

# Supplementary Materials: Lipophilicity, Pharmacokinetic Properties, and Molecular Docking Study on SARS-CoV-2 Target for Betulin Triazole Derivatives with Attached 1,4-Quinone

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**Table S1.** The experimental of lipophilicity ( $\log P_{\text{TLC}}$ ) and hydrophobic index ( $\phi_0$ ) for compounds **1–20**.

Compound	$\log P_{\text{TLC}}$	$\phi_0$	Compound	$\log P_{\text{TLC}}$	$\phi_0$
<b>1</b>	4.56	82.40	<b>11</b>	5.26	85.94
<b>2</b>	4.71	83.14	<b>12</b>	5.56	88.32
<b>3</b>	5.08	84.76	<b>13</b>	4.87	87.13
<b>4</b>	5.21	87.00	<b>14</b>	4.93	87.76
<b>5</b>	4.61	79.81	<b>15</b>	5.35	85.11
<b>6</b>	4.64	82.83	<b>16</b>	5.77	86.18
<b>7</b>	5.20	83.48	<b>17</b>	5.57	82.62
<b>8</b>	5.39	88.21	<b>18</b>	5.73	83.67
<b>9</b>	4.72	82.43	<b>19</b>	6.33	86.06
<b>10</b>	4.86	83.85	<b>20</b>	6.55	87.13

**Table S2.** The calculated lipophilicity for compounds **1–20**.

Com.	ALOGPs	AClogP	AlogP	XLOGP2	XLOGP3	milogP	iLOGP	WLOGP	MLOGP	SILICOS-IT
<b>1</b>	6.39	6.57	7.75	9.37	10.41	7.88	5.23	8.78	4.19	7.85
<b>2</b>	6.58	6.74	7.71	8.79	10.09	7.70	5.28	8.99	4.11	8.44
<b>3</b>	6.83	7.06	8.13	10.11	10.99	8.46	5.53	9.36	4.48	8.4
<b>4</b>	7.09	7.52	8.80	10.39	11.46	8.67	6.07	9.75	4.64	8.82
<b>5</b>	6.26	6.47	7.32	9.28	10.08	8.43	5.56	8.78	4.19	7.85
<b>6</b>	6.49	6.63	7.29	8.67	9.76	8.29	5.36	8.99	4.11	8.44
<b>7</b>	6.71	6.96	7.70	10.02	10.65	8.80	5.68	9.36	4.48	8.40
<b>8</b>	6.98	7.42	8.37	10.27	11.12	8.95	6.35	9.75	4.64	8.82
<b>9</b>	6.61	6.99	8.04	9.68	10.81	7.93	5.84	9.09	4.35	8.40
<b>10</b>	6.79	7.16	8.00	9.07	10.49	7.75	5.76	9.30	4.27	8.99
<b>11</b>	7.02	7.48	8.41	10.42	11.39	8.49	5.90	9.66	4.64	8.95
<b>12</b>	7.28	7.94	9.08	10.67	11.86	8.69	6.54	10.05	4.80	9.38
<b>13</b>	6.73	7.54	8.48	10.53	11.15	8.96	5.84	9.39	5.12	8.44
<b>14</b>	6.89	7.71	8.44	9.92	10.83	8.89	5.72	9.60	5.04	9.02
<b>15</b>	7.10	8.03	8.85	11.27	11.72	9.20	6.04	9.96	5.40	8.89
<b>16</b>	7.35	8.50	9.52	11.52	12.19	9.30	6.70	10.35	5.56	9.40
<b>17</b>	5.38	5.26	6.49	7.81	8.22	7.05	4.89	6.84	5.30	5.84
<b>18</b>	5.54	5.43	6.45	7.20	7.91	6.86	4.77	7.04	5.21	6.43
<b>19</b>	5.85	5.75	6.87	8.55	8.80	7.75	5.24	7.41	5.59	6.38
<b>20</b>	6.17	6.21	7.54	8.80	9.27	8.11	5.74	7.80	5.76	6.80

**Table S3.** The experimental and predicted logP.

Compound	Experimental logP <sub>TLC</sub>	Predicted logP	Residual
1	4.56	4.59	-0.027
2	4.71	4.64	0.064
3	5.08	5.09	-0.003
4	5.21	5.36	-0.156
5	4.61	4.59	0.024
6	4.64	4.64	0.005
7	5.20	5.09	0.113
8	5.39	5.36	0.024
9	4.72	4.78	-0.060
10	4.86	4.83	0.027
11	5.26	5.27	-0.016
12	5.56	5.55	0.013
13	4.87	4.90	-0.034
14	4.93	4.95	-0.022
15	5.35	5.40	-0.047
16	5.77	5.68	0.096

