

## Supporting Materials

**Table S1.** The PDB entries, the descriptions of the complexes and the experimental binding free energies

No.	PDB	$\Delta G_{\text{exp}}$	Descriptions	References
a1	1D3D	-12.39	alpha-thrombin/benzo[B]thiophene inhibitor 4	1
a2	1D3P	-10.08	alpha-thrombin/benzo[B]thiophene inhibitor 3	1
a3	1D3Q	-8.92	alpha-thrombin/benzo[B]thiophene inhibitor 2	1
a4	1D3T	-7.68	alpha-thrombin/benzo[B]thiophene inhibitor 1	1
a5	1DWB	-3.98	alpha-thrombin/benzamidine	2
a6	1DWC	-10.6	alpha-thrombin/MD-805	3
a7	1DWD	-11.57	alpha-thrombin/napap	4
b1	1AVD	-20.40	avidin/biotin	5
b2	×	-14.30	avidin/2'-iminobiotin	5
b3	×	-14.00	avidin/desthiobiotin	5
b4	×	-8.80	avidin/1'-Nmethoxycarbonylbiotinmethyl ester	5
b5	×	-8.20	avidin/D-4-n-hexyl-2-iminoimidazolidine	5
b6	×	-5.00	avidin/D-4-n-hexyloxazolidone	5
b7	×	-4.50	avidin/imidazolidone	5
c1	1AC4	-3.85	cytochrome C peroxidase/2,3,4-trimethyl-1,3-thiazole	6
c2	1AC8	-4.78	cytochrome C peroxidase/3,4,5-trimethylthiazole	6
c3	1AEB	-4.81	cytochrome C peroxidase/3-methylthiazole	6
c4	1AED	-5.86	cytochrome C peroxidase/3,4-dimethylthiazole	6
c5	1AEE	-3.96	cytochrome C peroxidase/aniline	6
c6	1AEF	-6.00	cytochrome C peroxidase/3-aminopyridine	6
c7	1AEG	-5.99	cytochrome C peroxidase/4-aminopyridine	6
c8	1AEH	-4.96	cytochrome C peroxidase/2-amino-4-methylthiazole	6
c9	1AEJ	-5.21	cytochrome C peroxidase/1-vinylimidazole	6
c10	1AEK	-4.92	cytochrome C peroxidase/indoline	6
c11	1AEM	-4.92	cytochrome C peroxidase/imidazo[1,2-A]pyridine	6
c12	1AEN	-7.07	cytochrome C peroxidase/2-amino-5-methylthiazole	6
c13	1AEO	-5.02	cytochrome C peroxidase/2-aminopyridine	6
c14	1AEQ	-4.73	cytochrome C peroxidase/2-ethylimidazole	6
c15	1AES	-4.33	cytochrome C peroxidase/imidazole	6
c16	1AET	-5.82	cytochrome C peroxidase/1-methylimidazole	6
c17	1AEU	-5.94	cytochrome C peroxidase/2-methylimidazole	6
c18	1AEV	-6.06	cytochrome C peroxidase/2-aminothiazole	6
d1	1NSC	-4.09	neuraminidase/sialic acid	7
d2	1NSD	-7.23	neuraminidase/dana	8
d3	2QWB	-3.74	mutated neuraminidase/sialic acid	9
d4	2QWC	-4.84	mutated neuraminidase/neu5ac2en	9
d5	2QWD	-6.61	mutated neuraminidase/4-amino-neu5ac2en	9

