alk, 1CHB, 2HB, 1UDD, 16VW
7	Trimethoprim (C ₁₄ H ₁₈ N ₄ O ₃), 5578	-6.1		Arg, Asp, Gln, 2Leu, Met, 2Phe, 2Pro, Thr, Tyr, 3Val	2Alk, 2CHB, 1Pi-Alk, 1Pi-Sig, 12VW
8	Ciprofloxacin (C ₁₇ H ₁₈ FN ₃ O ₃), 2764	-5.58		2Asn, 1Gln, 2Glu, 1Lys, 1Phe, 1Pro, 1Thr, 1Tyr, 1Val	3CHB, 1Hal, 3HB, 1Pi-Pi-T, 1Pi-Sig, 4VW
9	Gentamicin (C ₂₁ H ₄₃ N ₅ O ₇), 3467	-5.4		Arg, Asn, Asp, Cys, Gly, Ile, 2Leu, Met, Phe, Tyr, Val	5HB, 8VW
10	Levofloxacin (C ₁₈ H ₂₀ FN ₃ O ₄), 149096	-5.11		2Asn, 1Gln, 2Glu, 1Lys, 1Phe, 1Pro, 1Thr, 1Tyr, 1Val	5CHB, 2HB, 1Pi-Alk, 4VW

Table S3: List of the top hits antiparasitic drugs, flavonoids, and Vitamins against SARS-CoV-2 Spike protein by order of lowest to highest binding energy value. Ivermectin B1a seems to exhibit the lowest binding energy (-9.16 kcal/Mol) when docked with Spike protein, whereas Hydroxychloroquine exhibits the highest binding energy (-2.85 kcal/Mol). In terms of flavonoids Tetramethoxyflavone exhibit, the lowest binding energy (-4.91 kcal/Mol) and Galocatechin exhibits the highest binding energy (-2.88). Moreover, Vitamin D showed the lowest binding energy (-5.52 kcal/Mol) and Vitamin C the lowest binding energy (-2.95 kcal/Mol).

	Antiparasitic	Binding energy (kcal/Mol)	2D interaction	Interface amino acids	Type of interactions
1	Ivermectin B1a (C ₄₈ H ₇₄ O ₁₄), 6321424	-9.16		1Arg, 2Asn, 1Asp, 2Gly, 1His, 3Ile, 2Lys, 1Ser, 1Tyr, 4Val	4Alk, 1CHB, 2Pi-Alk, 1UPP, 12VW
2	Ivermectin B1b (C ₄₇ H ₇₂ O ₁₄), 6321425	-8.86		3Asn, 2Gln, 1Glu, 1Gly, 1Ile, 3Leu, 4Lys, 1Pro, 2Thr, 1Tyr, 1Val	2AC, 3Alk, 1CHB, 1Pi-Alk, 14VW
3	Hydroxychloroquine (C ₁₈ H ₂₆ ClN ₃ O), 3652	-2.85		Asn, 3Asp, Gln, Gly, 2Ile, Leu, Lys, Phe, Pro	2Alk, 1HB, 9VW
	Flavonoids	Binding energy (kcal/Mol)	2D interaction	Interface amino acids	Type of interactions
1	Tetramethoxyflavone (C ₁₉ H ₁₈ O ₆), 471721	-4.91		2Ala, 1Gln, 1Gly, 1Ile, 2Lys, 1Phe, 1Tyr, 1Val	2CHB, 2HB, 3Pi-Alk, 1Pi-Sig, 5VW