**Table S4.** The local minima of hybrids 1–16.

Compound	First area	Second area	Third area	Fourth area
1	-2.007	-2.721; -1.905	-2.231; -3.102; -2.612; -1.088	-2.993
2	-1.905	-2.721; -2.231	-2.286; -3.156; -2.612; -1.361	-2.612
3	-1.905	-2.721; -2.231	-2.395; -3.156; -2.667; -1.361	-1.905; -2.395
4	-1.905	-2.721; -2.231	-2.395; -3.156; -2.721; -1.361	-1.905; -2.503
5	-1.796	-2.612; -1.796	-2.068; -2.993; -2.503; -1.143	-2.721
6	-1.796	-2.503; -1.361	-2.286; -2.993; -2.612; -1.197	-2.612
7	-1.796	-2.612; -1.361	-2.286; -3.102; -2.612; -1.197	-1.905; -2.503
8	-1.796	-2.612; -1.361	-2.286; -3.102; -2.612; -1.252	-1.905; -2.503
9	-2.007	-2.721; -1.905	-2.231; -3.102; -2.612; -1.088	-2.993
10	-1.905	-2.721; -2.231	-2.286; -3.102; -2.612; -1.088	-2.612
11	-1.959	-2.721; -2.231	-2.449; -3.156; -2.721; -1.361	-1.905; -2.503
12	-1.959	-2.721; -2.231	-2.449; -3.156; -2.721; -1.361	-1.905; -2.449
13	-2.077	-2.068	-2.340; -3.102; -2.667; -1.252	-2.993
14	-2.068	-1.469	-2.340; -3.102; -2.667; -1.252	-2.667
15	-2.068	-1.633	-2.395; -3.102; -2.667; -1.252	-1.905; -2.340
16	-2.068	-1.469	-2.395; -3.102; -2.667; -1.361	-1.905; -2.503

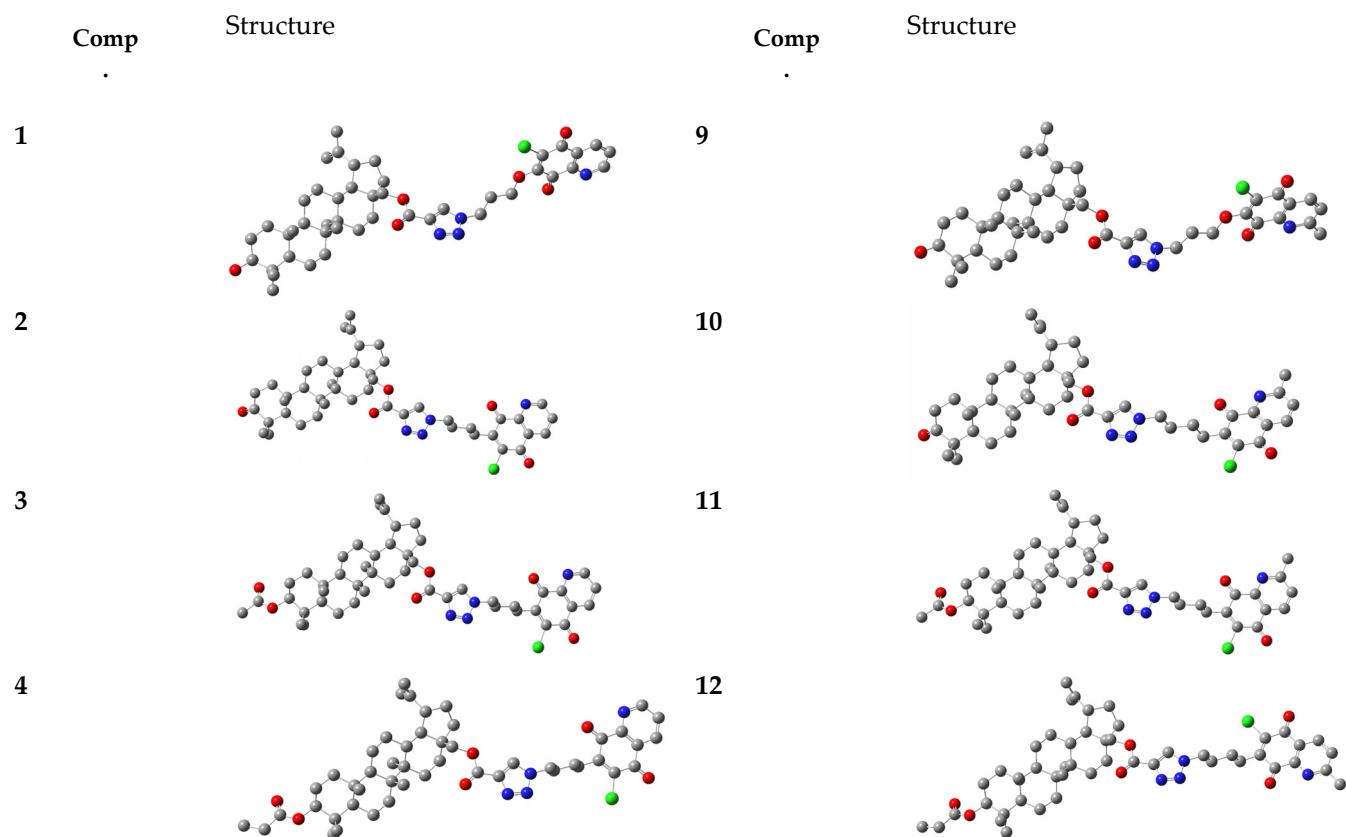
**Table S5.** Interaction of hybrids 1, 2, 6, 9, 10, 11 and 14 with active site of Mpro and PL protein.