d6	2QWE	-10.20	mutated neuraminidase/4-guanidino-neu5ac2en	9
d7	2QWF	-7.73	mutated neuraminidase/g20	9
d8	2QWG	-11.45	mutated neuraminidase/g28	9
e1	2CPP	-7.90	P450cam/Camphor	10
e2	6CPP	-5.91	P450cam/Camphane	10
e3	×	-6.54	P450cam/Camphoroquinone	10
e4	7CPP	-5.57	P450cam/Norcamphor	10
e5	×	-5.52	P450cam/ethylbenzene	10
e6	×	-5.93	P450cam/3,3,5,5-tetramethylcyclohexanone	10
e7	8CPP	-7.53	P450cam/thiocamphore	10
e8	4CPP	-5.90	P450cam/Adamantine	10
e9	5CPP	-7.40	P450cam/adamantanone	10
f1	1APT	-12.83	penicillopepsin/pepstatin analogue	11
f2	1APU	-10.51	penicillopepsin/pepstatin analogue	11
f3	1APV	-12.27	penicillopepsin/iva-val-val(H)Dfo-n-methylamide	12
f4	1APW	-10.91	penicillopepsin/iva-val-valDfo-n-methylamide	9
f5	2WEA	-8.37	penicillopepsin/pp6	13
f6	2WEB	-7.03	penicillopepsin/pp4	13
f7	2WEC	-6.80	penicillopepsin/pp5	13

Table S2. The binding free energies for the  $\alpha$ -thrombin ligands based on two different solute dielectric constants ( $\epsilon_{in}= 1$  and  $\epsilon_{in}= 2$ ) calculated by MM/PBSA (kcal/mol)

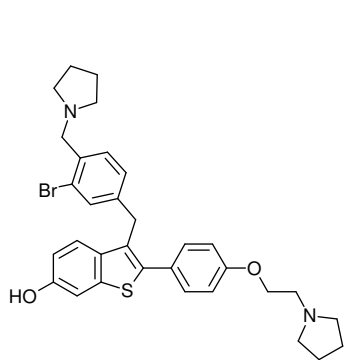
No.	$\Delta E_{vdw}$	$\Delta E_{ele}$	$\Delta G_{PB}$	$\Delta G_{SA}$	$\Delta E_{ele}+\Delta G_{PB}$	$\Delta E_{vdw}+\Delta G_{SA}$	T $\Delta S$	$\Delta G_{cal.}$	$\Delta G_{exp.}$
$\epsilon_{in}= 1$									
a1	-58.86 $\pm$ 1.09 <sup>a</sup>	-17.49 $\pm$ 0.98	57.19 $\pm$ 1.70	-7.61 $\pm$ 0.17	39.71 $\pm$ 0.73	-66.47 $\pm$ 1.27	-22.81 $\pm$ 1.57	-3.96 $\pm$ 1.03	-12.39
a2	-62.42 $\pm$ 1.69	-14.56 $\pm$ 9.43	60.57 $\pm$ 7.55	-7.65 $\pm$ 0.03	46.01 $\pm$ 1.88	-70.06 $\pm$ 1.72	20.20 $\pm$ 3.01	-3.85 $\pm$ 2.85	-10.08
a3	-52.69 $\pm$ 1.18	-8.61 $\pm$ 1.06	49.43 $\pm$ 1.78	-6.68 $\pm$ 0.24	40.82 $\pm$ 0.72	-59.37 $\pm$ 1.42	-20.01 $\pm$ 0.65	1.47 $\pm$ 1.36	-8.92