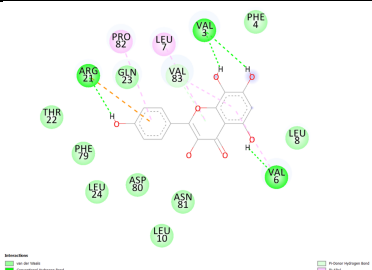
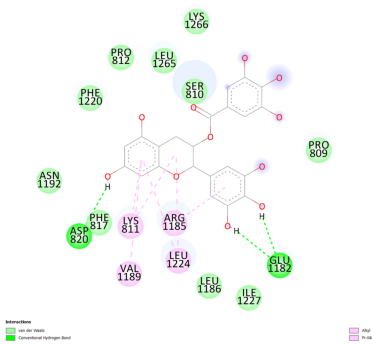
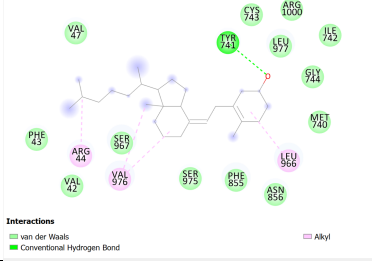
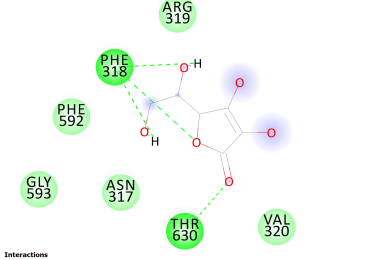
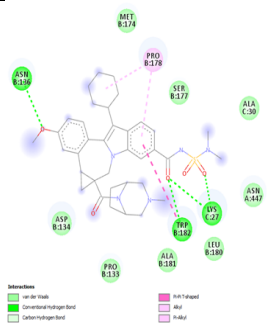
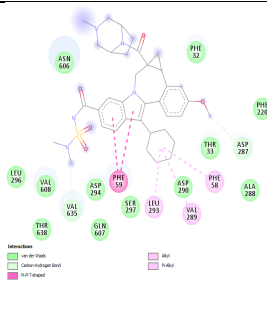
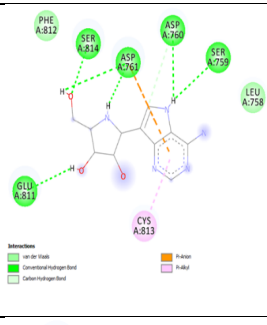
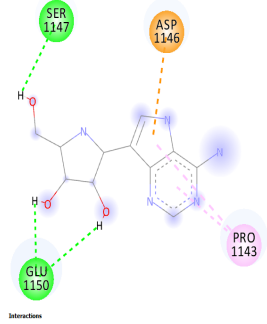
2	Herbacetin (C ₁₅ H ₁₀ O ₇), 5280544	-4.74		1Arg, 1Asn, 1Asp, 1Gln, 4Leu, 2Phe, 1Pro, 1Thr, 3Val	4HB, 4Pi-Alk, 1Pi-Cat, 1Pi-DHB, 9VW
3	Gallocatechin (C ₁₅ H ₁₄ O ₇), 65084	-2.88		1Arg, 1Asn, 1Asp, 1Glu, 1Ile, 3Leu, 2Lys, 1Ser, 2Phe, 2Pro, 1Val	2Alk, 3HB, 4Pi-Alk, 10VW
	Vitamins	Binding energy (kcal/Mol)	2D interaction	Interface amino acids	Type of interactions
1	Vitamin D (C ₂₇ H ₄₄ O), 5280795	-5.52		2Arg, Asp, Cys, Gly, Ile, 2Leu, Met, 2Phe, 2Ser, Tyr, 3Val	3Alk, 1HB, 12VW
2	Vitamin C (C ₆ H ₈ O ₆), 54670067	-2.95		Arg, Asn, Gly, 2Phe, Thr, Val	4HB, 5VW

Table S4: A comparative analysis between SARS-CoV-2 RdRp binding affinities and spike protein binding affinities in antivirals which primarily target viral RdRp. These comparisons include binding energy, the specific amino acids that are interacting with the drug, and the types of bonds formed during this interaction.