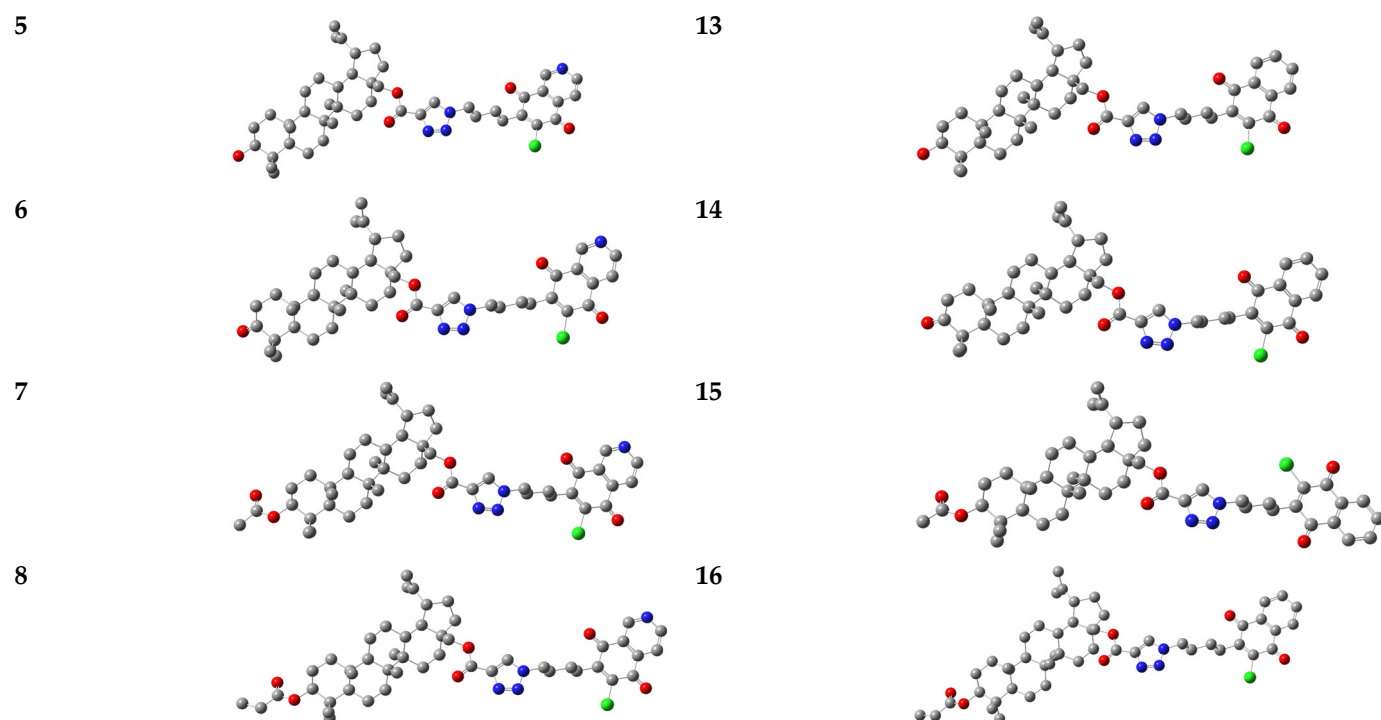
Protein.		Ligand		Interaction	
Name	Residue	Name	Residue	Type	Distance (Å)
Mpro	Gly143	2	carbonyl oxygen at C-8q	Conventional hydrogen bond	2.30
	His41		chlorine atom	Carbon hydrogen bond	2.97
	Asn142		carbonyl oxygen at C-8q	Carbon hydrogen bond	2.93
	Asn142		quinoline nitrogen	Carbon hydrogen bond	2.71
	Cys145		pyridine ring	$\pi$ -sulfur	3.62
	His41		pyridine ring	$\pi$ - $\pi$ stacked	4.74
	His41		benzoquinone ring	$\pi$ - $\pi$ stacked	5.73
	Cys145		benzoquinone ring	$\pi$ -alkyl	4.46
	Pro168		C-29	Alkyl-alkyl	3.62
	Pro168		C-30	Alkyl-alkyl	4.49