a4	-50.58 $\pm$ 2.58	-1.85 $\pm$ 0.27	37.73 $\pm$ 0.54	-6.42 $\pm$ 0.13	35.88 $\pm$ 0.27	57.00 $\pm$ 2.71	-17.97 $\pm$ 1.06	-3.16 $\pm$ 1.38	-7.68
a5	-18.73 $\pm$ 0.24	-133.38 $\pm$ 1.13	193.46 $\pm$ 0.23	-2.79 $\pm$ 0.00	60.08 $\pm$ 1.36	-21.51 $\pm$ 0.24	-15.50 $\pm$ 0.82	54.07 $\pm$ 0.79	-3.98
a6	-57.79 $\pm$ 0.46	-72.31 $\pm$ 0.44	121.35 $\pm$ 0.61	-7.11 $\pm$ 0.01	49.04 $\pm$ 0.17	-64.89 $\pm$ 0.45	-24.99 $\pm$ 1.73	9.14 $\pm$ 2.01	-10.6
a7	-58.27 $\pm$ 0.17	-162.49 $\pm$ 0.25	249.30 $\pm$ 0.92	-7.02 $\pm$ 0.04	86.81 $\pm$ 1.18	-65.29 $\pm$ 0.13	-22.53 $\pm$ 0.33	44.05 $\pm$ 0.98	-11.57
$\epsilon_{in}= 2$									
a1	-58.86 $\pm$ 1.09 <sup>a</sup>	-8.74 $\pm$ 0.49	28.29 $\pm$ 0.70	-7.61 $\pm$ 0.17	19.54 $\pm$ 0.22	-66.47 $\pm$ 1.27	-22.81 $\pm$ 1.57	-24.12 $\pm$ 0.52	-12.39
a2	-62.42 $\pm$ 1.69	-7.28 $\pm$ 4.71	29.74 $\pm$ 3.64	-7.65 $\pm$ 0.03	22.46 $\pm$ 1.07	-70.06 $\pm$ 1.72	20.20 $\pm$ 3.01	-27.40 $\pm$ 3.65	-10.08
a3	-52.69 $\pm$ 1.18	-4.30 $\pm$ 0.53	23.97 $\pm$ 0.88	-6.68 $\pm$ 0.24	19.66 $\pm$ 0.35	-59.37 $\pm$ 1.42	-20.01 $\pm$ 0.65	-19.69 $\pm$ 1.73	-8.92
a4	-50.58 $\pm$ 2.58	-0.93 $\pm$ 0.13	18.23 $\pm$ 0.31	-6.42 $\pm$ 0.13	17.31 $\pm$ 0.17	57.00 $\pm$ 2.71	-17.97 $\pm$ 1.06	-21.73 $\pm$ 1.47	-7.68
a5	-18.73 $\pm$ 0.24	-66.69 $\pm$ 0.56	93.87 $\pm$ 0.16	-2.79 $\pm$ 0.00	27.18 $\pm$ 0.72	-21.51 $\pm$ 0.24	-15.50 $\pm$ 0.82	21.16 $\pm$ 0.15	-3.98
a6	-57.79 $\pm$ 0.46	-36.15 $\pm$ 0.22	59.07 $\pm$ 0.27	-7.11 $\pm$ 0.01	22.92 $\pm$ 0.05	-64.89 $\pm$ 0.45	-24.99 $\pm$ 1.73	-16.99 $\pm$ 2.13	-10.6
a7	-58.27 $\pm$ 0.17	-81.24 $\pm$ 0.13	120.87 $\pm$ 0.44	-7.02 $\pm$ 0.04	39.63 $\pm$ 0.57	-65.29 $\pm$ 0.13	-22.53 $\pm$ 0.33	-3.14 $\pm$ 0.37	-11.57