	Name and PubChem ID	Bound molecule	Binding Energy (kcal)	2D Interaction Plot	Interface Amino Acid	Type of Interactions
1	Beclabuvir (C36H45N5O5S), 49773361	RdRp	-5.63		2Ala, 2Asn, Asp, Leu, Lys, Met, 2Pro, Ser, Trp	4HB, 2Pi-Alk, 1Pi-Pi-T, 8VW
		Spike	-6.64		Ala, Asn, 3Asp, Gln, 2Leu, 4Phe, Ser, 2Thr, 3Val	2Alk, 2CHB, 1Pi-Alk, 2Pi-Pi-T; 12VW
2	Galidesivir (C11H15N5O3), 10445549	RdRp	-4.38		2Asp, Cys, Glu, Leu, Phe, 2Ser	1CHB, 6HB, 1Pi-Alk, 1Pi-Ani, 2VW
		Spike	-3.11		Asp, Glu, Pro, Ser	3HB, 2Pi-Alk, 1Pi-An

3	Ribavirin (C ₈ H ₁₂ N ₄ O ₅), 37542	RdRp	-4.2	Interaktionen: - von der Waals - Conventional hydrogen bond - Hydrophobic - Pi-H-T stacking - Pi-Allyl	2Ala, Arg, Asp, Ile, Leu, Lys, 2Phe, Val	5HB, 2Pi-Alk, 1Pi-Cat, 1Pi-Pi, 4VW
		Spike	-3.29	Interaktionen: - von der Waals - Conventional hydrogen bond	Arg, Cys, Gly, Ile, Leu, Phe, Pro, 3Thr	3HB, 7VW
4	Favipiravir (C ₅ H ₄ FN ₃ O ₂), 492405	RdRp	-3.09	Interaktionen: - von der Waals - Conventional hydrogen bond - Carbon hydrogen bond - Hydrogen (Pi-stacking) - Pi-Allyl - Pi-Allyl	Gly, His, Leu, Lys, 3Met, Phe, Ser, Thr	1CHB, 1Hal, 4HB, 1Pi-Alk, 1Pi-Sul, 5VW
		Spike	-2.94	Interaktionen: - von der Waals - Conventional hydrogen bond - Carbon hydrogen bond - Pi-Allyl	Gly, His, Ile, Pro, Ser, Thr, Tyr, Val	1CHB, 4HB, 1Pi-Alk, 4VW

5	Sofosbuvir (C ₂₂ H ₂₉ FN 3O ₉ P), 45375808	RdRp	-2.95		Ala, His, Lys, Phe, 3Ser, Thr	1Alk, 2HB, 2Pi-Alk, 3VW, 1UPP
		Spike	-3.7		Arg, Asn, Gln, Gly, Ile, Phe, Pro, Ser, Thr	1APS, 2HB, 1Pi-DHB, 5VW
6	Tenofovir (C ₉ H ₁₄ N ₅ O 4P), 464205	RdRp	-2.34		Ala, 3Asn, 2Asp, Gly, His, Lys, 2Ser	1CHB, 3HB, 1Pi-Alk, 9VW
		Spike	-2.75		Lys, 2Trp, 2Tyr	1CHB, 1HB, 2Pi-Alk, 1Pi- Cat, 2Pi-Pi- Stk, 2VW

