

# **Upgrading nirmatrelvir to prevent SARS-CoV-2 Mpro via DeepFrag and free energy calculations**

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Table S1. FPL results of the modified compounds versus PF-07321332 (nirmatrelvir).

Source	N°	Code	F <sub>Max</sub>	W	ΔG <sub>FPL</sub> <sup>Pre a</sup>
		<b>PF-07321332</b>	788.3 ± 38.1	103.7 ± 3.6	-11.32
	1	<b>PF_9b</b>	871.1 ± 43.7	122.1 ± 5.0	-12.35
	2	<b>PF_9a</b>	896.9 ± 55.5	119.8 ± 8.2	-12.22
	3	<b>PF_3</b>	890.7 ± 48.4	116.0 ± 4.3	-12.01
	4	<b>PF_35</b>	776.6 ± 61.3	115.4 ± 8.0	-11.97
<b>PF-07321332</b>	5	<b>PF_12</b>	759.7 ± 15.5	107.3 ± 3.0	-11.52
	6	<b>PF_33</b>	779.0 ± 56.3	104.5 ± 5.8	-11.37
	7	<b>PF_34</b>	787.6 ± 55.8	103.7 ± 6.8	-11.32
	8	<b>PF_2</b>	763.7 ± 69.0	97.5 ± 4.8	-10.97
	9	<b>PF_14</b>	689.8 ± 26.6	94.3 ± 4.7	-10.79

Table S2. FPL results of the modified compounds.

Source	N°	Code	F <sub>Max</sub>	W	ΔG <sub>FPL</sub> <sup>Pre a</sup>
<b>PF_9b</b>	1	<b>PF_9b_3</b>	730.9 ± 26.1	113.6 ± 3.4	-11.87
	2	<b>PF_9b_13</b>	788.2 ± 15.7	105.1 ± 1.9	-11.40
<b>PF_9a</b>	1	<b>PF_9a_33</b>	882.6 ± 56.9	130.1 ± 7.3	-12.80
	2	<b>PF_9a_11</b>	849.2 ± 18.2	117.3 ± 4.2	-12.08
<b>PF_9a</b>	3	<b>PF_9a_32</b>	781.5 ± 33.0	113.5 ± 5.3	-11.87
	4	<b>PF_9a_2</b>	803.9 ± 58.9	112.4 ± 4.0	-11.81
<b>PF_9a</b>	5	<b>PF_9a_13</b>	669.9 ± 21.5	92.0 ± 3.2	-10.67
	1	<b>PF_3_4</b>	827.5 ± 35.0	116.3 ± 2.7	-12.03
<b>PF_3</b>	2	<b>PF_3_34</b>	852.9 ± 35.1	114.9 ± 3.4	-11.95
	3	<b>PF_3_2</b>	764.6 ± 56.8	104.5 ± 5.0	-11.36
<b>PF_3</b>	4	<b>PF_3_13</b>	713.8 ± 30.2	98.30 ± 4.7	-11.02
	1	<b>PF_35_14</b>	627.1 ± 29.8	92.0 ± 3.8	-10.66
<b>PF_12</b>	1	<b>PF_12_33</b>	869.2 ± 6.6	121.9 ± 1.6	-12.34
	2	<b>PF_12_9</b>	877.5 ± 42.9	116.5 ± 5.0	-12.04
<b>PF_12</b>	3	<b>PF_12_3</b>	836.7 ± 42.5	114.6 ± 6.4	-11.93
	4	<b>PF_12_32</b>	833.3 ± 28.6	113.2 ± 4.2	-11.85
<b>PF_12</b>	5	<b>PF_12_34</b>	842.0 ± 44.8	112.0 ± 3.5	-11.79
	6	<b>PF_12_13</b>	654.9 ± 22.2	88.4 ± 2.3	-10.46

Table S3. FPL results of the modified compounds.

<b>Source</b>	<b>N<sup>0</sup></b>	<b>Code</b>	<b>F<sub>Max</sub></b>	<b>W</b>	<b>ΔG<sub>FPL</sub><sup>Pre a</sup></b>
<b>PF_9a_33</b>	1	<b>PF_9a_33_13</b>	828.4 ± 14.4	121.0 ± 2.6	-12.29
	2	<b>PF_9a_33_2</b>	787.7 ± 47.7	107.3 ± 6.2	-11.52
	3	<b>PF_9a_33_3</b>	806.5 ± 10.4	101.3 ± 5.1	-11.18
	4	<b>PF_9a_33_32</b>	689.0 ± 36.5	95.3 ± 6.0	-10.85
<b>PF_3_4</b>	1	<b>PF_3_4_62</b>	1032.2 ± 33.5	146.5 ± 5.2	-13.71
	2	<b>PF_3_4_32</b>	978.9 ± 57.6	144.5 ± 4.1	-13.60
	3	<b>PF_3_4_7</b>	957.6 ± 44.0	131.2 ± 3.0	-12.86
	4	<b>PF_3_4_33</b>	922.9 ± 28.1	128.5 ± 3.4	-12.70
	5	<b>PF_3_4_31</b>	908.5 ± 30.7	124.5 ± 4.6	-12.48
	6	<b>PF_3_4_12</b>	811.6 ± 5.6	113.6 ± 2.0	-11.87
	7	<b>PF_3_4_10</b>	717.1 ± 35.8	101.8 ± 2.1	-11.21
<b>PF_12_33</b>	1	<b>PF_12_33_3</b>	882.7 ± 12.2	125.4 ± 5.5	-12.53
	2	<b>PF_12_33_9</b>	839.7 ± 68.3	118.0 ± 9.6	-12.12
	3	<b>PF_12_33_32</b>	802.0 ± 47.3	115.2 ± 3.0	-11.96
	4	<b>PF_12_33_33</b>	816.6 ± 41.5	106.3 ± 5.0	-11.47
	5	<b>PF_12_33_5</b>	706.9 ± 49.0	105.7 ± 1.7	-11.43
<b>PF_12_9</b>	1	<b>PF_12_9_31</b>	940.2 ± 15.4	133.5 ± 2.2	-12.99
	2	<b>PF_12_9_3</b>	906.7 ± 34.0	125.8 ± 4.6	-12.55
	3	<b>PF_12_9_5</b>	885.3 ± 24.0	116.8 ± 5.2	-12.06
	4	<b>PF_12_9_32</b>	726.0 ± 44.7	107.3 ± 4.1	-11.52
	5	<b>PF_12_9_33</b>	760.3 ± 8.8	101.2 ± 2.5	-11.18
	6	<b>PF_12_9_12</b>	683.1 ± 8.6	97.5 ± 1.4	-10.97
	7	<b>PF_12_9_2</b>	721.8 ± 25.9	90.5 ± 4.0	-10.58

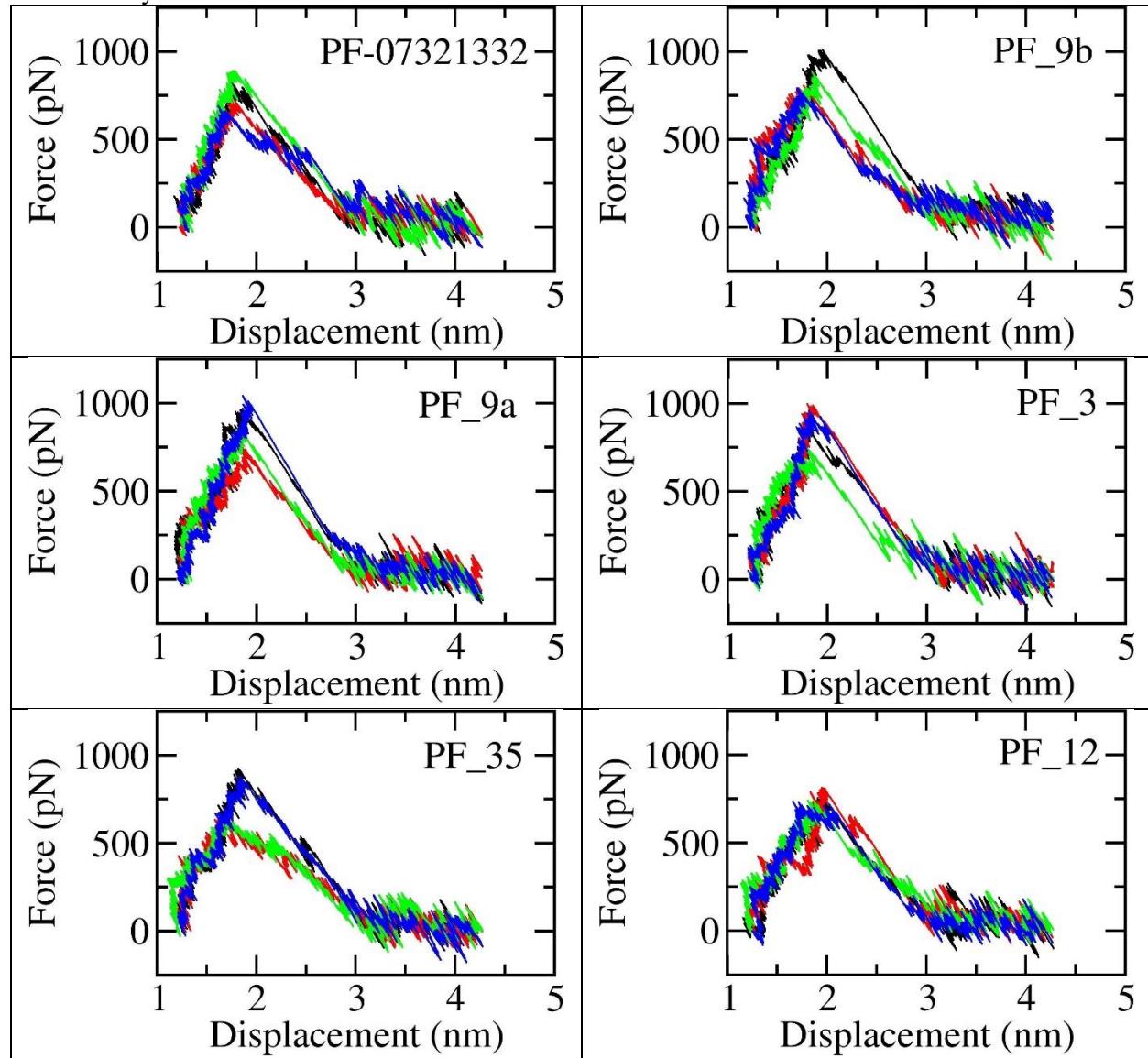
**Table S4.** FPL results of the modified compounds.

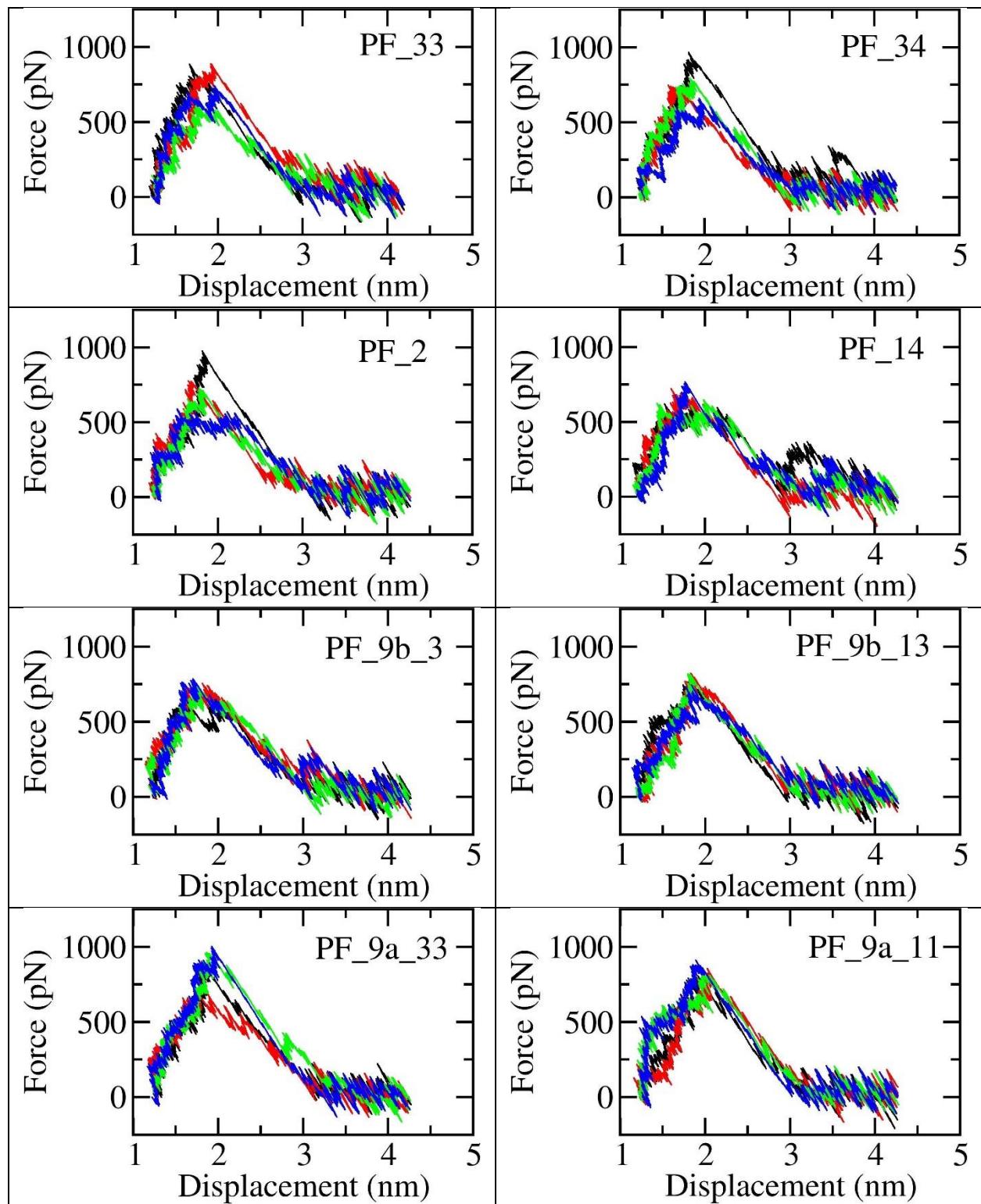
<b>Source</b>	<b>N<sup>0</sup></b>	<b>Code</b>	<b>F<sub>Max</sub></b>	<b>W</b>	<b>ΔG<sub>FPL</sub><sup>Pre a</sup></b>
<b>PF_3_4_62</b>	1	<b>PF_3_4_62_33</b>	927.7 ± 28.0	122.6 ± 4.2	-12.38
	2	<b>PF_3_4_62_12</b>	785.7 ± 32.5	101.8 ± 6.3	-11.21
	3	<b>PF_3_4_62_32</b>	731.7 ± 60.3	100.4 ± 5.7	-11.13
	4	<b>PF_3_4_62_10</b>	661.4 ± 31.4	86.7 ± 4.1	-10.37
<b>PF_3_4_32</b>	1	<b>PF_3_4_32_32</b>	911.3 ± 24.8	126.3 ± 3.8	-12.58
	2	<b>PF_3_4_32_10</b>	837.0 ± 60.4	121.1 ± 7.1	-12.30
<b>PF_3_4_7</b>	1	<b>PF_3_4_7_30</b>	792.3 ± 66.8	119.4 ± 8.1	-12.20
	2	<b>PF_3_4_7_9</b>	792.7 ± 26.7	103.5 ± 6.9	-11.31
	3	<b>PF_3_4_7_31</b>	640.4 ± 38.9	89.9 ± 2.5	-10.55
	4	<b>PF_3_4_7_32</b>	660.3 ± 28.9	89.6 ± 5.8	-10.53
	5	<b>PF_3_4_7_11</b>	611.1 ± 43.3	76.1 ± 4.3	-9.77
<b>PF_3_4_33</b>	1	<b>PF_3_4_33_32</b>	927.2 ± 39.3	142.2 ± 6.2	-13.47
	2	<b>PF_3_4_33_31</b>	908.0 ± 63.5	134.7 ± 8.1	-13.05
	3	<b>PF_3_4_33_12</b>	676.8 ± 22.7	98.3 ± 4.1	-11.01
<b>PF_3_4_31</b>	1	<b>PF_3_4_31_32</b>	863.1 ± 45.1	123.3 ± 6.1	-12.42
	2	<b>PF_3_4_31_10</b>	795.4 ± 49.9	109.5 ± 4.1	-11.65
	3	<b>PF_3_4_31_12</b>	729.7 ± 51.4	105.4 ± 5.4	-11.41
<b>PF_12_33_3</b>	1	<b>PF_12_33_3_32</b>	804.3 ± 28.3	111.8 ± 4.0	-11.77
	2	<b>PF_12_33_3_31</b>	727.1 ± 53.1	100.2 ± 5.0	-11.12
	3	<b>PF_12_33_3_8</b>	691.5 ± 32.2	94.9 ± 4.9	-10.83
	4	<b>PF_12_33_3_2</b>	729.6 ± 21.6	89.9 ± 4.9	-10.55
	5	<b>PF_12_33_3_4</b>	656.4 ± 36.4	87.6 ± 4.6	-10.42
<b>PF_12_9_31</b>	1	<b>PF_12_9_31_32</b>	836.6 ± 28.1	126.8 ± 3.9	-12.61
	2	<b>PF_12_9_31_3</b>	829.9 ± 43.2	112.7 ± 5.9	-11.82
	3	<b>PF_12_9_31_2</b>	705.3 ± 29.1	89.6 ± 3.6	-10.53
<b>PF_12_9_3</b>	1	<b>PF_12_9_3_31</b>	690.2 ± 44.7	97.7 ± 4.0	-10.98
	2	<b>PF_12_9_3_30</b>	587.0 ± 43.2	87.6 ± 3.9	-10.42
	3	<b>PF_12_9_3_32</b>	569.0 ± 35.0	73.7 ± 4.1	-9.64
	4	<b>PF_12_9_3_11</b>	532.9 ± 40.3	72.9 ± 3.7	-9.59