	Pro168	C-12	Alkyl-alkyl	4.77	
	Pro168	quinoline nitrogen	Carbon hydrogen bond	3.08	
	Asn142	triazole ring	Carbon hydrogen bond	3.24	
	Glu166	triazole ring	$\pi$ -anion	3.23	
	Leu141	triazole ring	Amide- $\pi$ stacked	4.59	
	Pro168	pyridine ring	$\pi$ -alkyl	4.96	
	Met49	6	C-30	Alkyl-alkyl	5.20
	Met165	C-30	Alkyl-alkyl	5.44	
	Met49	C-12	Alkyl-alkyl	3.71	
	Met49	C-29	Alkyl-alkyl	4.44	
	Met165	C-22	Alkyl-alkyl	4.83	
	His41	C-30	$\pi$ -alkyl	4.03	
	Gly143	carbonyl oxygen at C-8q	Conventional hydrogen bond	2.17	
	His41	chlorine atom	Carbon hydrogen bond	3.02	
	Asn142	carbonyl oxygen at C-8q	Carbon hydrogen bond	2.96	
	Asn142	quinoline nitrogen	Carbon hydrogen bond	2.59	
	Cys145	pyridine ring	$\pi$ -sulfur	3.55	
	His41	10	pyridine ring	$\pi$ - $\pi$ stacked	4.80
	His41	benzoquinone ring	$\pi$ - $\pi$ stacked	5.66	
	Cys145	benzoquinone ring	$\pi$ -alkyl	4.42	
	Pro168	C-29	Alkyl-alkyl	4.87	
	Pro168	C-30	Alkyl-alkyl	4.78	
	Pro168	C-12	Alkyl-alkyl	4.81	
	Asn142	C-28	Carbon hydrogen bond	3.49	
	Glu166	triazole ring	$\pi$ -anion	3.01	
	Leu167	pyridine ring	Amide- $\pi$ stacked	4.28	
	Leu167	benzoquinone ring	Amide- $\pi$ stacked	4.81	
	Pro168	pyridine ring	$\pi$ -Alkyl	3.96	
	Pro168	11	benzoquinone ring	$\pi$ -Alkyl	4.20
	Met49	C-30	Alkyl-alkyl	5.37	
	Met49	C-12	Alkyl-alkyl	4.13	
	Met165	C-29	Alkyl-alkyl	5.27	
	Met49	C-29	Alkyl-alkyl	3.62	
	His41	C-29	$\pi$ -Alkyl	3.85	
	Gln189	triazole ring	$\pi$ -sigma	2.45	
	His41	pyridine ring	$\pi$ - $\pi$ -stacked	4.53	
	His41	benzoquinone ring	$\pi$ - $\pi$ stacked	5.36	
	Met165	14	triazole ring	$\pi$ -Alkyl	4.80
	Cys145	pyridine ring	$\pi$ -Alkyl	4.81	
	Cys145	benzoquinone ring	$\pi$ -Alkyl	4.98	
	Pro168	C-21	Alkyl-alkyl	5.11	
PLpro	Thr301	betulin hydroxyl group	Conventional hydrogen bond	2.92	
	Asn267	carbonyl oxygen at C-8q	Carbon hydrogen bond	2.63	
	Tyr268	triazole ring	$\pi$ -sigma	2.62	
	Tyr268	triazole ring	$\pi$ - $\pi$ T-shaped	5.42	
	Pro248	1	pyridine ring	Alkyl-alkyl	4.22
	Pro248	benzoquinone ring	Alkyl-alkyl	4.27	
	Pro248	C-24	$\pi$ -alkyl	4.11	
	Pro248	C-23	$\pi$ -alkyl	4.21	
	Leu162	C-16	Alkyl-alkyl	4.78	
	Arg166	6	betulin carbonyl group	Conventional hydrogen bond	2.16
Tyr268	triazole ring	$\pi$ - $\pi$ T-shaped	5.38		

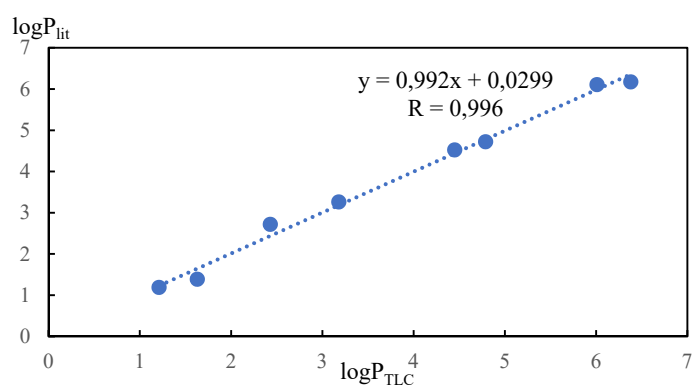
Pro247		benzoquinone ring	Amide- $\pi$ stacked	5.56
Pro248		pyridine ring	Alkyl-alkyl	4.40
Pro248		benzoquinone ring	Alkyl-alkyl	4.47
Pro248		C-24	$\pi$ -alkyl	4.13
Pro248		C-23	$\pi$ -alkyl	3.80
Leu162		C-16	Alkyl-alkyl	4.63
Leu162		C-22	Alkyl-alkyl	5.16
Arg166		betulin hydroxyl group	Conventional hydrogen bond	2.20
Tyr268		triazole ring	Carbon hydrogen bond	2.10
Tyr268		triazole ring	$\pi$ - $\pi$ T-shaped	4.96
Pro248		pyridine ring	$\pi$ -alkyl	4.55
Pro248	9	benzoquinone ring	$\pi$ -alkyl	5.14
Pro248		C-24	Alkyl-alkyl	4.42
Pro248		C-23	Alkyl-alkyl	4.54
Leu162		C-21	Alkyl-alkyl	5.22
Leu162		C-16	Alkyl-alkyl	4.77
Lys157		carbonyl oxygen at C-5q	Conventional hydrogen bond	2.17
Lys157		oxygen at C-7q	Conventional hydrogen bond	2.32
Gly163		carbonyl oxygen at C-5q	Carbon hydrogen bond	2.82
Leu162		benzene ring	$\pi$ -alkyl	4.75
Leu162		benzoquinone ring	$\pi$ -alkyl	4.07
Pro248	14	C-26	Alkyl-alkyl	4.33
Pro248		C-12	Alkyl-alkyl	4.99
Pro248		C-6	Alkyl-alkyl	3.81
Met208		C-30	Alkyl-alkyl	5.46
Tyr264		C-26	$\pi$ -alkyl	5.29
Tyr264		C-16	$\pi$ -alkyl	4.08

Figure S1. The optimized structure of hybrids 1-16.