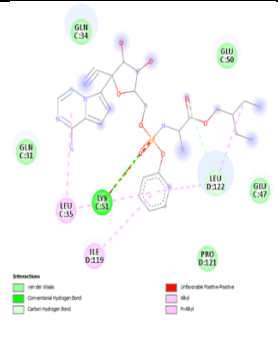
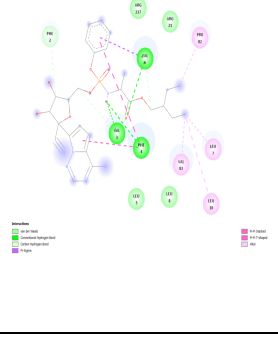
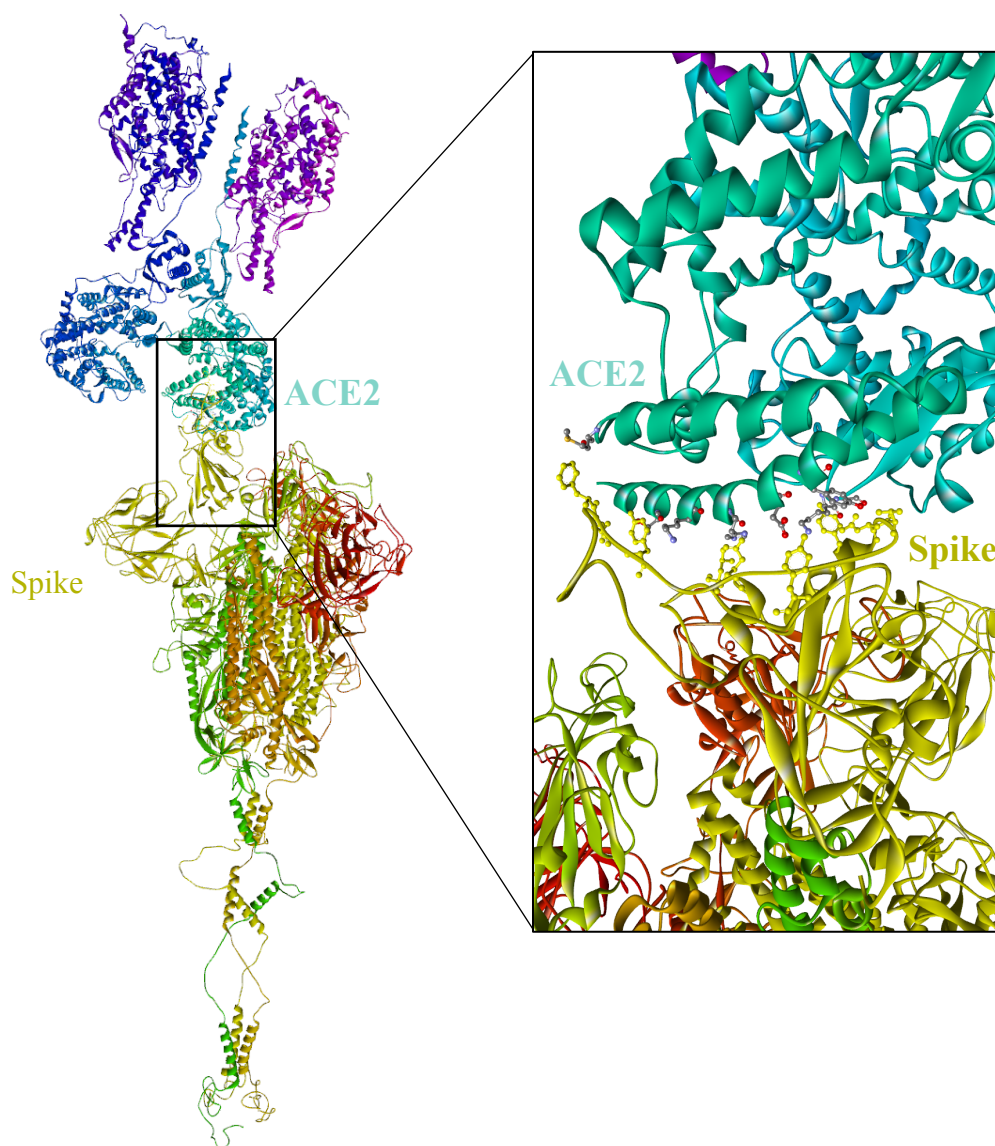
7	Remdesivir (C ₂₇ H ₃₅ N ₆ O ₈ P), 121304016	RdRp	-1.28		2Gln, 2Glu, Ile, 2Leu, Lys, Pro	1Alk, 1CHB, 1HB, 4Pi-Alk, 5VW, 1UPP
		Spike	-2.65		2Arg, 4Leu, 2Phe, 1Pro, 3Val	4Alk, 1CHB, 3HB, 1Pi-Pi-T, 1Pi-Pi-Stk, 1Pi-Sig, 4VW