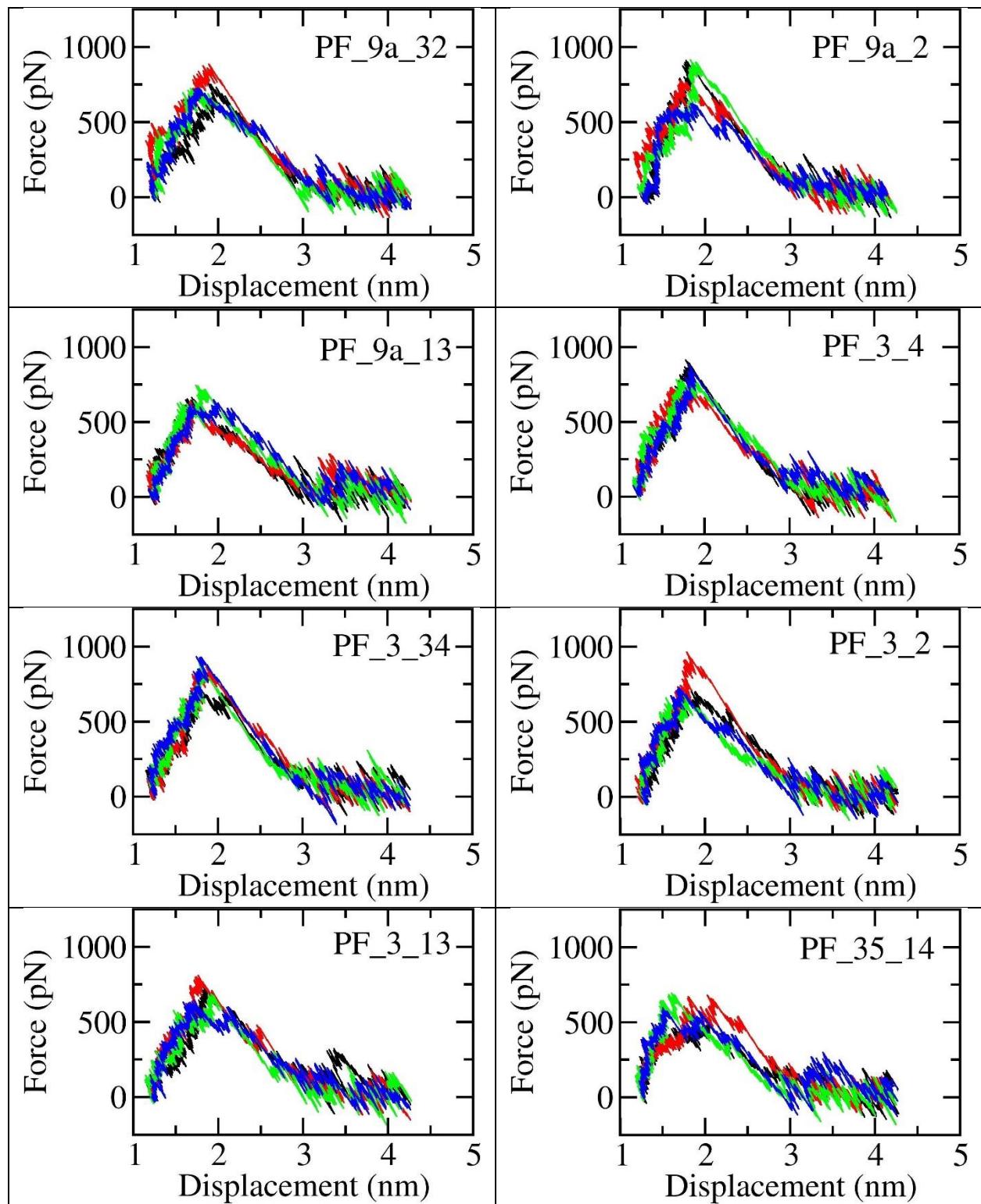
**Table S5.** FPL results of the modified compounds.

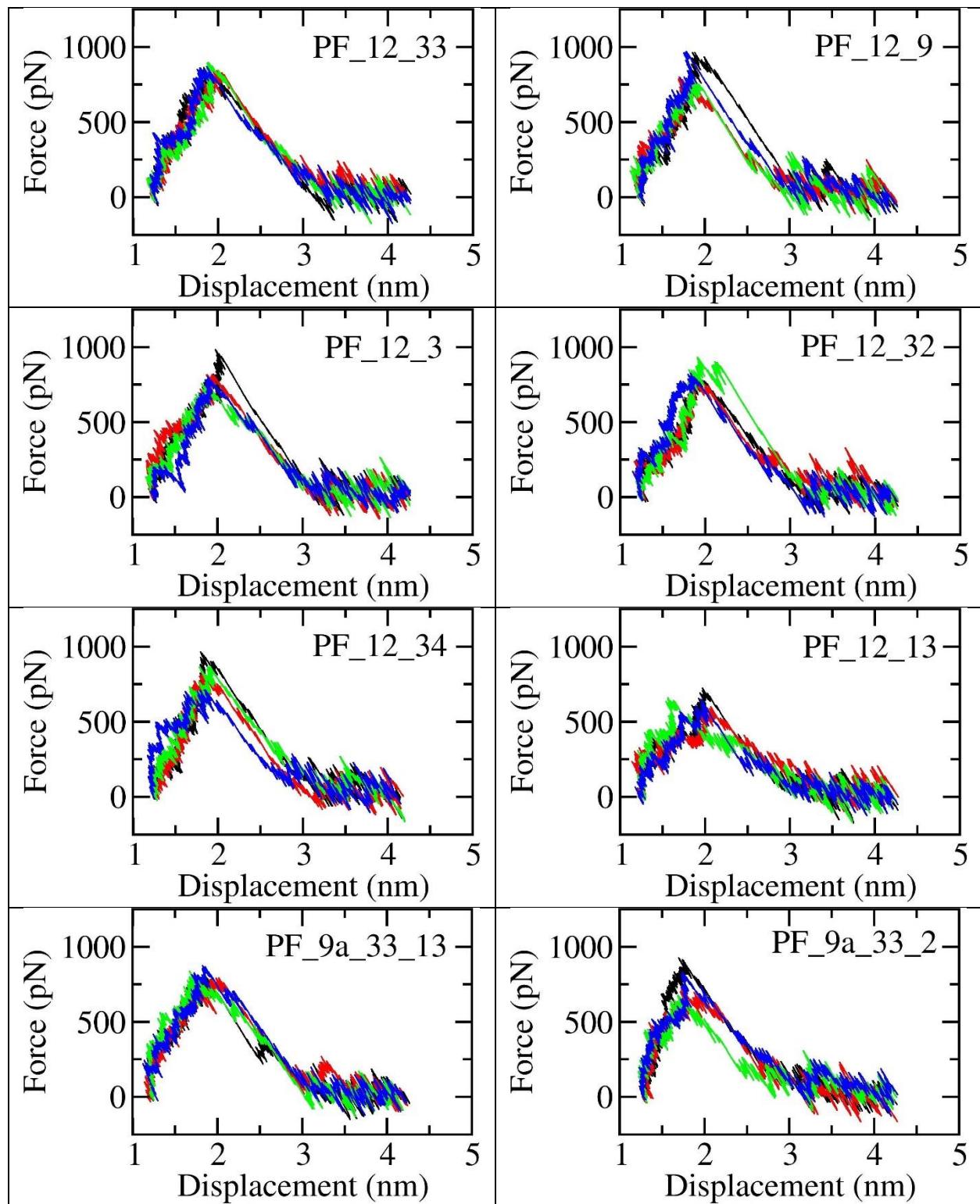
Source	N <sup>0</sup>	Code	F <sub>Max</sub>	W	ΔG <sub>FPL</sub> <sup>Pre a</sup>
<b>PF_3_4_33_32</b>	1	<b>PF_3_4_33_32_10</b>	658.0 ± 37.5	95.3 ± 5.5	-10.85
	2	<b>PF_3_4_33_32_60</b>	737.6 ± 19.2	95.2 ± 2.52	-10.84
<b>PF_3_4_31</b>	1	<b>PF_3_4_33_31_60</b>	813.8 ± 12.4	105.0 ± 1.4	-11.39
	2	<b>PF_3_4_33_31_10</b>	646.1 ± 31.5	95.9 ± 3.4	-10.88

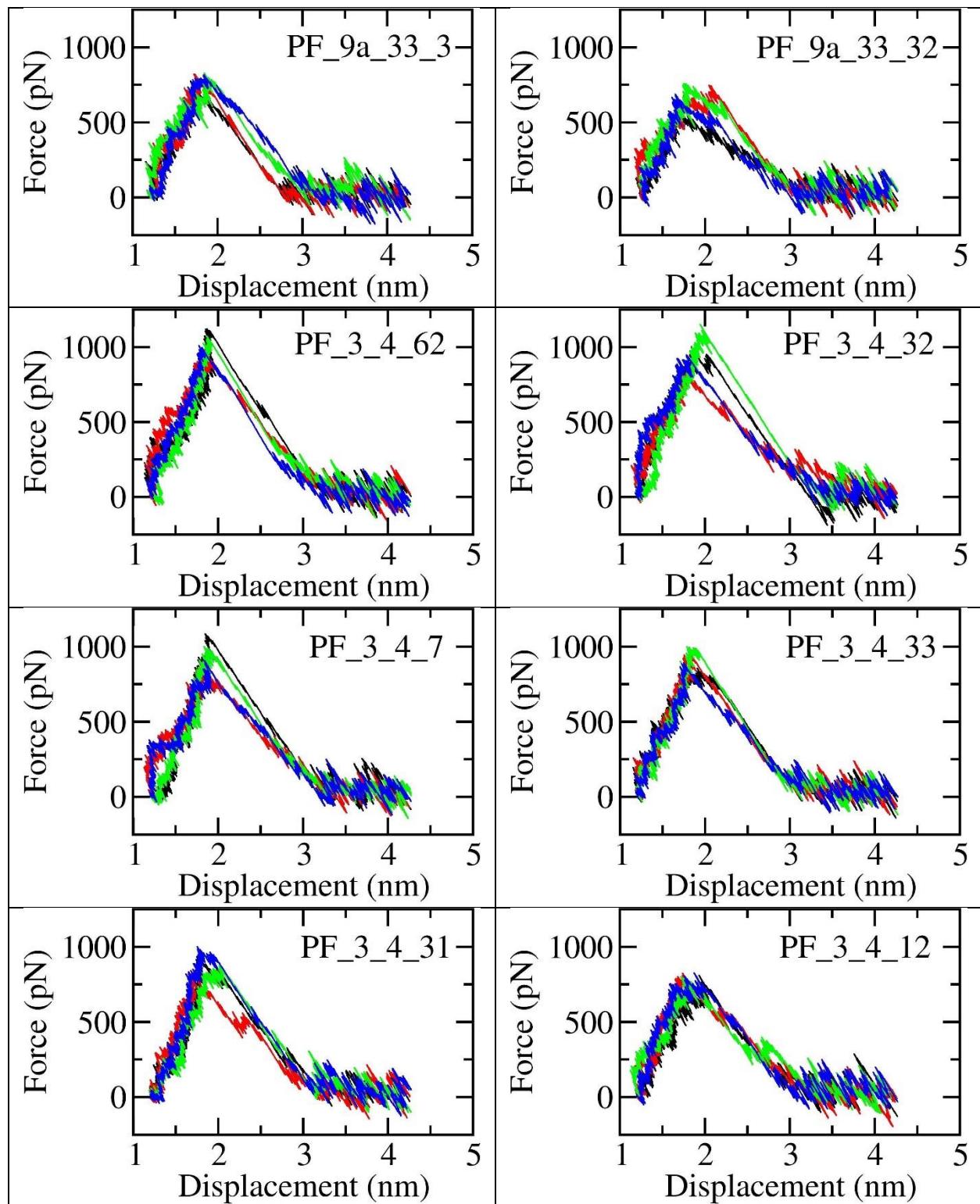
**Table S6.** The profile of pulling work over the ligand displacement. The calculations were repeated 4 times each system.