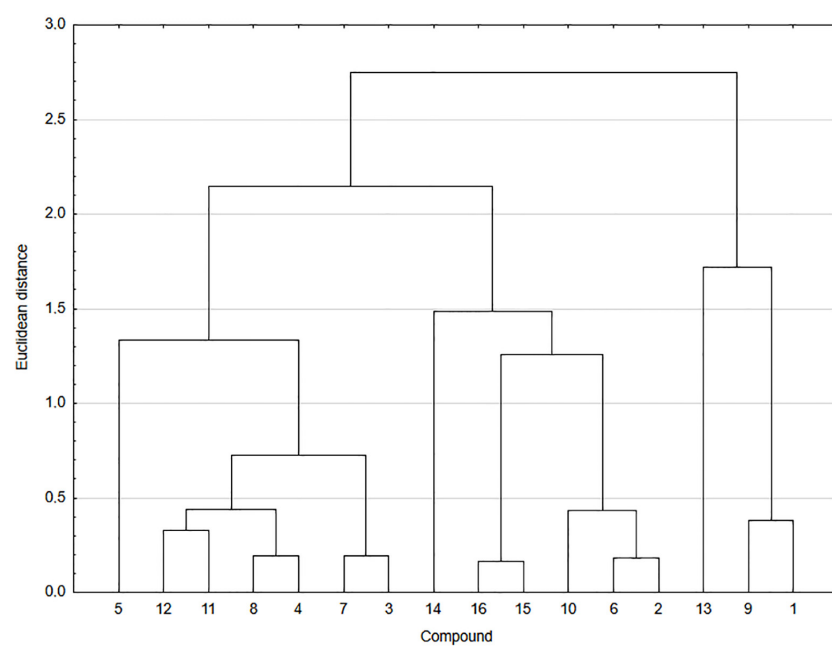




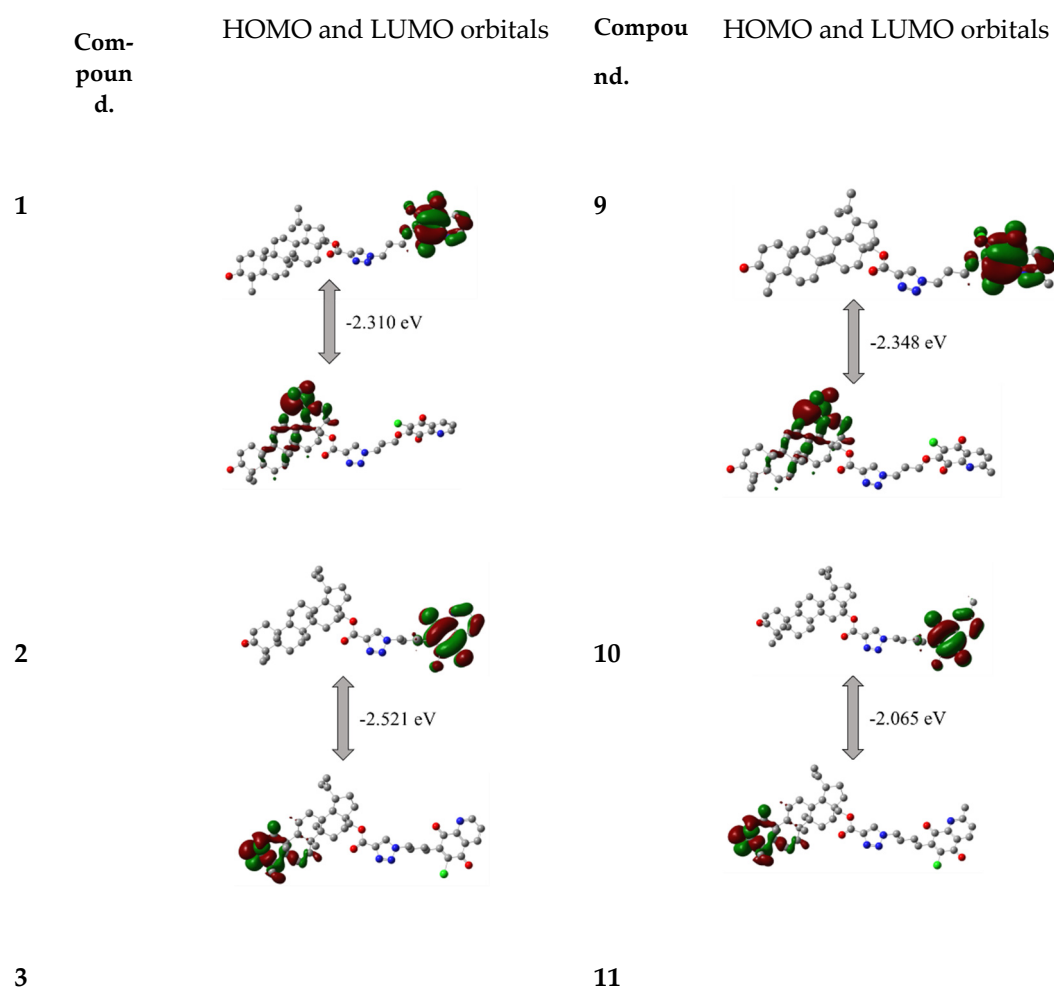
**Figure S2.** The linear regression between the experimental and literature lipophilicity for standard substance.