Figure S1

A



B

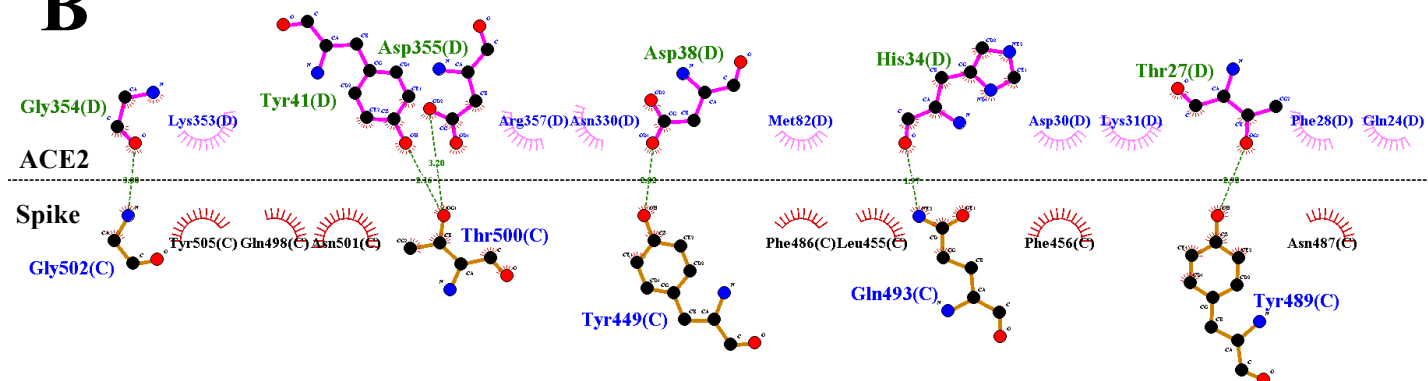
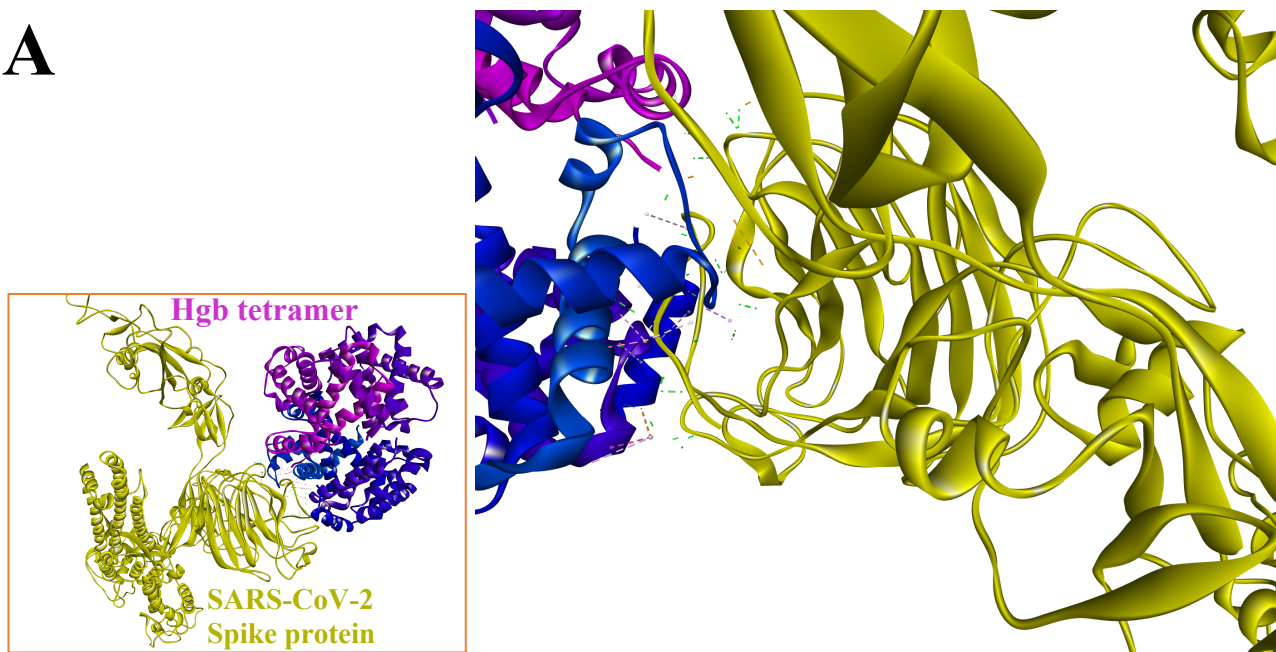