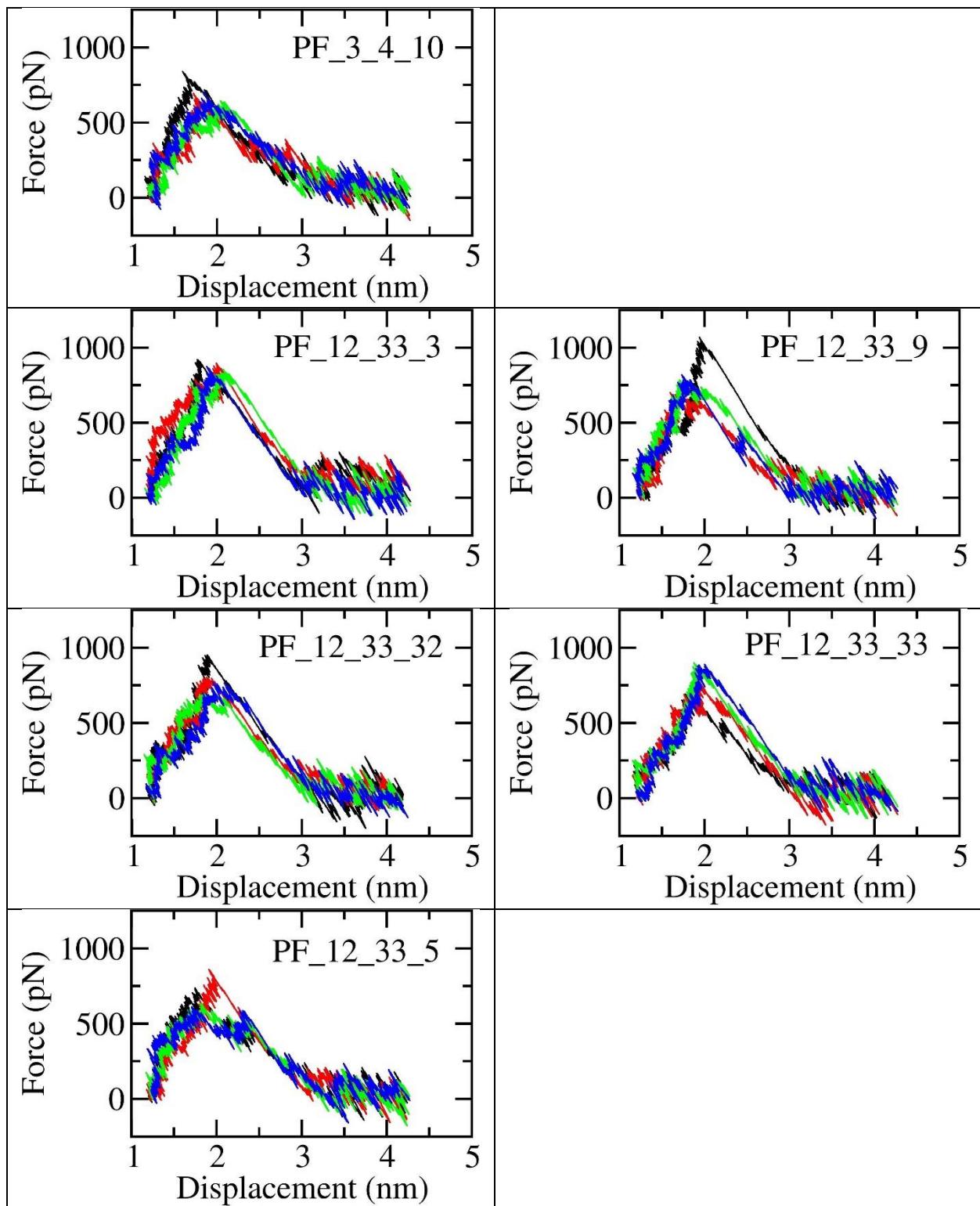


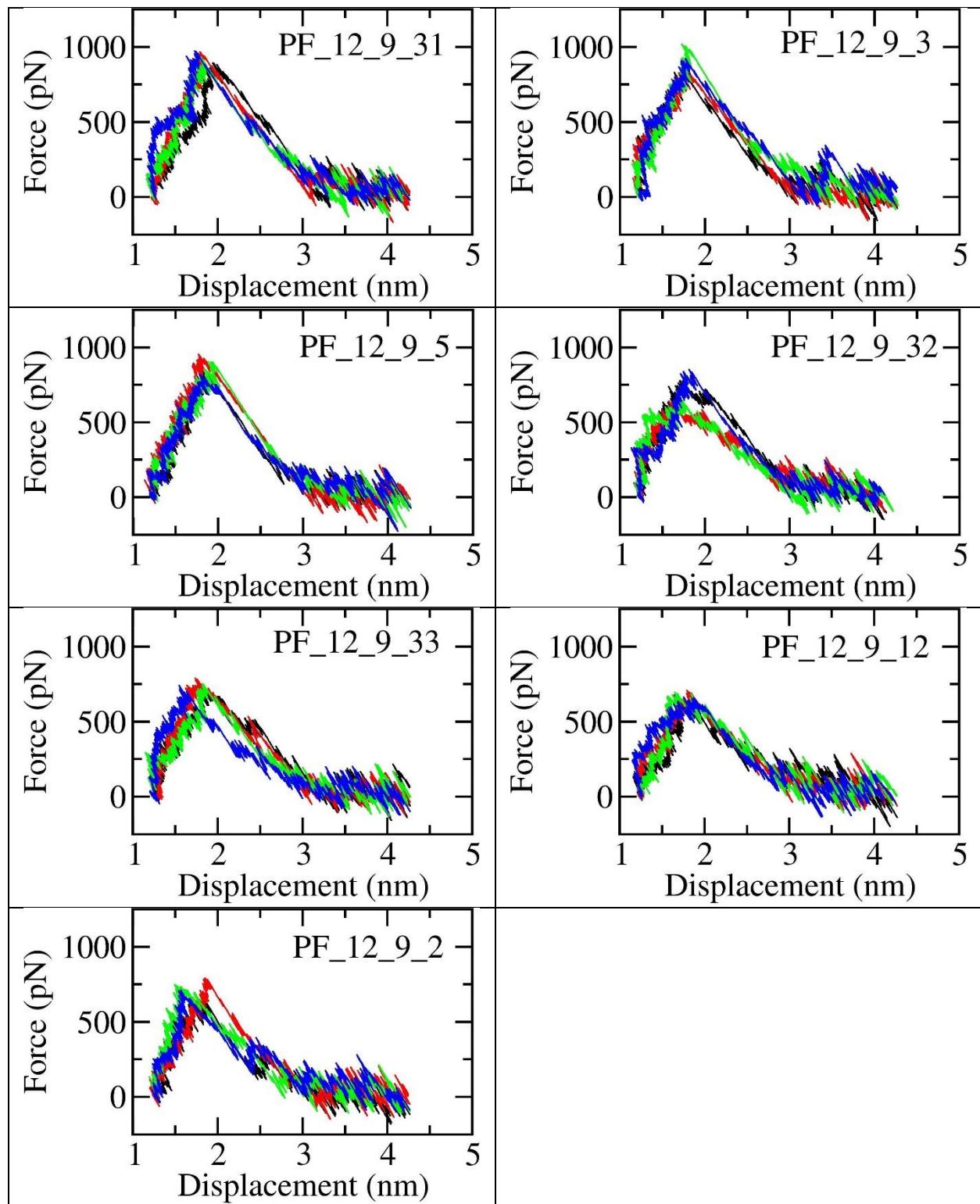


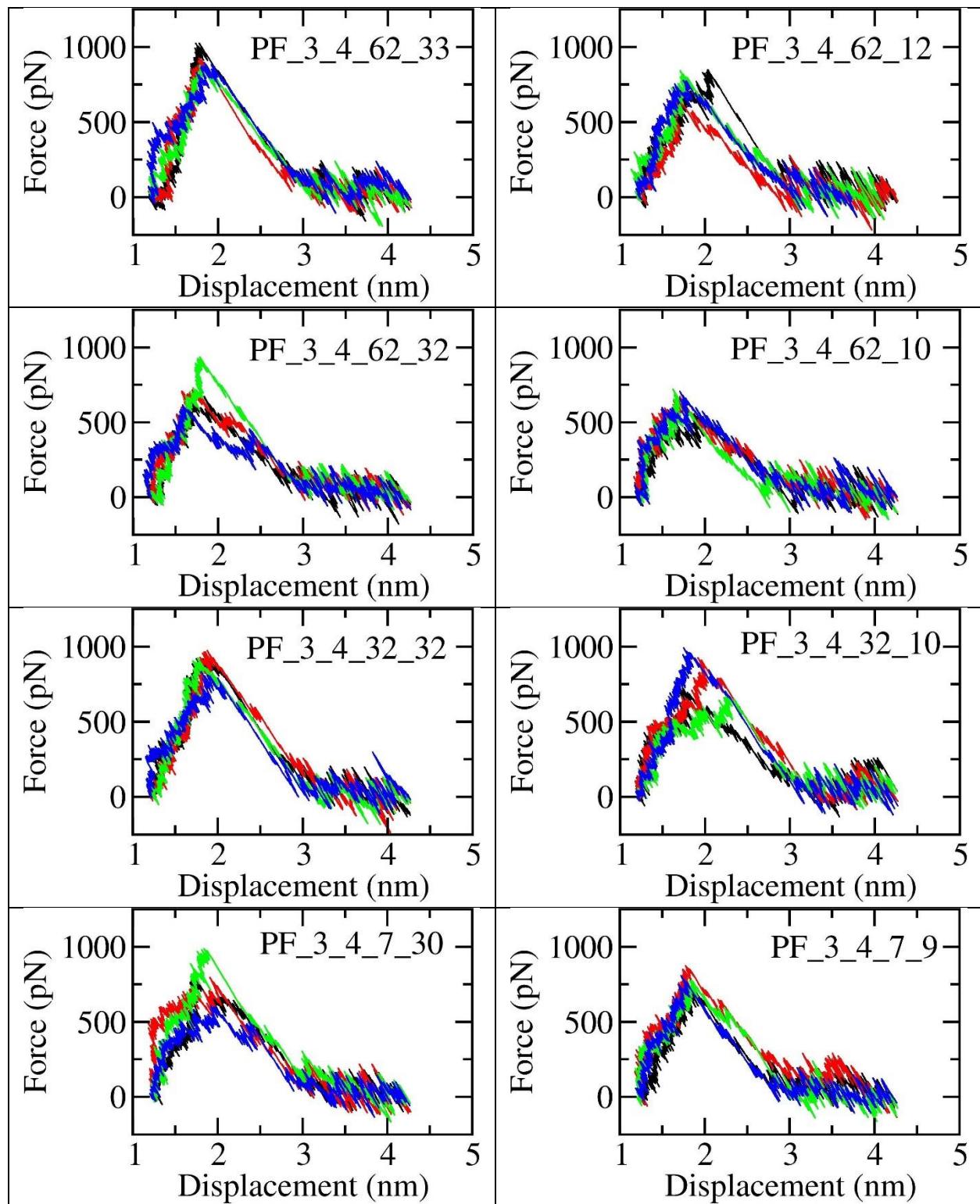


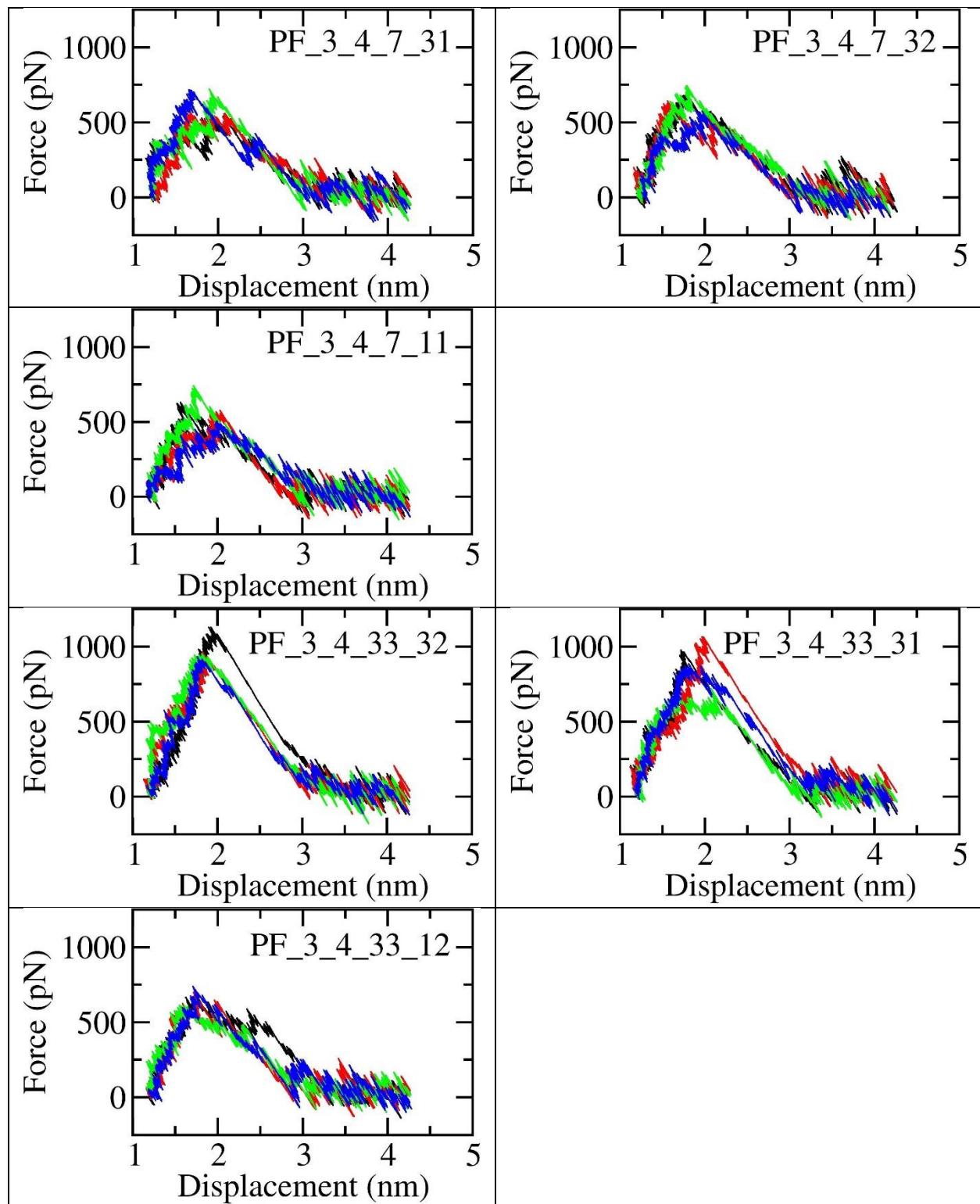


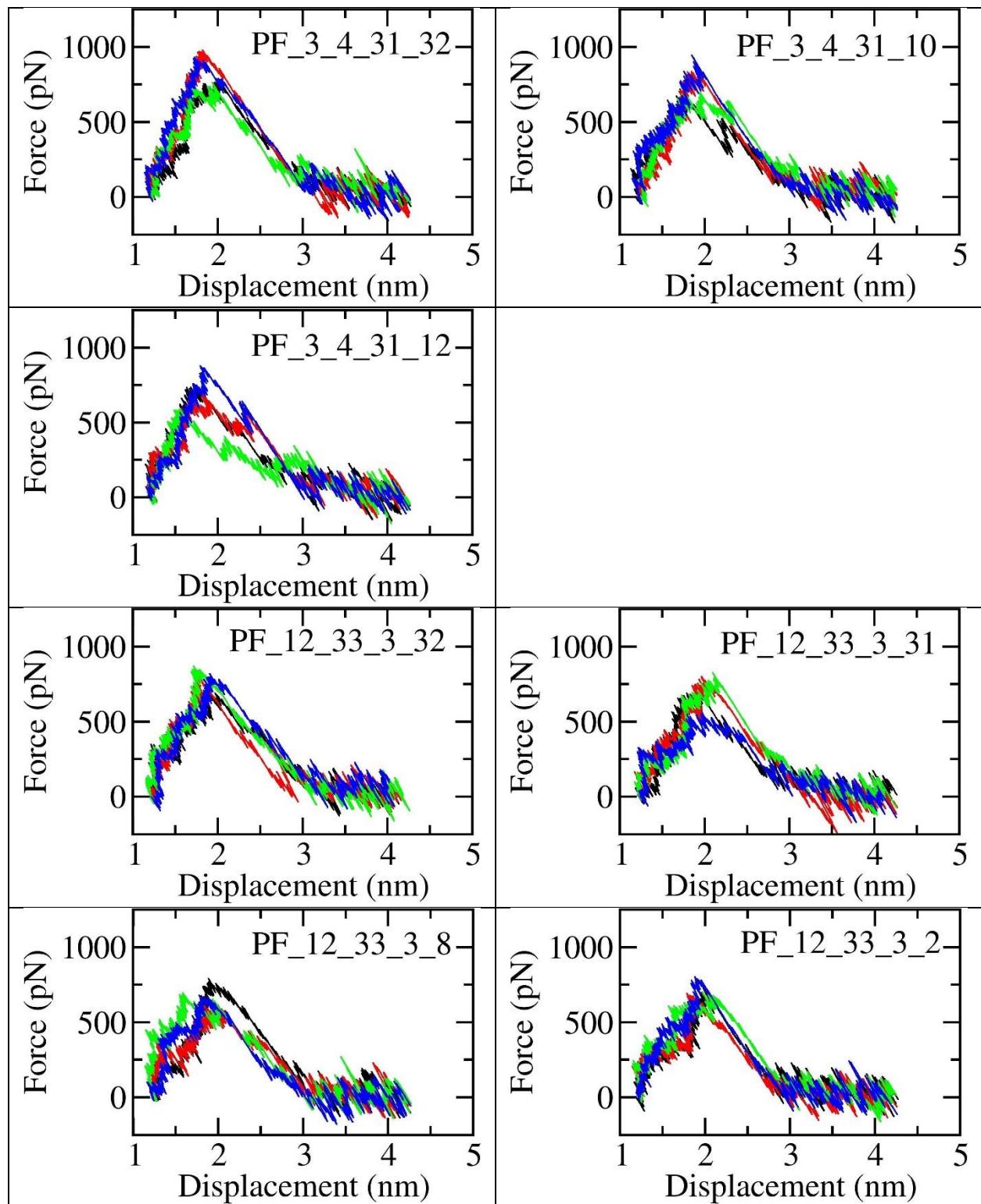


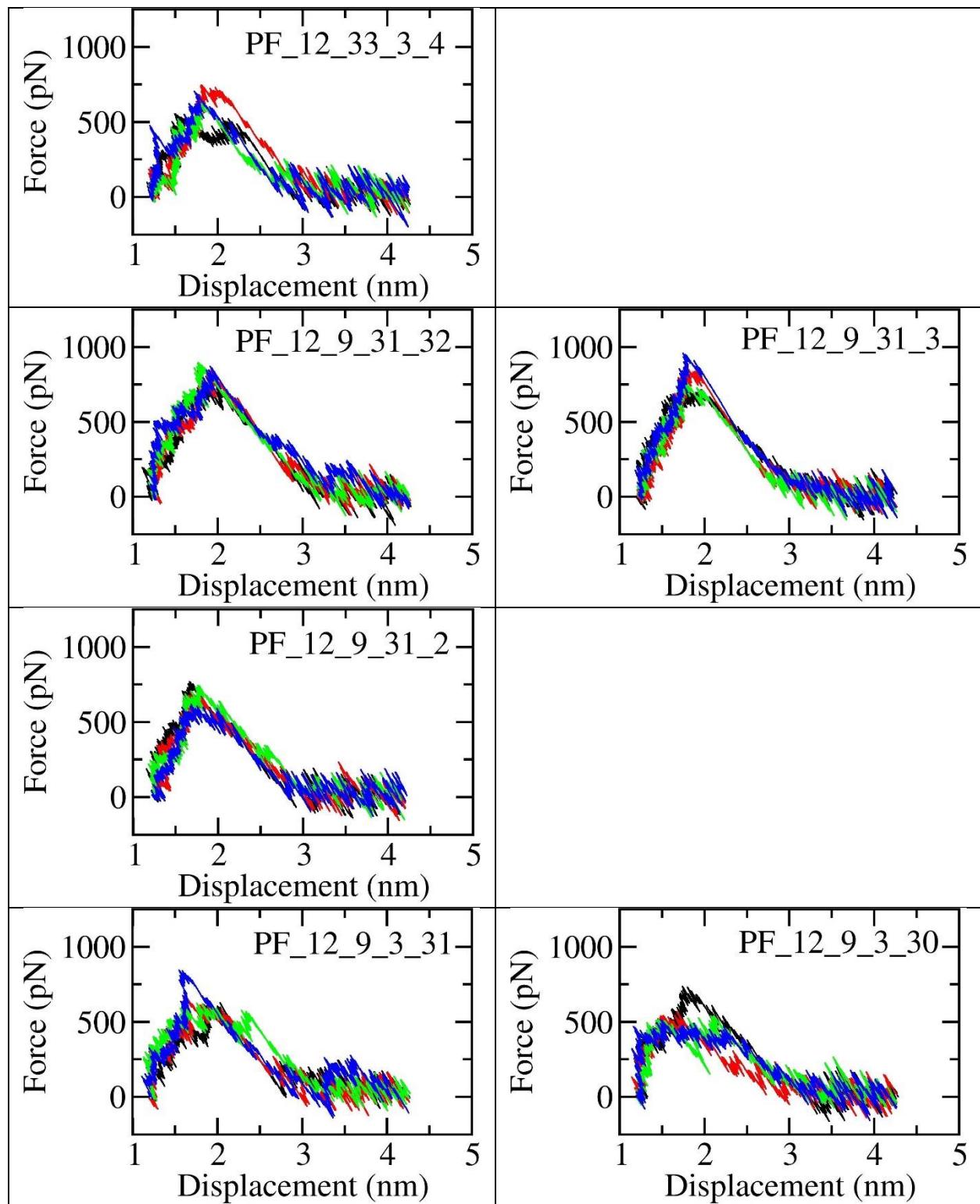


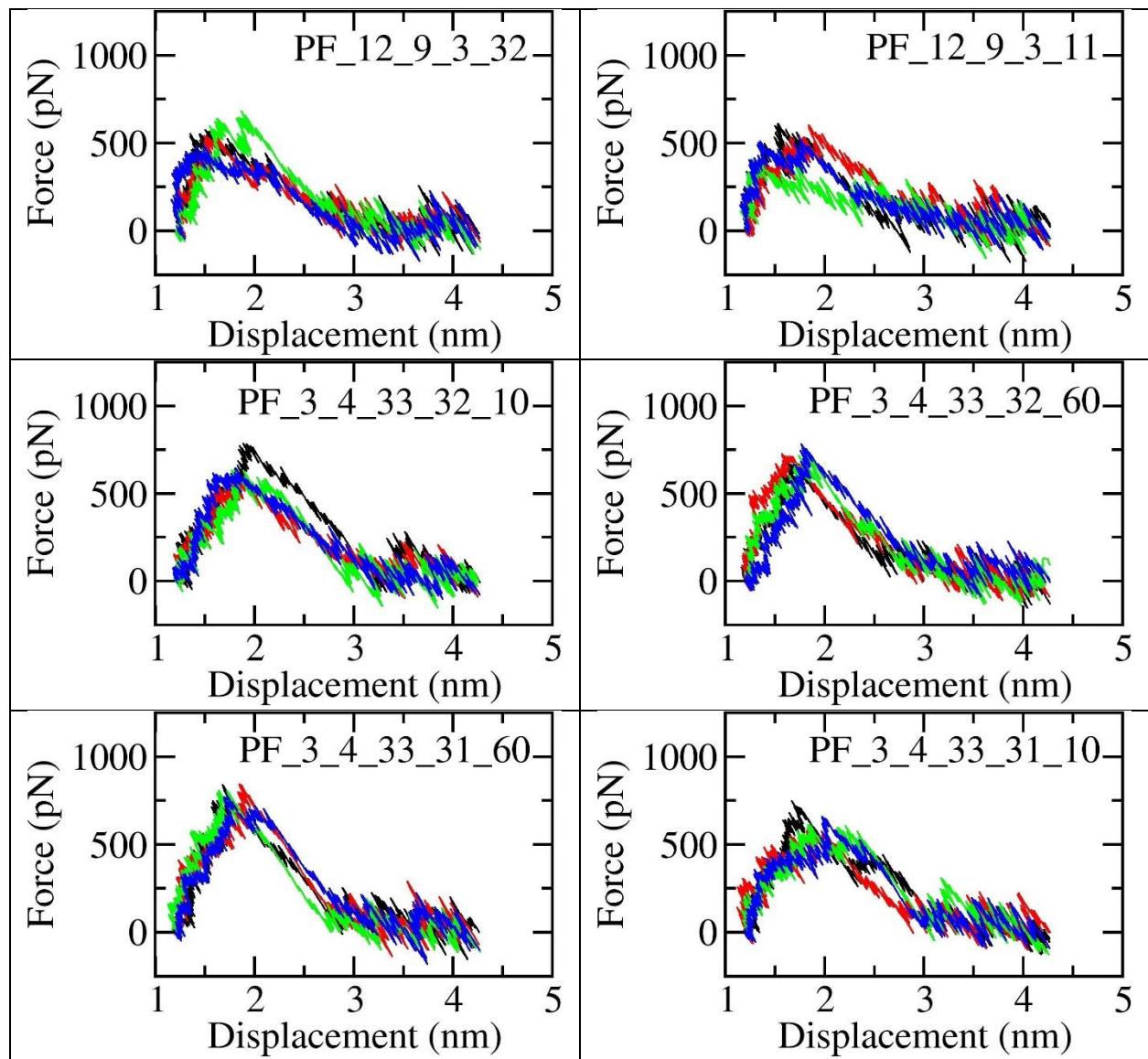




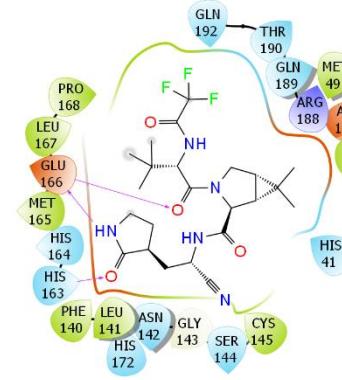
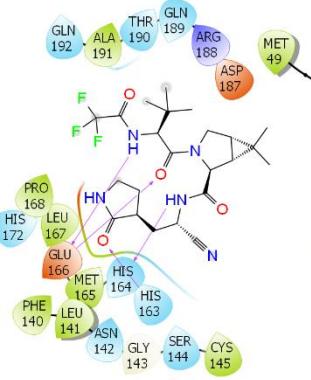
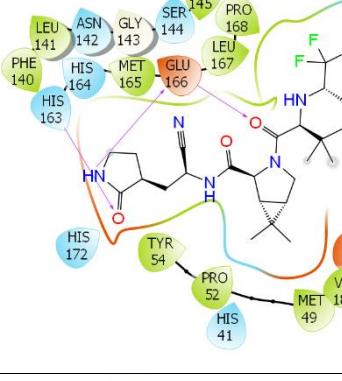
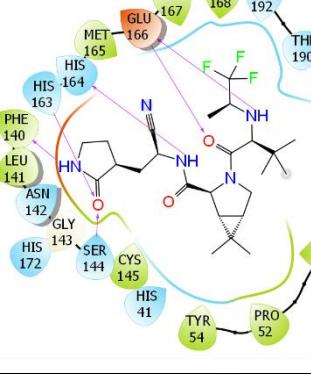
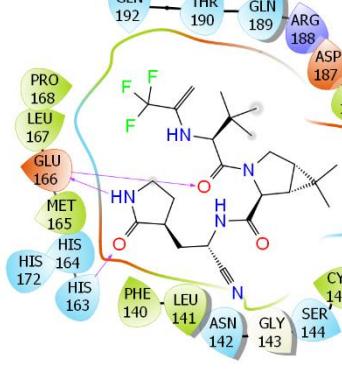
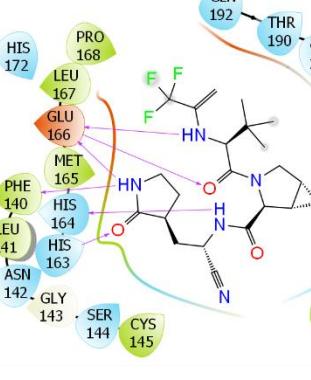