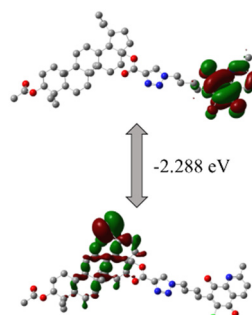
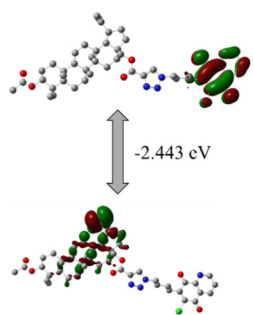


**Figure S3.** The similarity analysis of pharmacokinetic parameters of hybrids **1–16**.

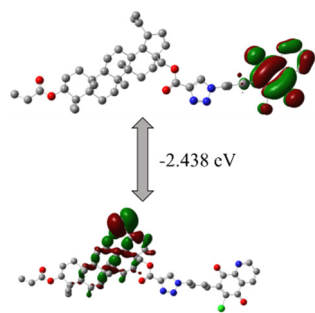


**Figure S4.** The HOMO and LUMO orbitals for hybrids **1–16**.

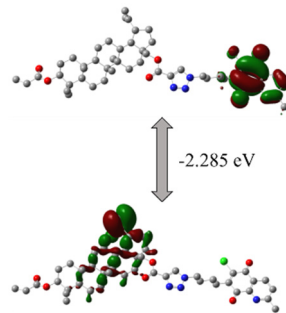




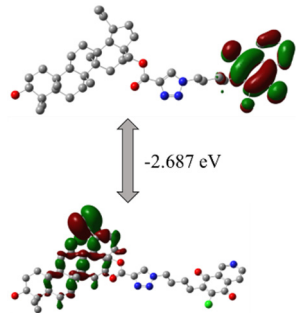
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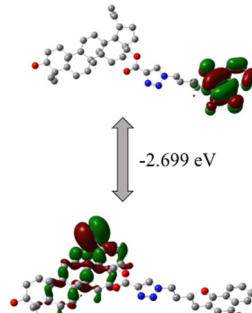
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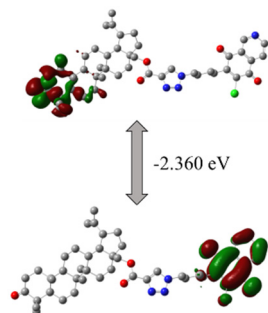
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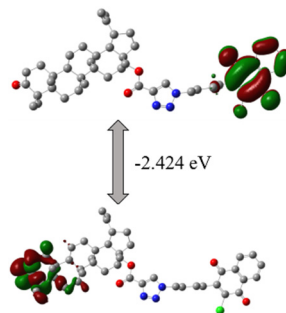
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14