Table S3. The experimental and the predicted binding free energies and the corresponding energy terms calculated by MM/GBSA (kcal/mol)

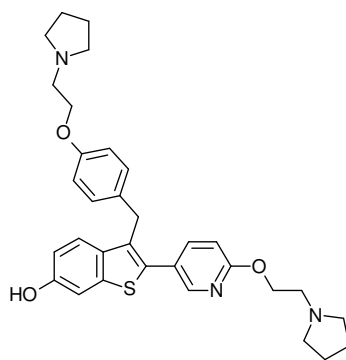
No.	$\Delta E_{\text{vdw}}$	$\Delta E_{\text{ele}}$	$\Delta G_{\text{GB}}$	$\Delta G_{\text{SA}}$	$\Delta E_{\text{ele}}+\Delta G_{\text{GB}}$	$\Delta E_{\text{vdw}}+\Delta G_{\text{SA}}$	TAS	$\Delta G_{\text{cal.}}$	$\Delta G_{\text{exp.}}$
$\alpha$ -thrombin ( $\epsilon_{\text{in}} = 4$ )									
a1	-58.86±1.09 <sup>a</sup>	-8.75±0.49	17.62±0.25	-7.61±0.17	8.87±0.23	-66.47±1.27	-22.81±1.57	-34.79±0.07	-12.39
a2	-62.42±1.69	-7.28±4.71	18.84±2.34	-7.65±0.03	11.56±2.37	-70.06±1.72	20.20±3.01	-38.31±2.35	-10.08
a3	-52.69±1.18	-4.30±0.53	15.17±0.75	-6.68±0.24	10.86±0.22	-59.37±1.42	-20.01±0.65	-28.49±1.85	-8.92
a4	-50.58±2.58	-0.93±0.13	11.48±0.39	-6.42±0.13	10.56±0.25	57.00±2.71	-17.97±1.06	-28.48±1.39	-7.68
a5	-18.73±0.24	-66.69±0.56	61.51±0.59	-2.79±0.00	5.18±0.02	-21.51±0.24	-15.50±0.82	-11.20±0.60	-3.98
a6	-57.79±0.46	-36.15±0.22	44.74±0.21	-7.11±0.01	8.58±0.01	-64.89±0.45	-24.99±1.73	-31.32±2.19	-10.6
a7	-58.27±0.17	-81.24±0.13	82.18±0.10	-7.02±0.04	0.94±0.02	-65.29±0.13	-22.53±0.33	-41.83±0.17	-11.57
avidin ( $\epsilon_{\text{in}} = 1$ )									
b1	-30.31±0.30	-172.54±4.72	157.25±4.70	-4.21±0.01	-15.29±0.02	-34.52±0.31	-18.69±0.22	-31.12±0.50	-20.4
b2	-30.10±0.83	-156.76±4.74	145.28±4.07	-4.27±0.04	-11.48±0.67	-34.57±0.79	-20.44±0.65	-25.41±0.81	-14.3
b3	-28.80±0.19	-165.47±5.49	147.76±3.32	-4.21±0.01	-17.71±2.17	-33.01±0.17	-21.33±0.43	-29.39±1.92	-14.0
b4	-40.31±0.14	-28.99±0.50	37.95±0.22	-5.55±0.06	8.96±0.72	-45.86±0.19	-17.64±1.61	-19.44±2.14	-8.8
b5	-28.52±1.39	-21.67±7.19	23.89±3.15	-4.09±0.04	2.22±4.03	-32.61±1.43	-15.19±1.03	-15.21±3.63	-8.2
b6	-25.53±1.00	-21.97±0.20	22.87±0.01	-3.69±0.28	0.90±0.19	-29.22±1.28	-14.30±1.61	-14.01±0.14	-5.0
b7	-11.99±0.16	-25.11±0.30	19.93±0.11	-2.12±0.01	-5.18±0.14	-14.10±0.16	-13.57±1.08	-5.71±0.78	-4.5
cytochrome C peroxidase ( $\epsilon_{\text{in}} = 1$ )									
c1	-25.60±0.02	-352.29±0.21	354.69±0.20	-2.81±0.01	2.40±0.01	-28.41±0.03	-15.50±0.09	-10.51±0.06	-3.85
c2	-25.84±0.06	-354.14±0.11	354.21±0.08	-2.79±0.02	0.06±0.03	-28.63±0.04	-16.87±1.10	-11.69±1.10	-4.78
c3	-17.90±0.91	-361.70±3.03	361.23±2.79	-1.93±0.43	-0.47±0.24	-19.82±1.34	-14.53±0.06	-5.76±1.16	-4.81
c4	-22.28±0.15	-353.88±1.88	353.41±1.97	-2.62±0.00	-0.47±0.10	-24.90±0.16	-15.05±1.02	-10.31±0.96	-5.86
c5	-18.07±0.07	-365.62±6.40	362.08±4.85	-2.47±0.02	-3.54±1.55	-20.53±0.05	-13.19±1.28	-10.88±2.88	-3.96
c6	-17.74±0.48	-377.44±0.95	372.47±0.56	-2.86±0.44	-4.97±0.39	-20.60±0.05	-15.63±0.03	-9.94±0.41	-6.00
c7	-17.94±0.12	-372.64±3.54	369.19±3.19	-2.40±0.01	-3.45±0.35	-20.34±0.12	-15.73±0.04	-8.06±0.50	-5.99
c8	-21.33±0.03	-360.68±2.15	359.78±2.11	-2.49±0.00	-0.90±0.04	-23.83±0.03	-15.50±0.40	-9.23±0.47	-4.96
c9	-17.09±0.41	-372.32±1.53	367.54±1.25	-2.44±0.00	-4.77±0.28	-19.52±0.41	-15.44±0.14	-8.86±0.01	-5.21
c10	-20.72±0.31	-372.36±0.79	366.06±1.03	-3.29±0.43	-6.30±0.24	-24.01±0.74	-15.45±2.08	-14.86±1.11	-4.92
c11	-21.89±0.08	-363.77±3.15	358.94±3.22	-2.64±0.01	-4.84±0.08	-24.53±0.06	-14.50±0.33	-14.86±0.19	-4.92