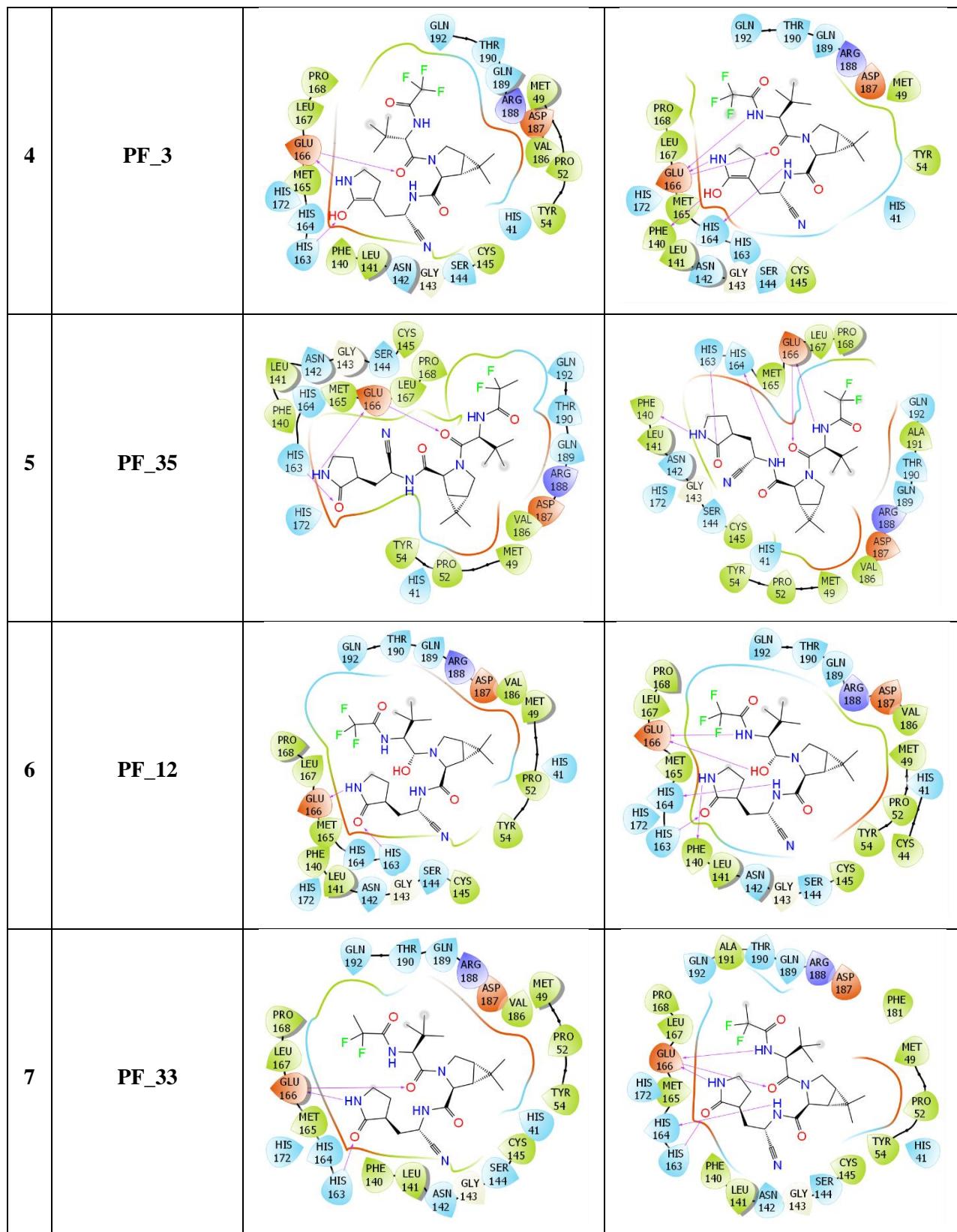


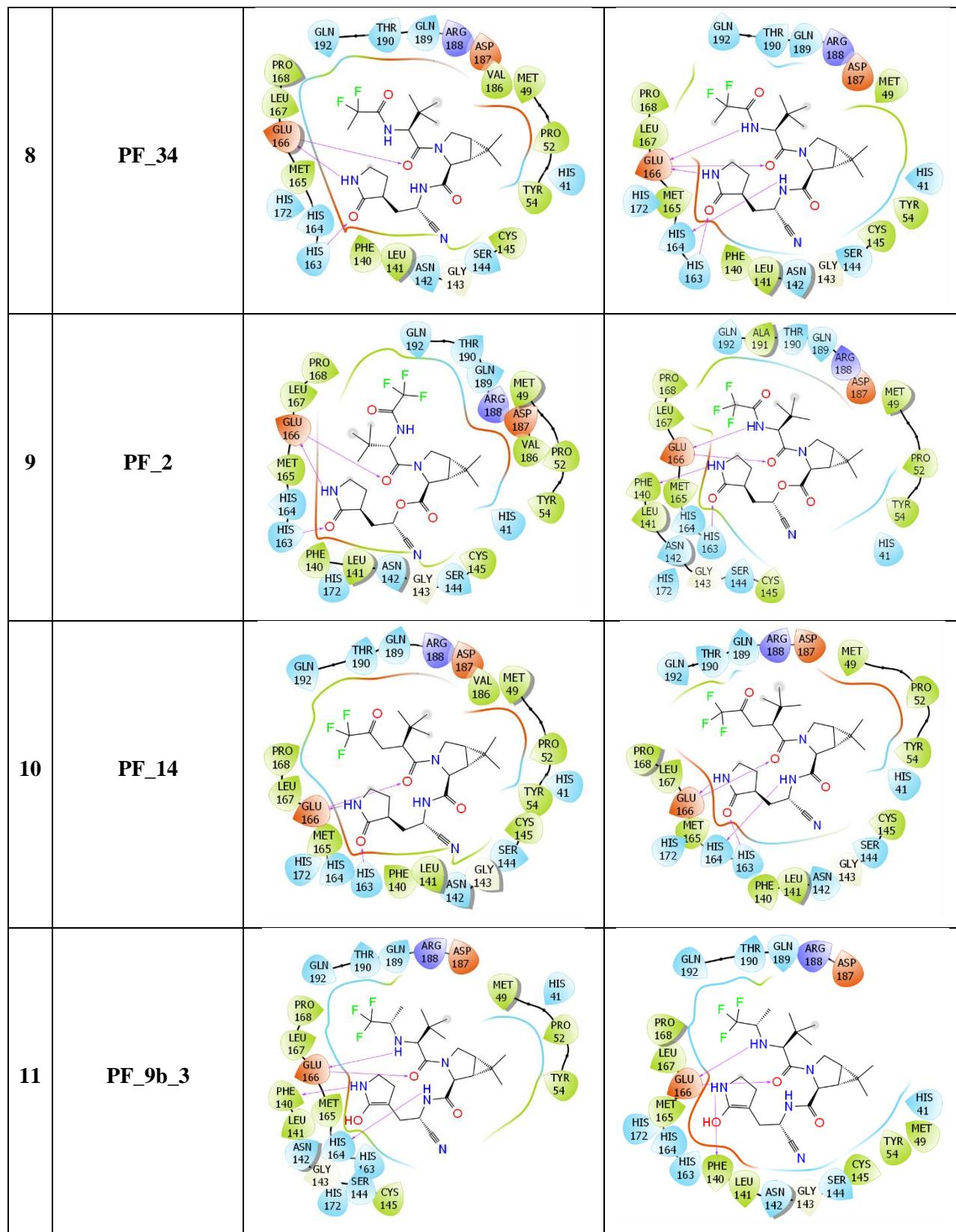


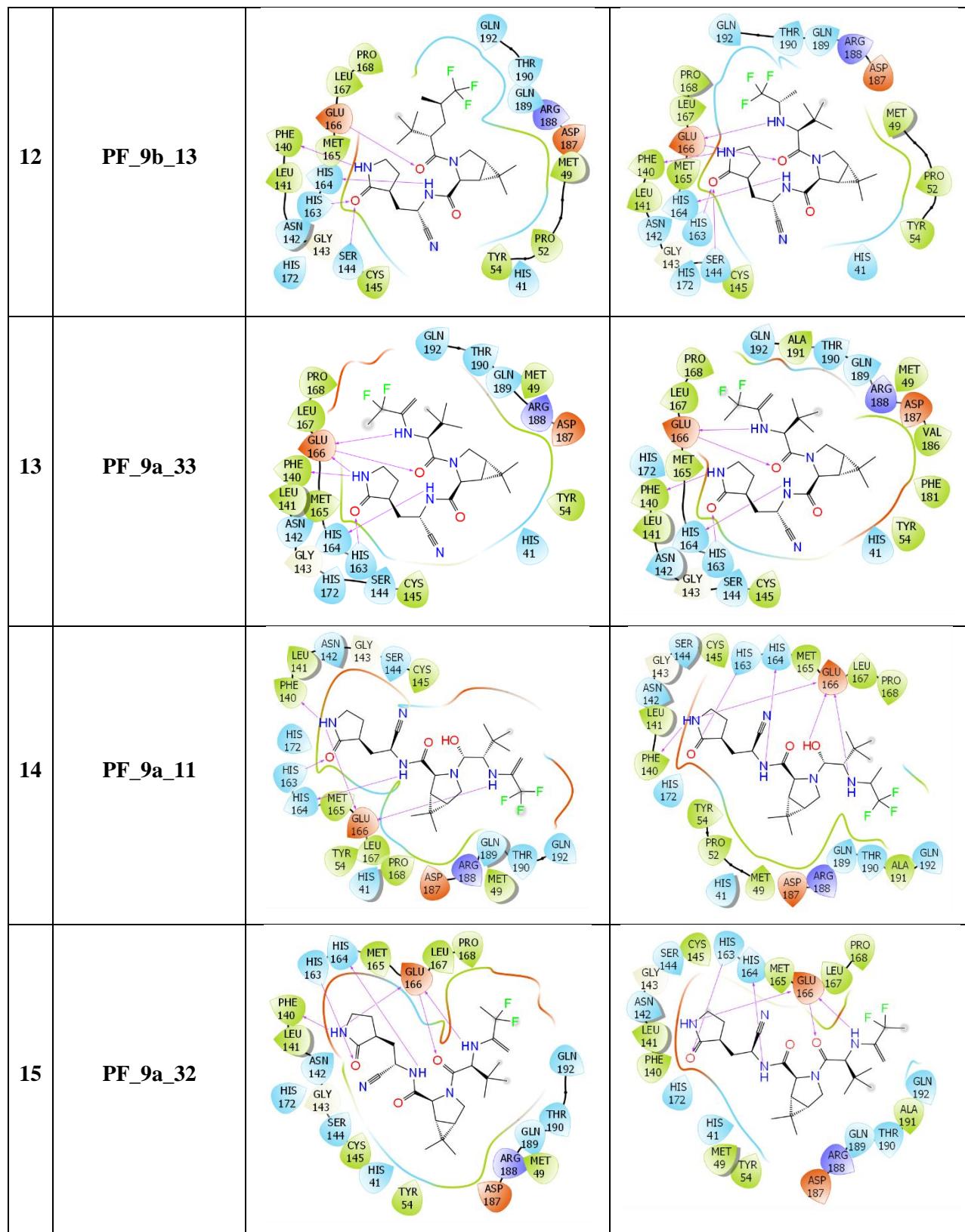


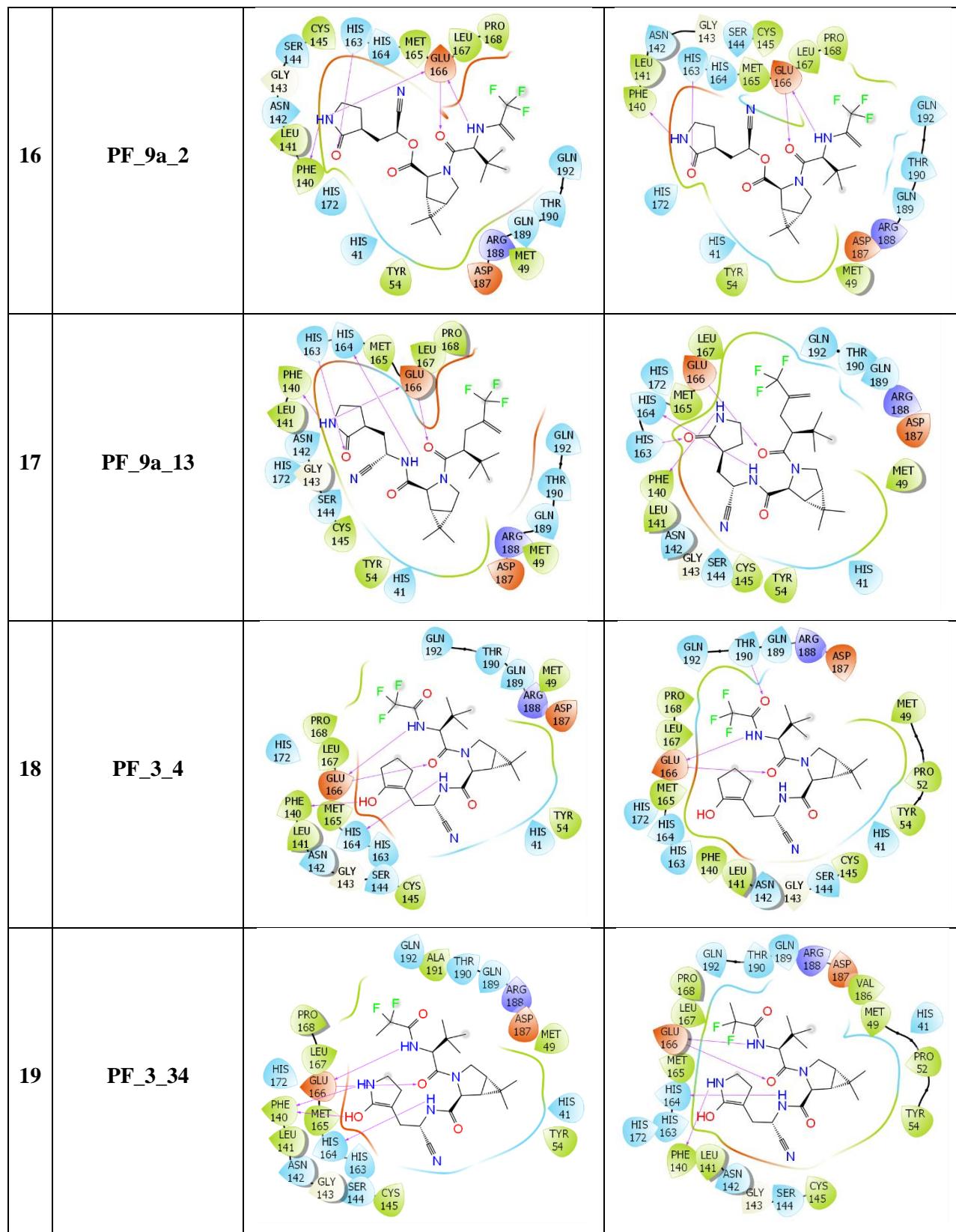
**Table S7.** The 2D interaction diagram between SARS-CoV-2 Mpro + ligand.

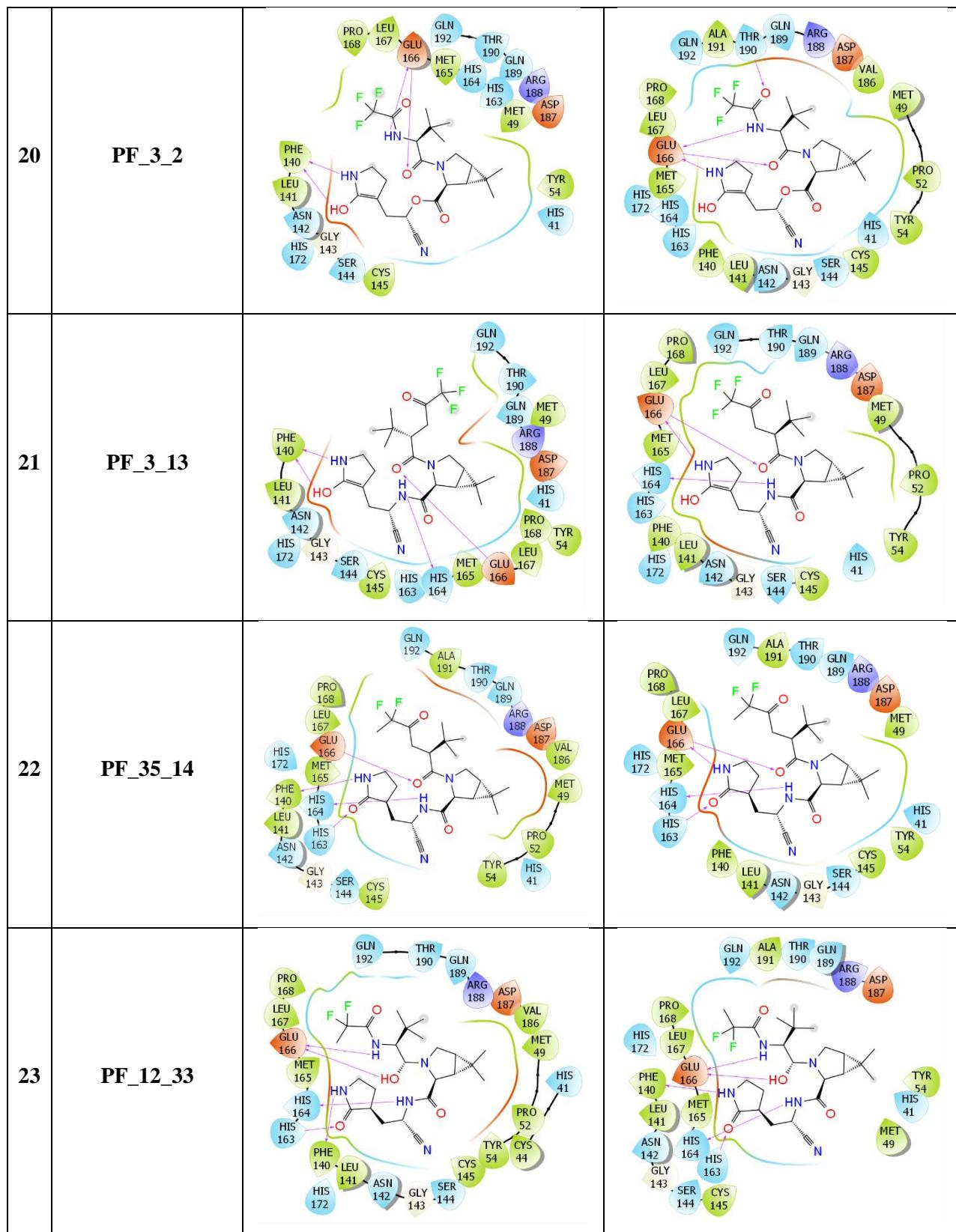
Nº	Name	Initial Structure <sup>a</sup>	MD-refined structure <sup>b</sup>
1	PF-07321332		
2	PF_9b		
3	PF_9a		