Figure S2

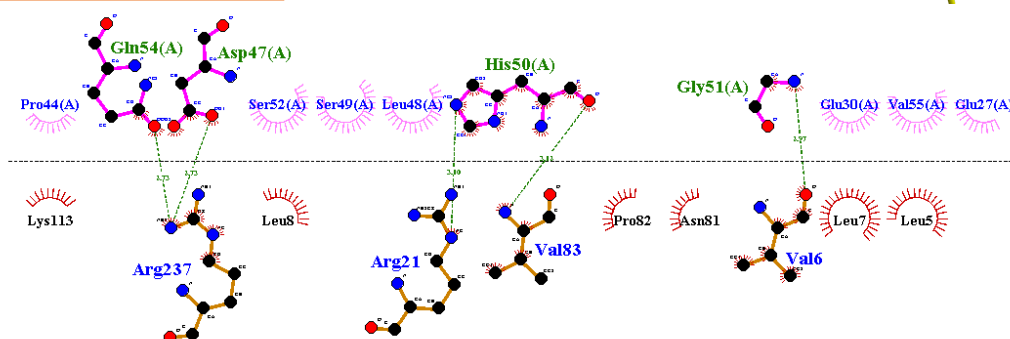
A



B

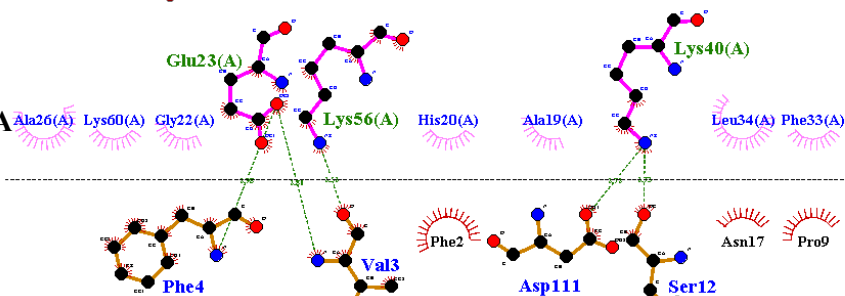
Hgb Chain A

Spike



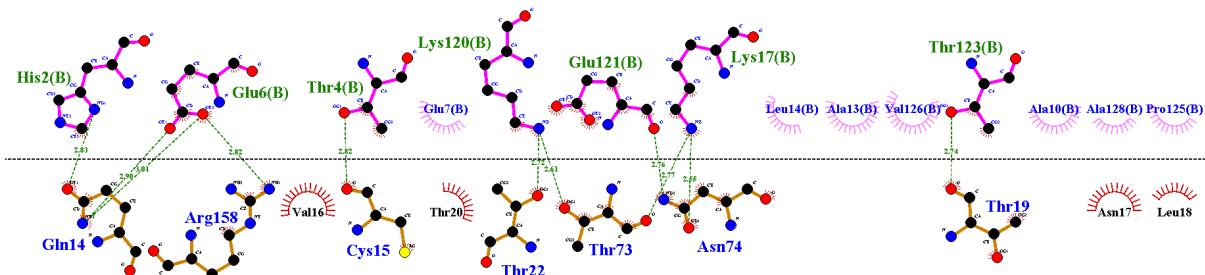