c12	-18.40±0.08	-379.84±2.79	366.23±3.09	-2.48±0.00	-13.61±0.30	-20.88±0.08	-14.39±0.48	-20.10±0.69	-7.07
c13	-16.64±0.04	-381.35±2.81	367.75±2.68	-2.78±0.4	-13.61±0.13	-19.42±0.46	-13.81±0.31	-19.22±0.02	-5.02
c14	-19.34±0.12	-366.62±1.23	362.71±1.06	-2.62±0.0	-3.91±0.17	-21.96±0.12	-15.22±0.93	-10.64±0.88	-4.73
c15	-11.42±0.03	-372.89±0.16	366.57±0.51	-2.42±0.4	-6.32±0.35	-13.84±0.42	-12.09±0.29	-8.07±1.05	-4.33
c16	-15.83±0.06	-375.74±2.87	369.63±2.73	-2.26±0.0	-6.10±0.14	-18.09±0.06	-15.19±0.04	-9.00±0.16	-5.82
c17	-16.35±0.00	-370.52±1.42	365.21±1.33	-2.34±0.0	-5.31±0.09	-18.69±0.01	-14.95±0.68	-9.05±0.77	-5.94
c18	-15.64±0.13	-369.60±2.73	363.27±1.55	-2.35±0.0	-6.33±1.18	-17.98±0.14	-16.93±0.07	-7.38±0.96	-6.06
neuraminidase ( $\epsilon_{in} = 4$ )									
d1	-22.63±0.30	-29.30±0.41	23.53±0.13	-3.99±0.3	-5.77±0.54	-26.61±0.08	-21.97±1.02	-10.41±0.56	-4.09
d2	-22.95±0.57	-33.69±1.21	27.53±0.89	-4.65±0.0	-6.16±0.31	-27.60±0.59	-15.78±2.65	-17.97±2.93	-7.23
d3	-21.19±0.17	-26.07±0.54	22.73±0.49	-4.45±0.0	-3.34±1.02	-25.64±0.17	-21.54±0.07	-7.43±0.78	-3.74
d4	-21.84±0.52	-22.36±0.17	20.43±0.30	-4.64±0.0	-1.93±0.13	-26.49±0.53	-20.75±2.30	-7.67±2.96	-4.84
d5	-20.12±0.64	-23.62±0.04	21.98±1.10	-4.58±0.0	-1.64±1.06	-24.71±0.68	-19.03±0.08	-7.31±1.66	-6.61
d6	-28.40±0.03	-22.53±3.39	20.18±2.05	-4.99±0.0	-2.35±1.33	-33.39±0.05	-19.29±1.83	-16.45±3.11	-10.2
d7	-27.83±0.38	-3.53±0.35	7.02±0.37	-4.98±1.1	3.69±0.02	-32.81±1.55	-22.86±2.01	-6.26±0.48	-7.73
d8	-29.43±0.01	-13.36±0.42	14.97±0.53	-4.49±0.4	1.42±0.95	-33.92±0.46	-17.72±2.16	-14.77±2.65	-11.45
P450cam ( $\epsilon_{in} = 2$ )									
e1	-27.57±0.61	-4.55±0.13	5.37±0.06	-2.78±0.5	0.82±0.07	-30.35±0.03	-15.25±0.85	-14.29±0.75	-7.90
e2	-26.94±0.28	-0.07±0.02	1.16±0.05	-3.33±0.0	1.09±0.03	-30.27±0.25	-16.63±0.62	-12.55±0.34	-5.91
e3	-29.03±0.24	-4.30±0.06	6.86±0.19	-3.38±0.0	2.57±0.13	-31.41±0.24	-14.45±1.76	-15.39±1.39	-6.54
e4	-20.35±0.05	-0.93±0.05	3.61±0.12	-2.79±0.0	2.69±0.07	-23.14±0.03	-15.04±0.27	-5.41±0.31	-5.57
e5	-21.13±0.24	-0.12±0.04	3.89±0.03	-2.37±0.5	3.78±0.07	-23.50±0.78	-14.40±0.32	-5.32±0.39	-5.52
e6	-27.68±0.36	-4.51±1.17	5.00±0.57	-3.56±0.0	0.49±0.60	-31.25±0.35	-15.31±0.54	-15.44±0.30	-5.93
e7	-29.42±0.10	-2.61±0.08	4.21±0.04	-3.48±0.0	2.05±0.04	-32.89±0.10	-13.24±0.42	-17.61±0.28	-7.53
e8	-26.36±0.04	-0.01±0.0	1.95±0.01	-2.60±0.5	1.94±0.01	-28.96±0.51	-13.99±0.56	-13.03±0.04	-5.90
e9	-28.09±0.11	-1.44±0.04	4.54±0.03	-3.15±0.0	3.10±0.01	-31.23±0.10	-15.10±0.60	-13.03±0.71	-7.40
penicillopepsin ( $\epsilon_{in} = 2$ )									
f1	-50.93±1.01	-17.01±2.12	28.28±1.51	-8.59±0.0	11.27±0.61	-59.52±1.02	-22.43±1.30	-22.81±0.33	-12.83
f2	-41.34±1.58	-12.23±3.30	21.88±2.62	-6.78±0.1	9.65±0.68	-48.12±1.76	-22.01±0.44	-16.64±1.52	-10.51
f3	-48.82±1.24	-33.59±0.82	34.00±1.20	-6.34±0.1	0.42±0.42	-55.16±1.37	-24.89±0.64	-29.86±1.63	-12.27