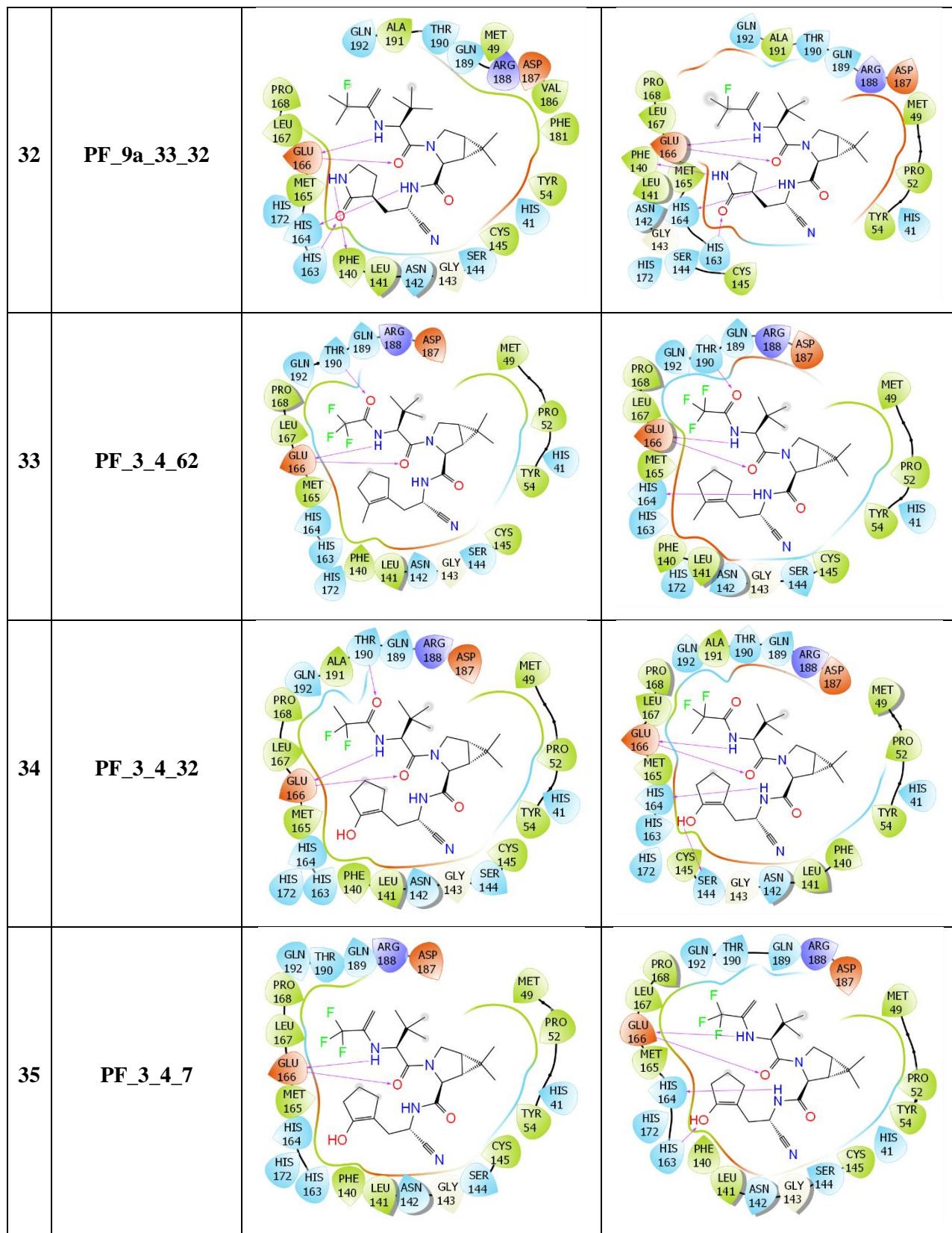


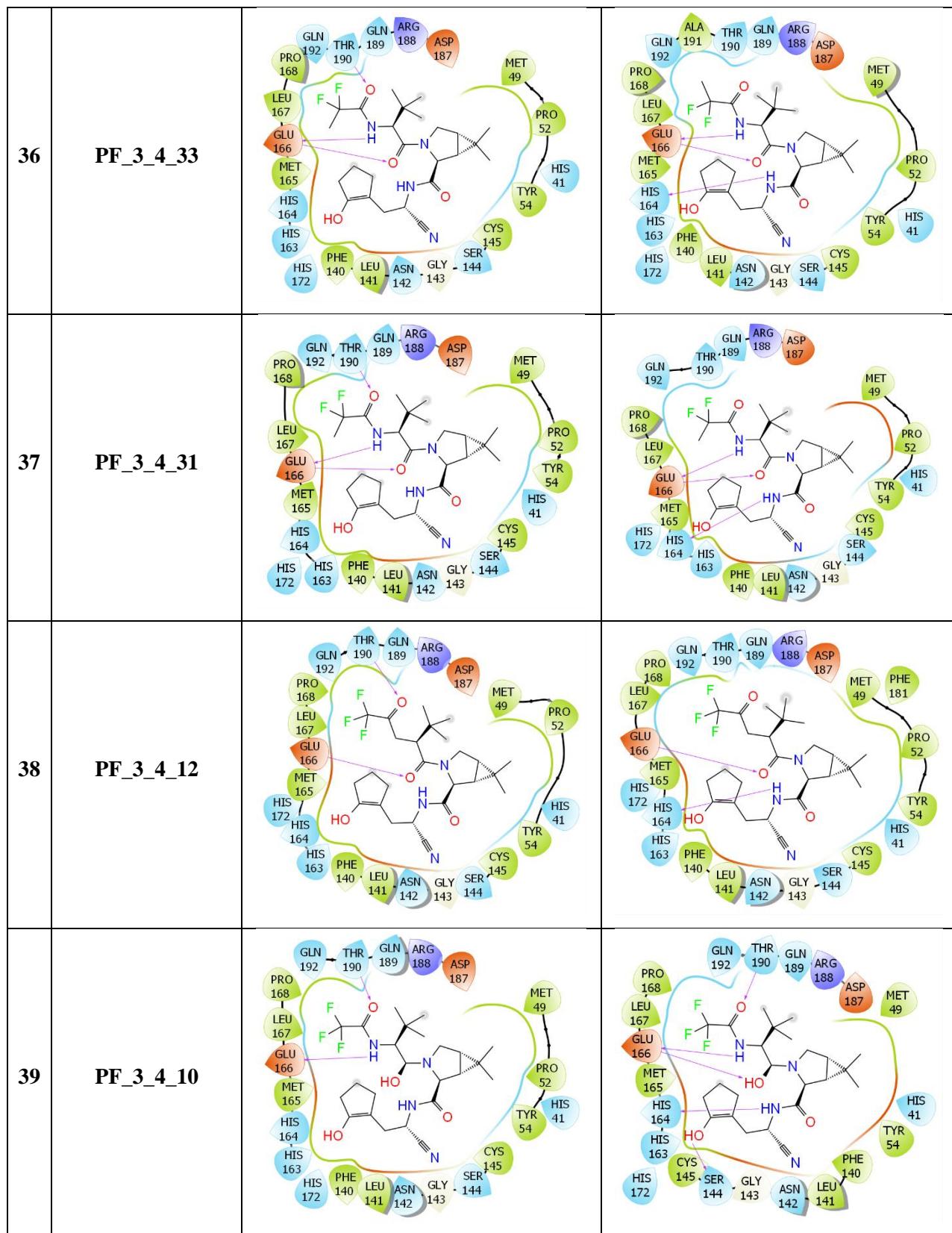


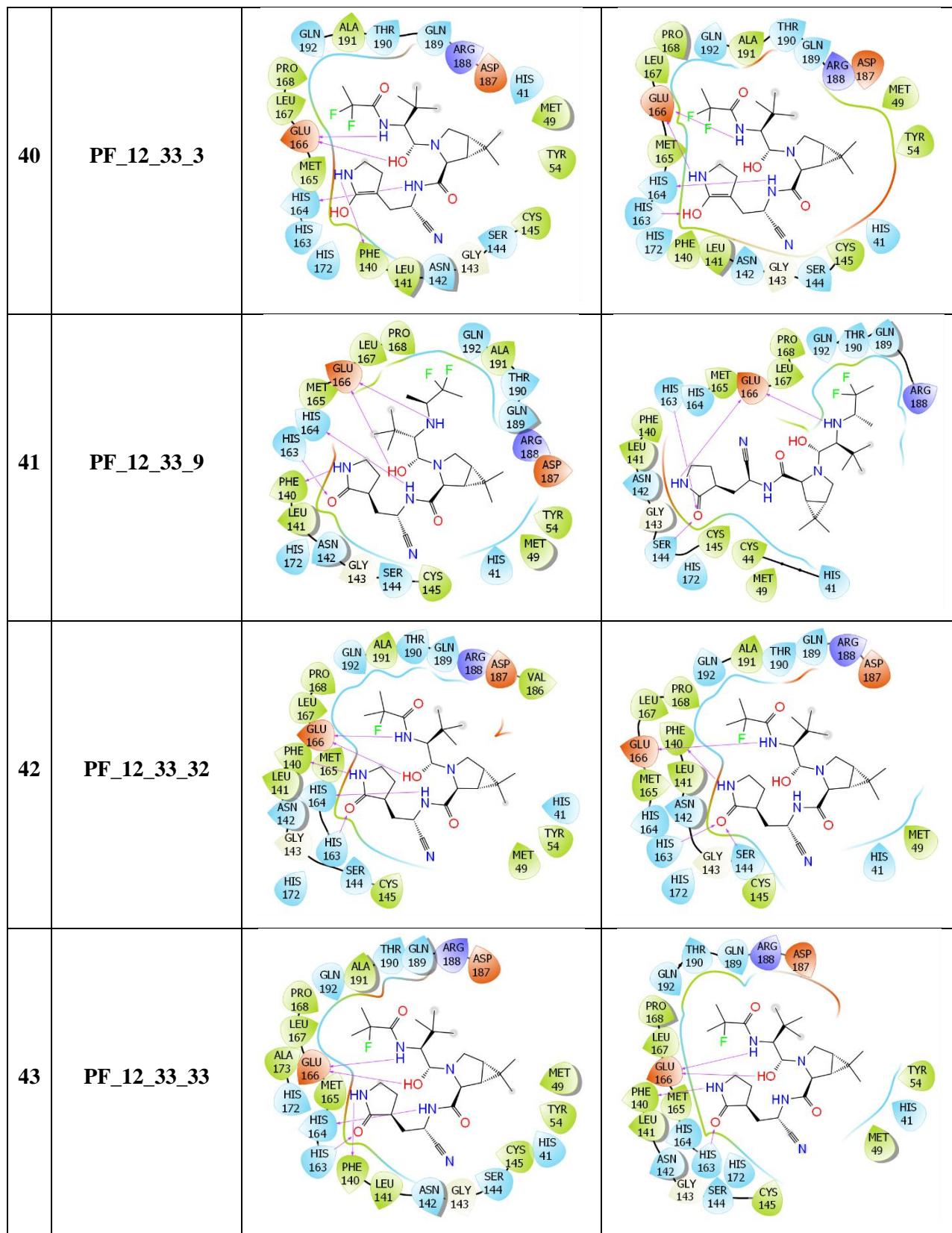


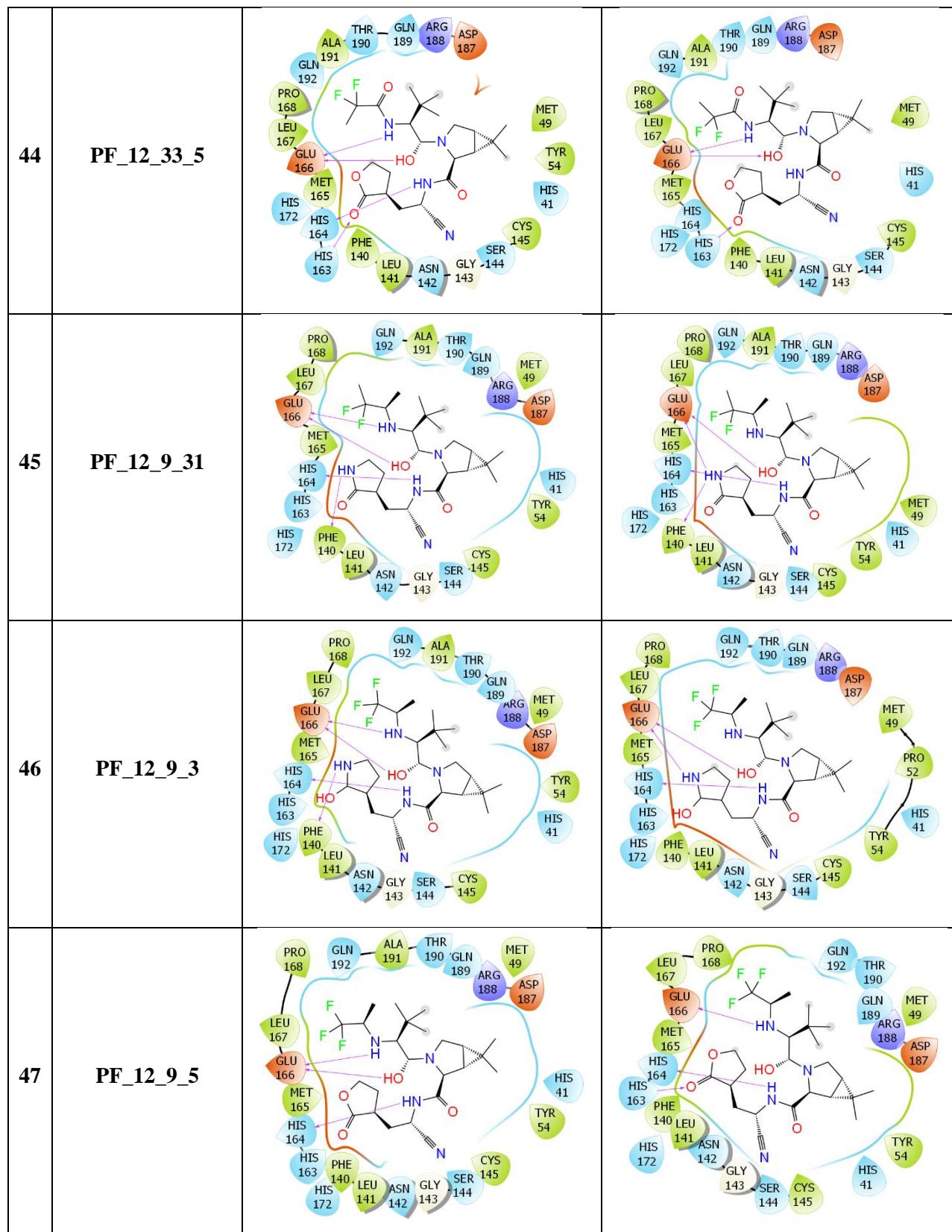
24	PF_12_9		
25	PF_12_3		
26	PF_12_32		
27	PF_12_34		