7

15



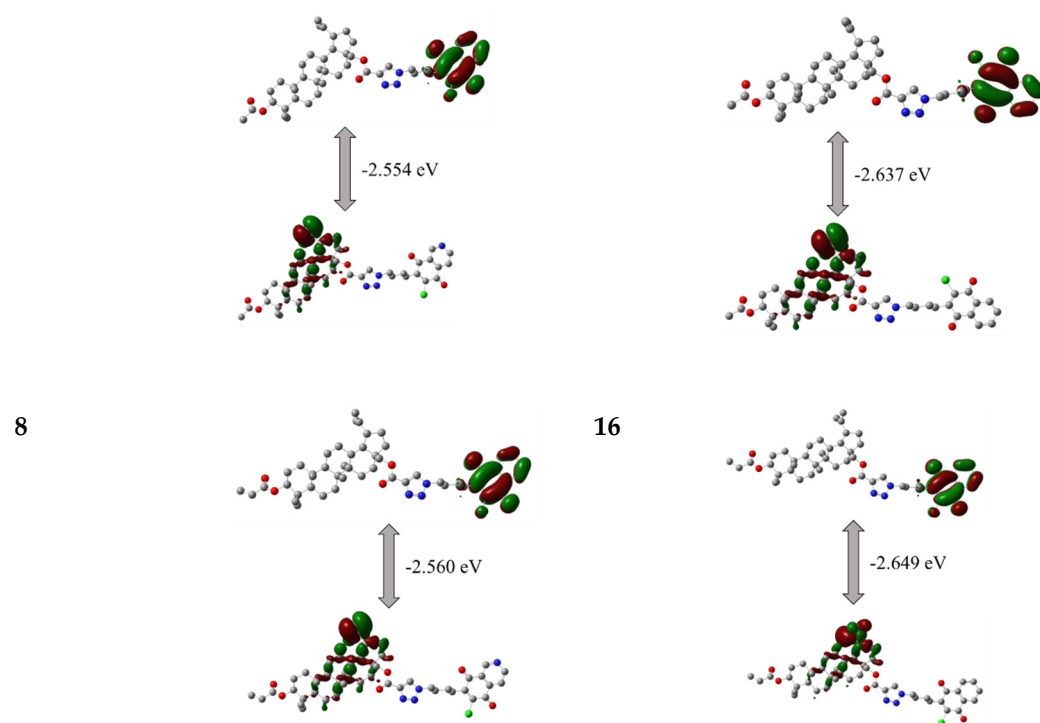
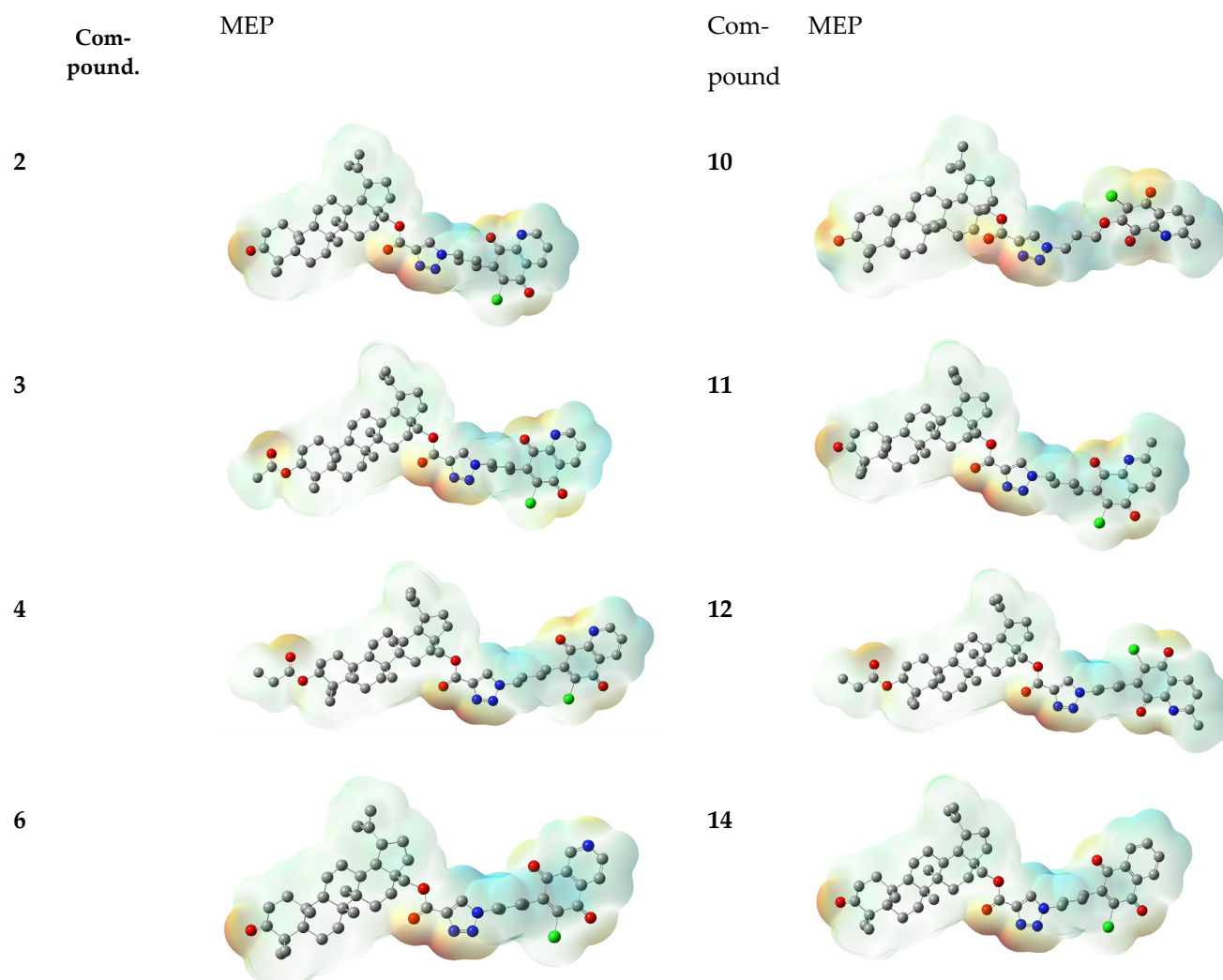
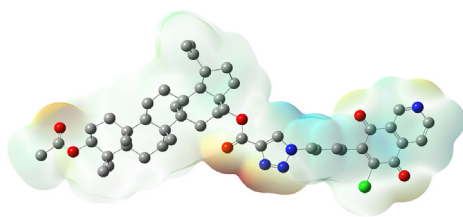


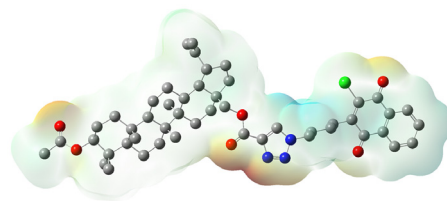
Figure S5. The MEP for hybrids 1–16.