Hgb Chain A

Spike



Hgb Chain B

Spike



Hgb Chain D

Spike

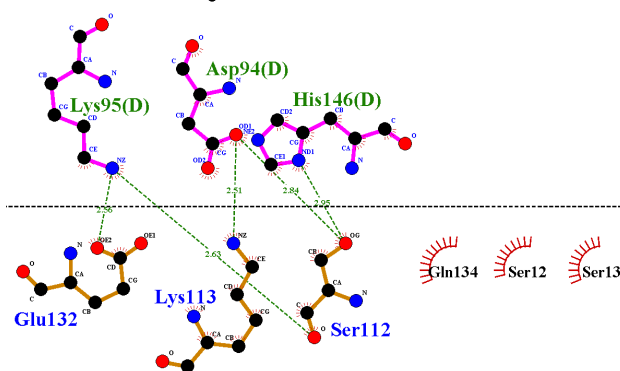
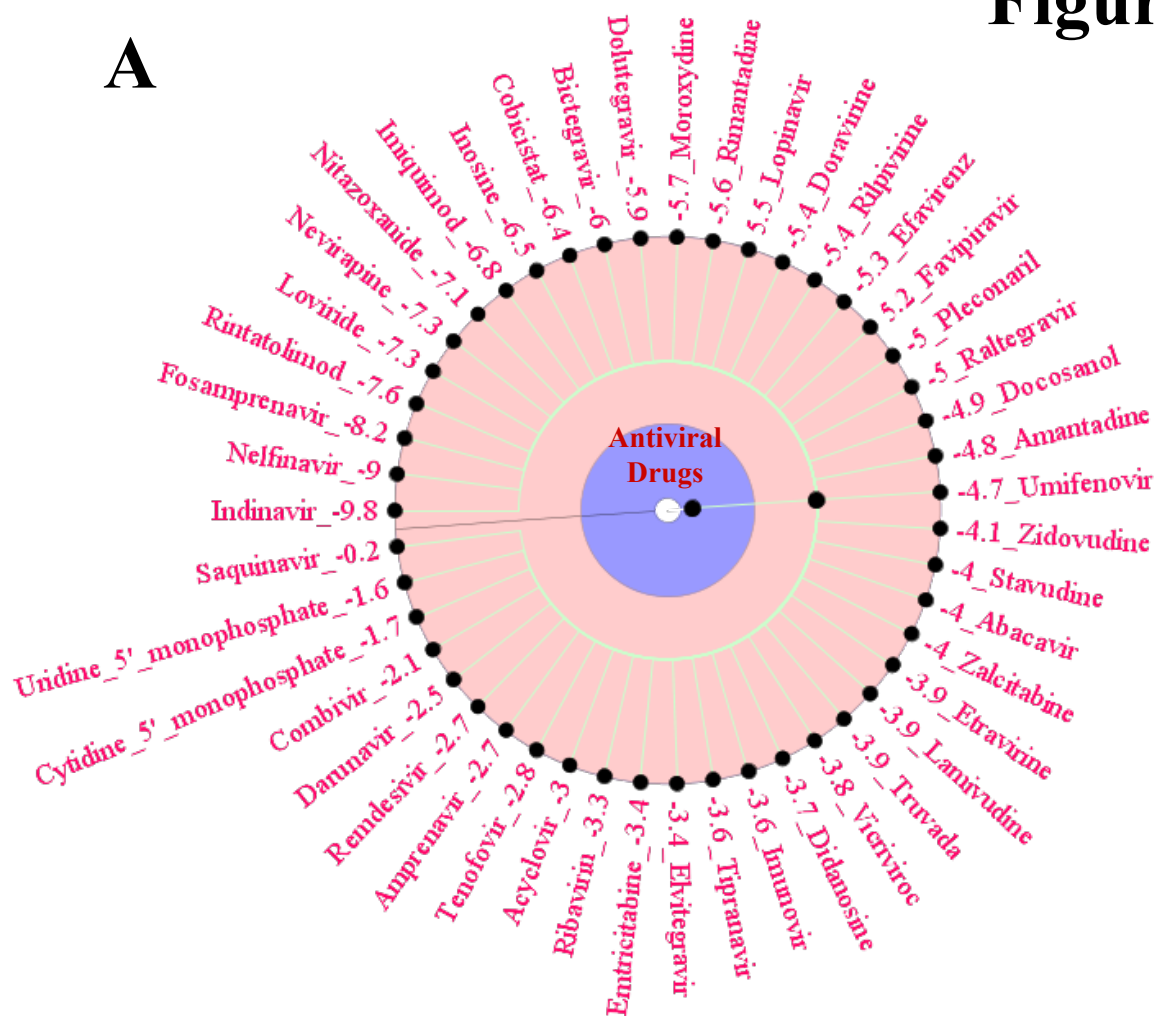


Figure S3

A



B