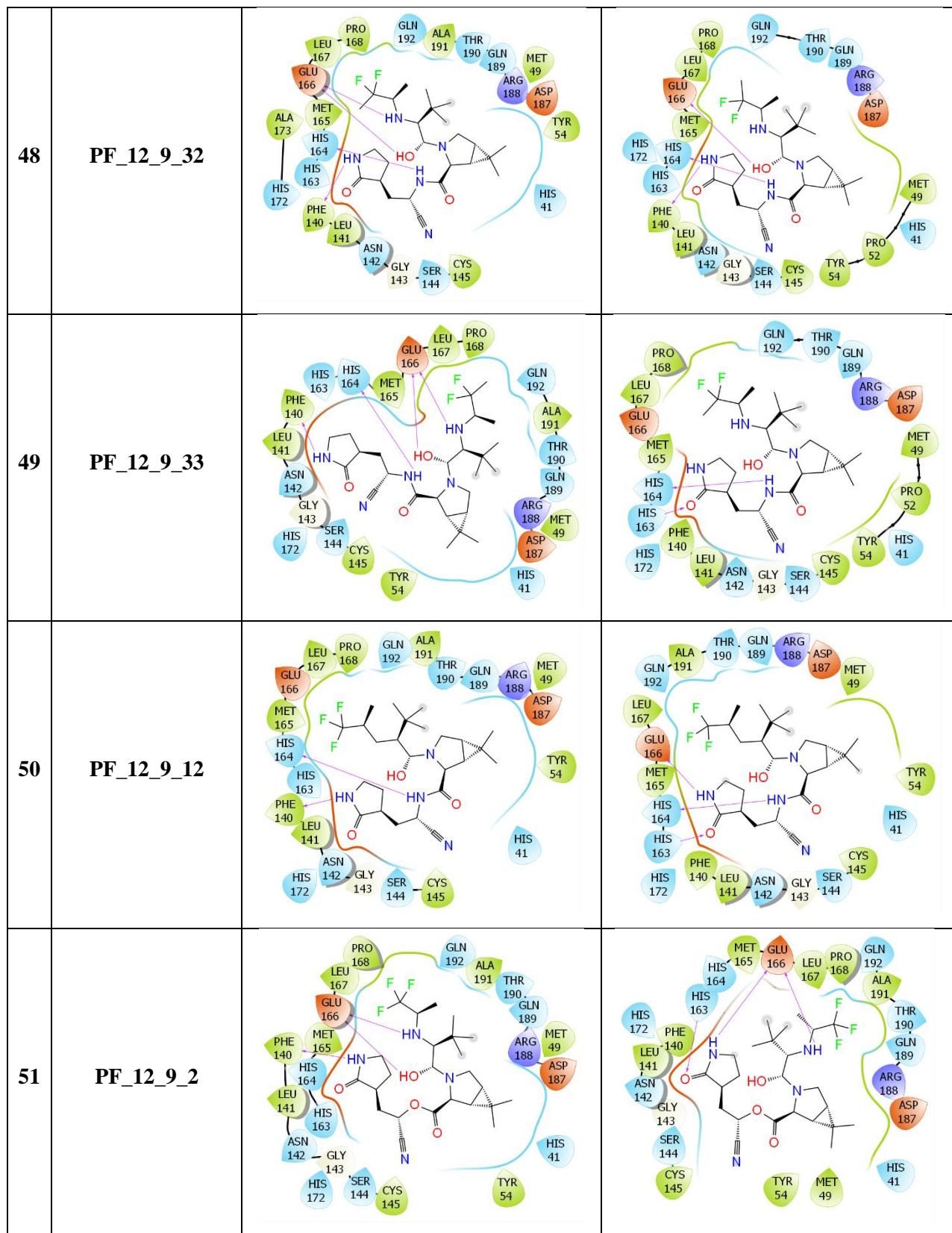
28	PF_12_13		
29	PF_9a_33_13		
30	PF_9a_33_2		
31	PF_9a_33_3		