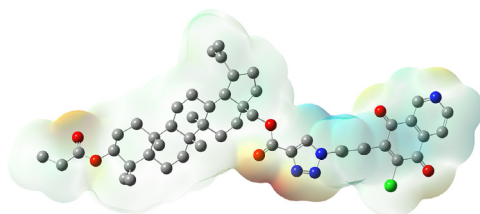
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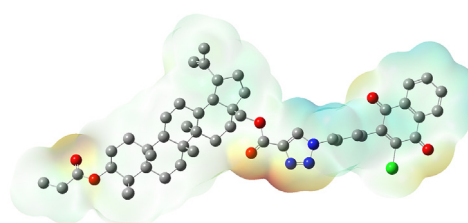
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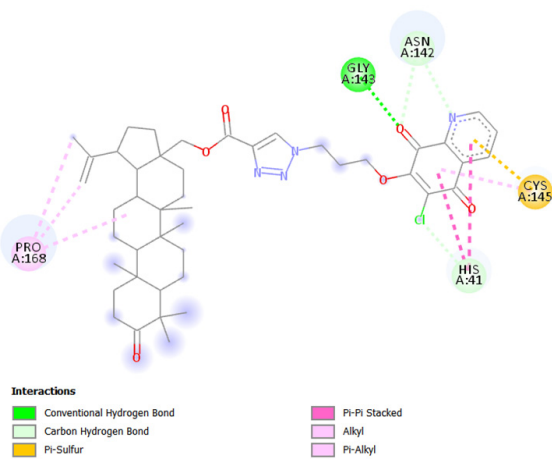


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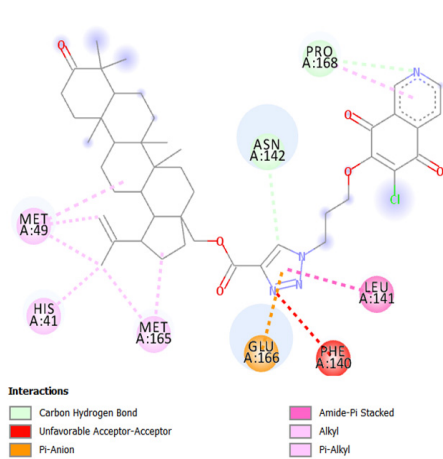


**Figure S6.** Docking pose of COVID-19 Mpro protein complex with hybrids 2 (A.), 6 (B.), 10 (C.), 11 (D.), and 14 (E.).

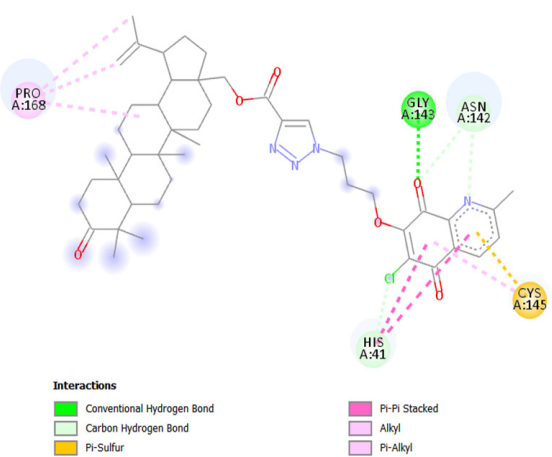
A.



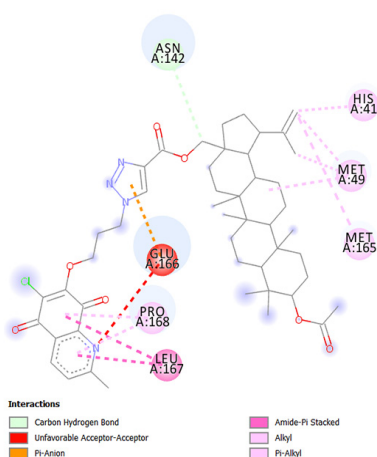
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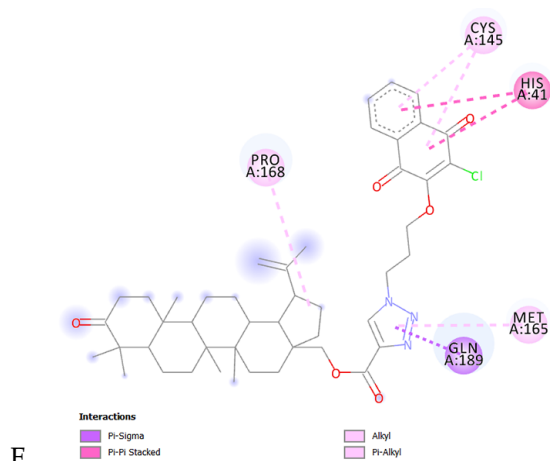


C.



D.





**Figure S7.** Docking pose of COVID-19 PLpro protein complex with hybrids **1** (A.), **6** (B.), **9** (C.), and **14** (D.).