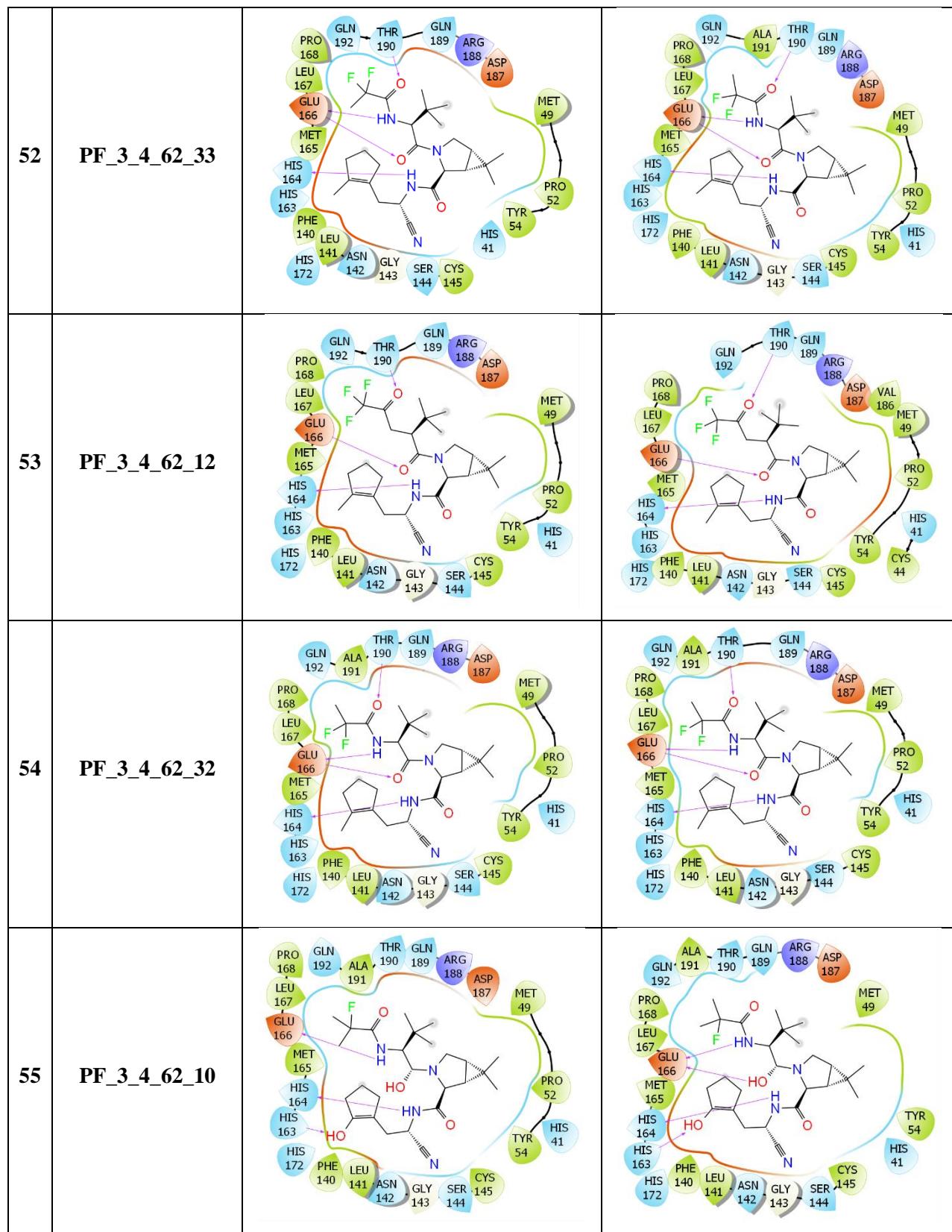


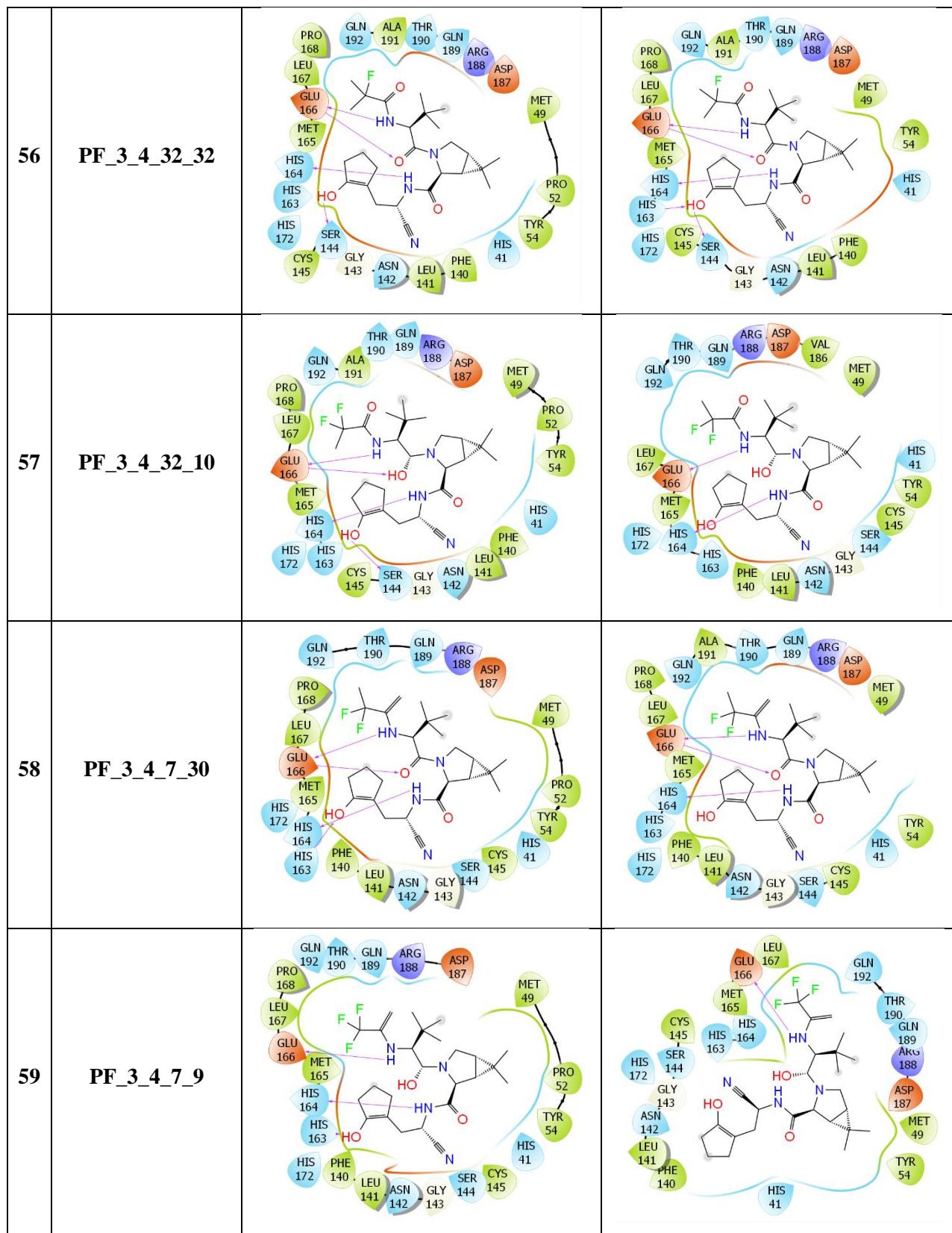


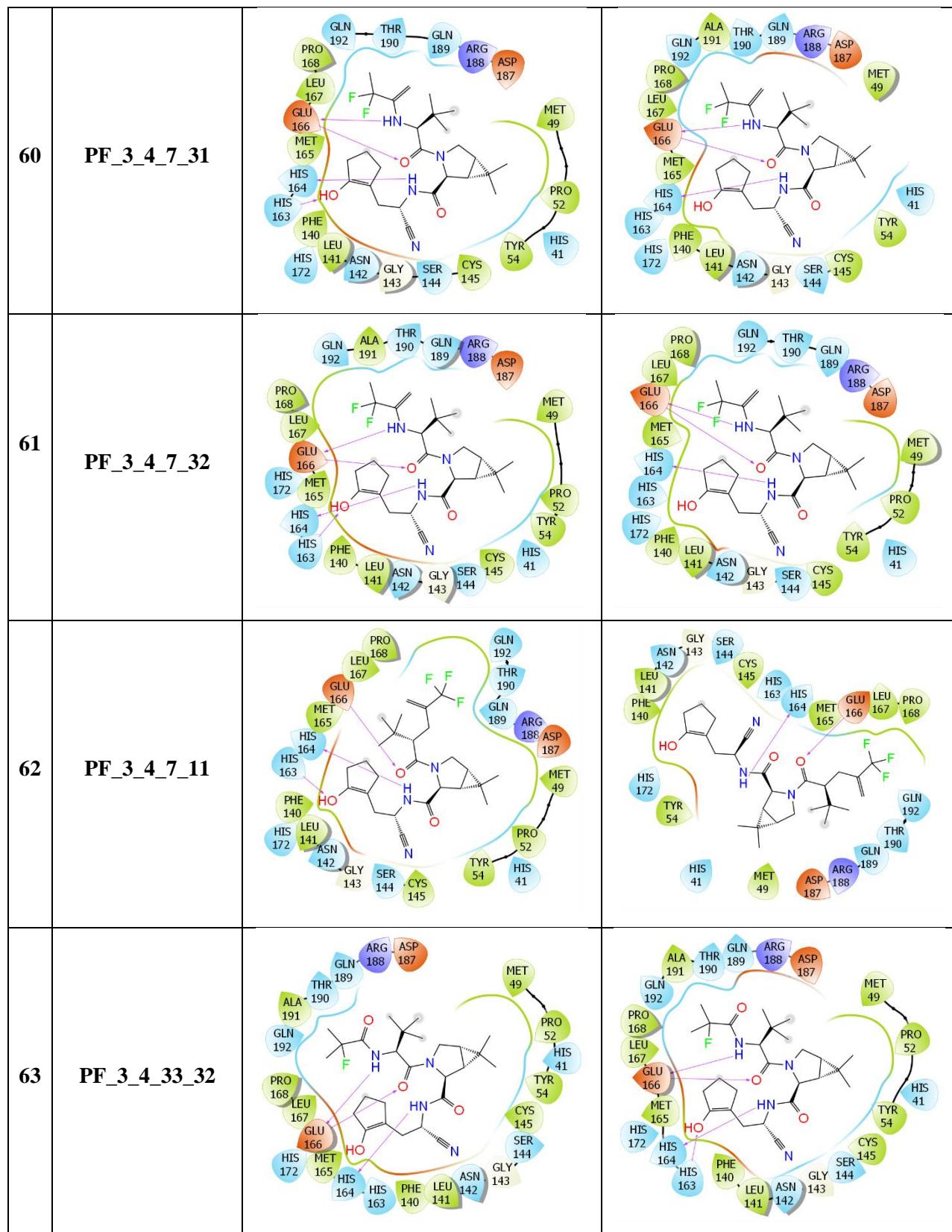




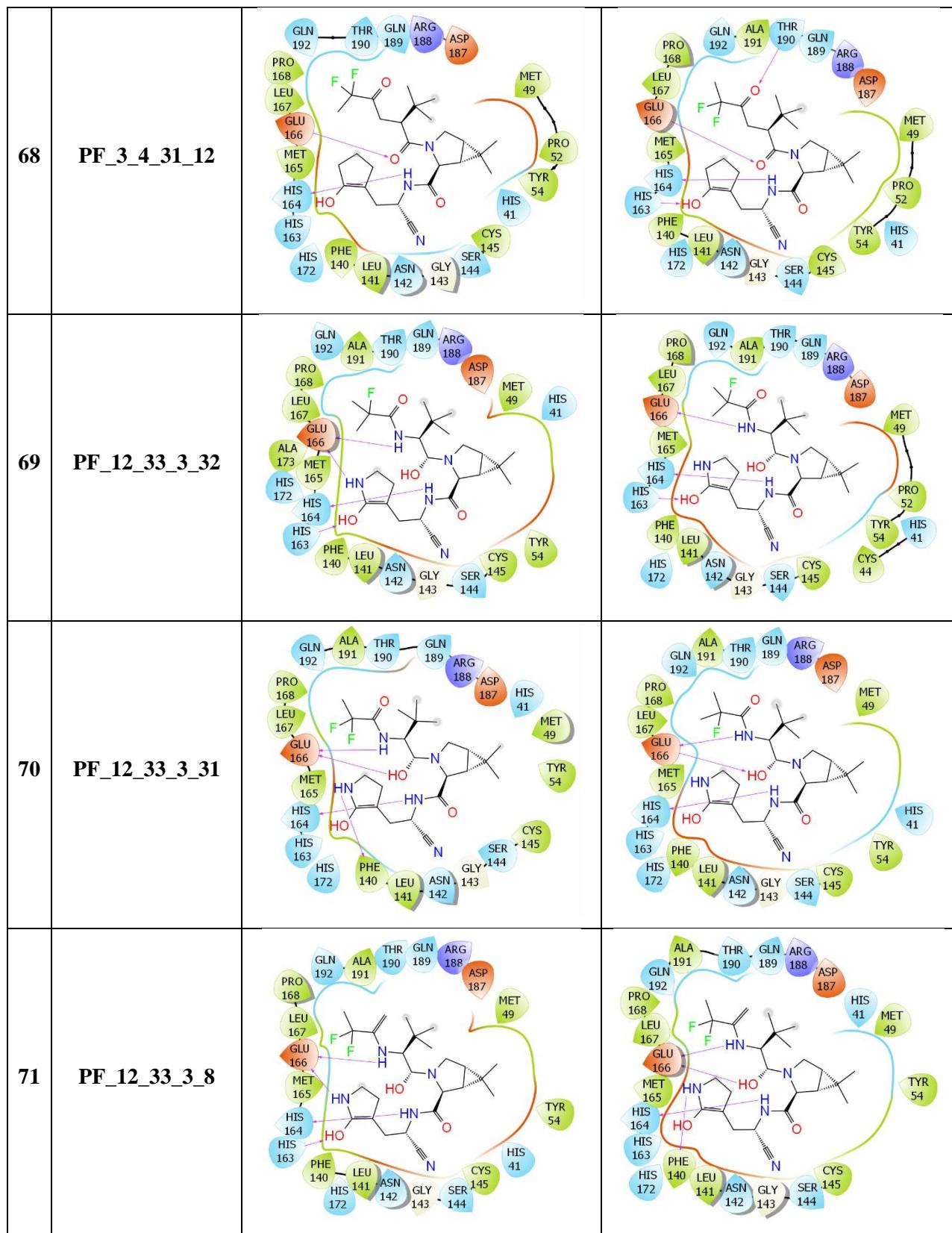


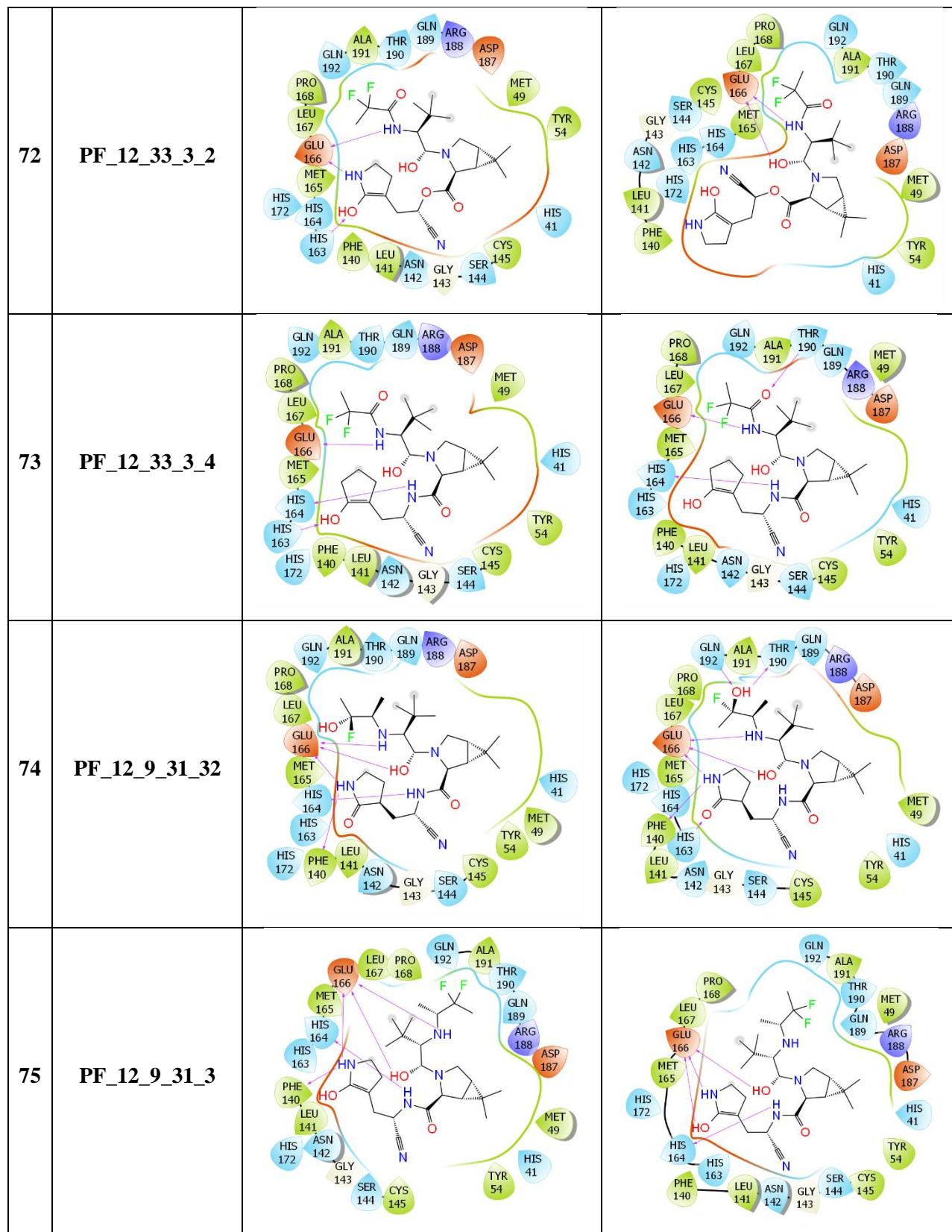


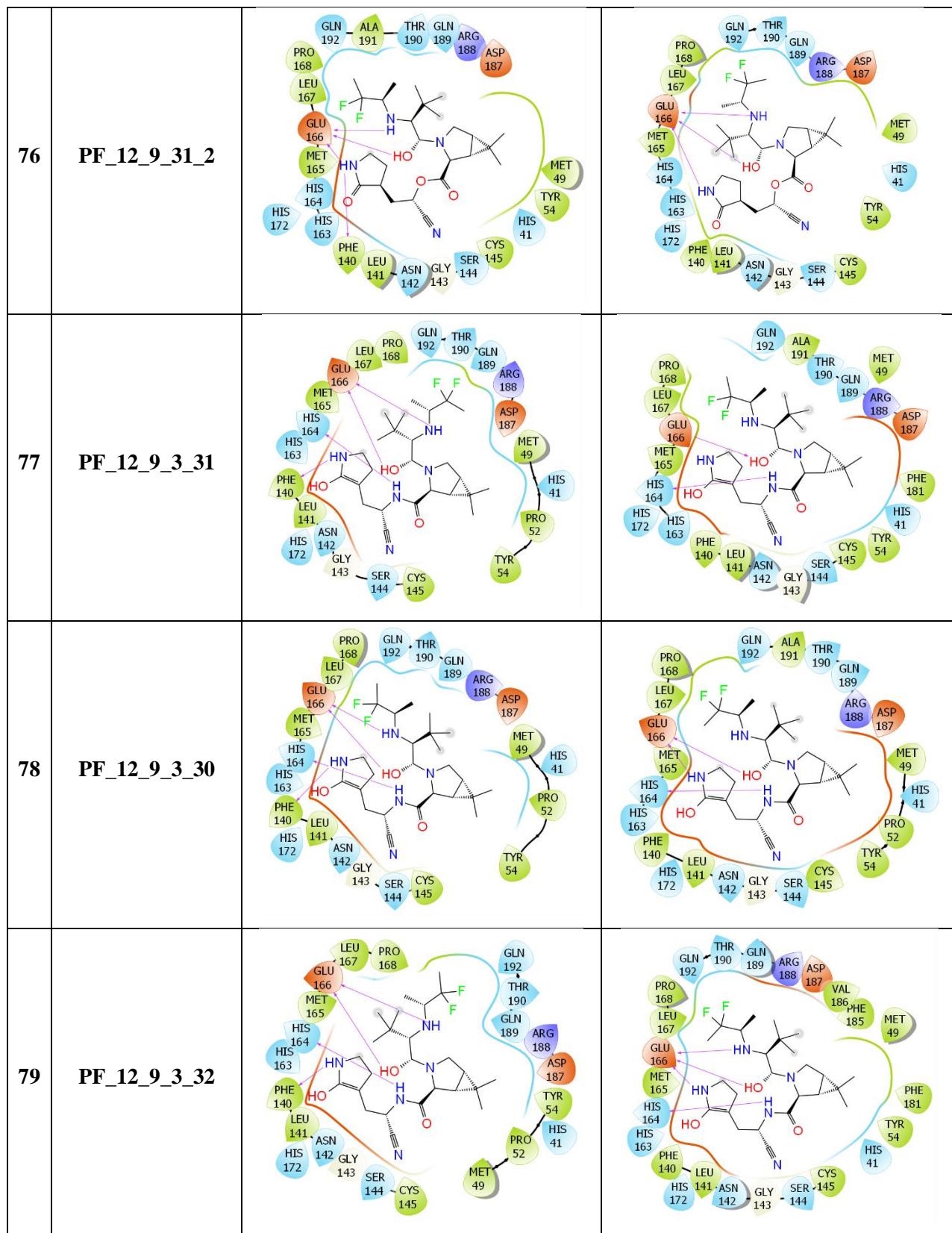


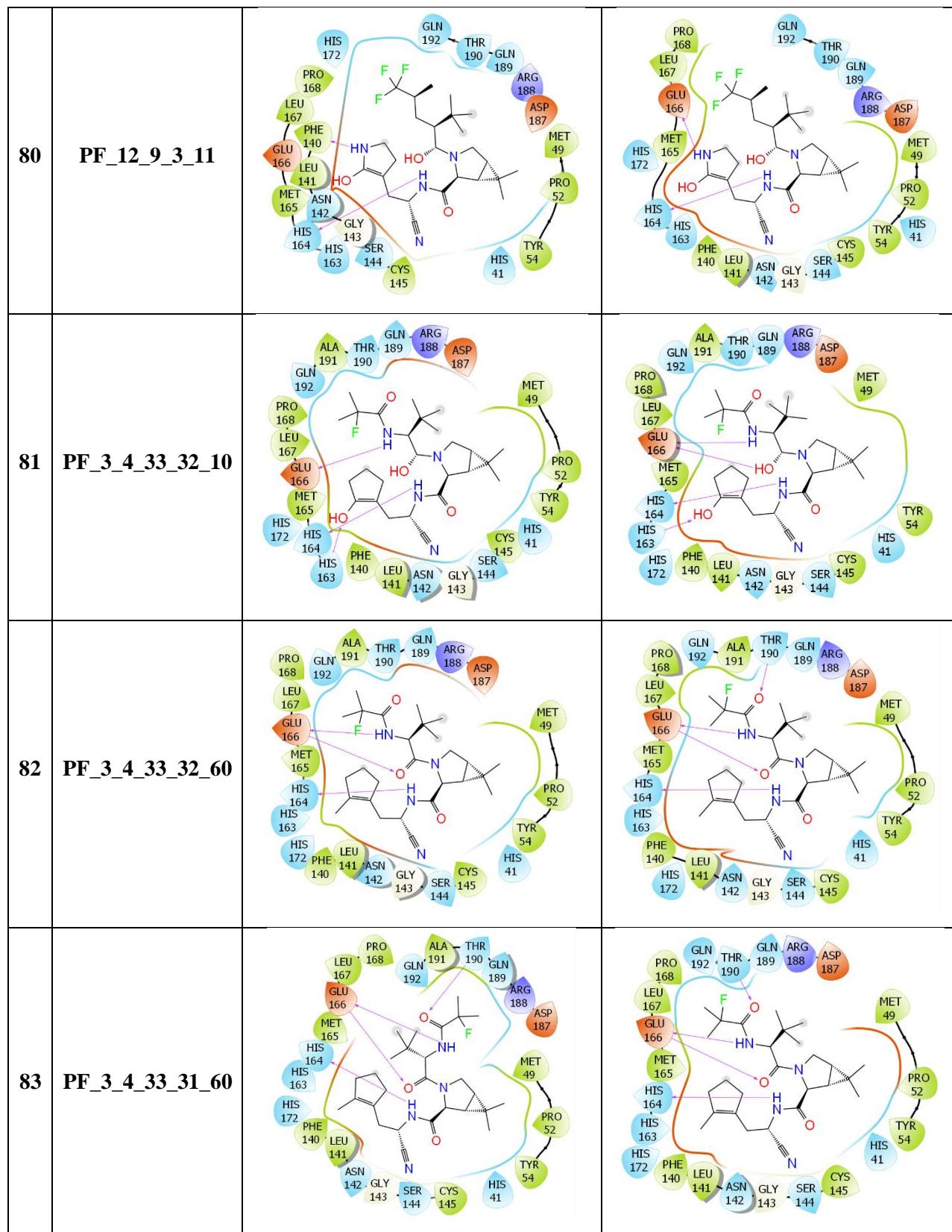


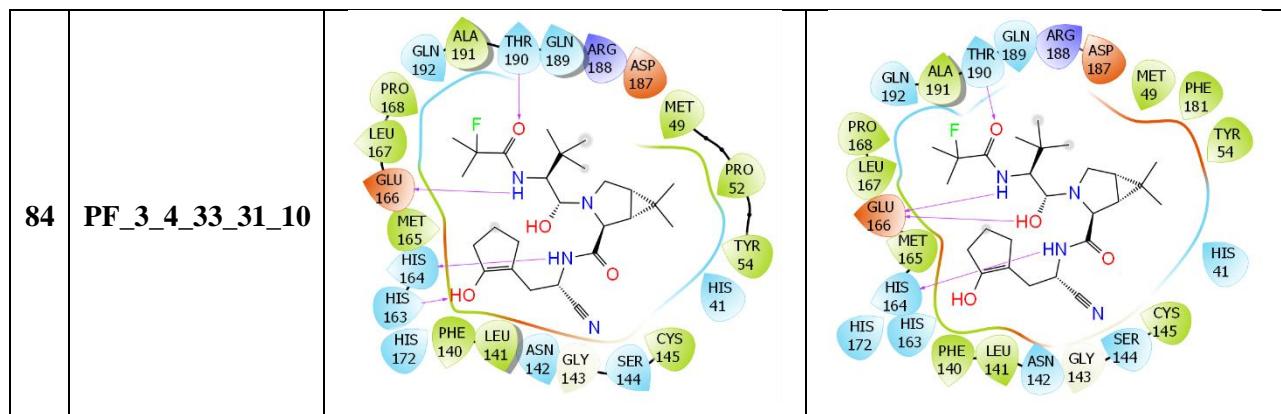
64	PF_3_4_33_31		
65	PF_3_4_33_12		
66	PF_3_4_31_32		
67	PF_3_4_31_10		











<sup>a</sup>The initial conformation of the complex, which was provided by DeepFrag application. <sup>b</sup>The MD-refined structure of the complex.

Table S8. Toxicity prediction via PreADMET.

Nº	Compound	CarcinoMouse	Ames <sub>test</sub>	TA100 <sub>10RLI</sub>	TA100 <sub>NA</sub>	TA1535 <sub>10RLI</sub>	TA1535 <sub>NA</sub>
1	<b>PF_3_4_62</b>	negative	non-mutagen	negative	negative	negative	negative
2	<b>PF_3_4_32</b>	negative	non-mutagen	negative	negative	negative	negative
3	<b>PF_3_4_33_32</b>	negative	non-mutagen	negative	negative	negative	negative
4	<b>PF_3_4_33_31</b>	negative	non-mutagen	negative	negative	negative	negative
5	<b>PF_12_9_31</b>	negative	non-mutagen	negative	negative	negative	negative
6	<b>PF_3_4_7</b>	negative	non-mutagen	negative	negative	negative	negative
7	<b>PF_9a_33</b>	negative	non-mutagen	negative	negative	negative	negative
8	<b>PF_3_4_33</b>	negative	non-mutagen	negative	negative	negative	negative
9	<b>PF_12_9_31_32</b>	negative	non-mutagen	negative	negative	negative	negative
10	<b>PF_3_4_32_32</b>	negative	non-mutagen	negative	negative	negative	negative
11	<b>PF_12_9_3</b>	negative	non-mutagen	negative	negative	negative	negative
12	<b>PF_12_33_3</b>	negative	non-mutagen	negative	negative	negative	negative
13	<b>PF_3_4_31</b>	negative	non-mutagen	negative	negative	negative	negative
14	<b>PF_3_4_31_32</b>	negative	non-mutagen	negative	negative	negative	negative
15	<b>PF-07321332</b>	positive	non-mutagen	positive	positive	positive	positive