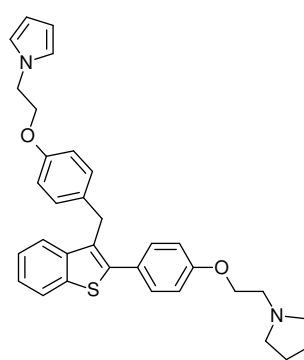
f4	-40.62±2. 30	-13.06±3. 58	19.72±1.9 2	-6.79±0.2 9	6.66±1.66	-47.41±2.59	-16.09±1. 52	-24.65±5. 77	-10.91
f5	-38.63±4. 78	-7.32±0.6 2	16.34±1.5 5	-5.48±0.8 5	9.02±0.93	-44.10±5.63	-20.06±0. 16	-15.02±4. 86	-8.37
f6	-42.23±4. 33	-8.75±0.9 9	20.74±1.2 1	-6.60±0.0 5	11.99±0.22	-48.83±4.28	-21.31±0. 23	-15.53±3. 83	-7.03
f7	-45.28±2. 23	-15.07±2. 75	25.78±3.2 3	-7.51±0.4 8	10.72±0.48	-52.79±2.72	-23.67±2. 29	-18.41±0. 06	-6.80



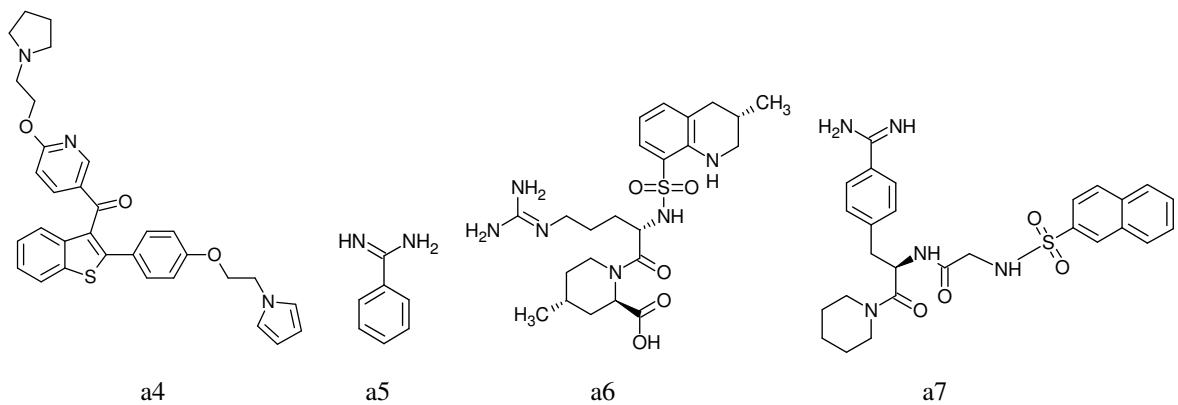
a1



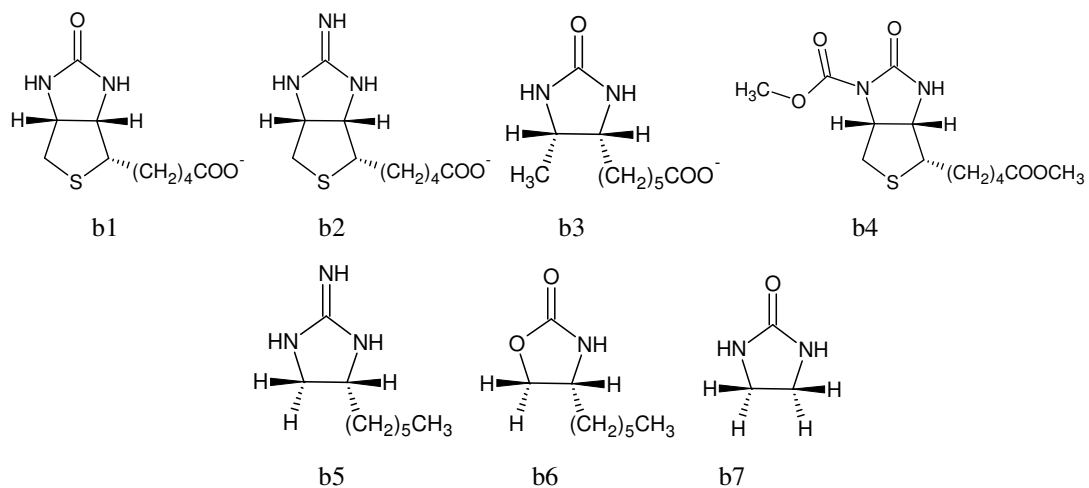
a2



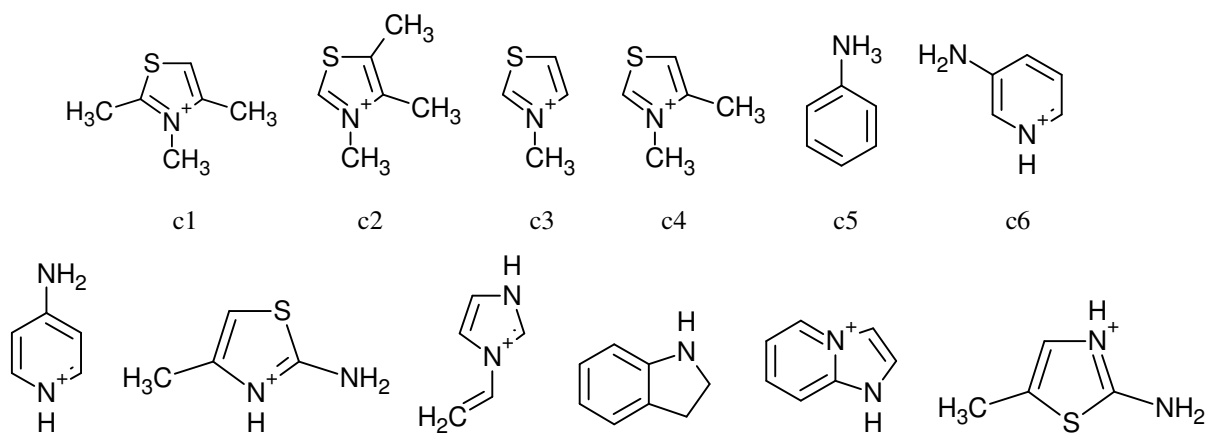
a3



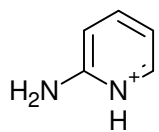
$\alpha$ -thrombin inhibitors



avidin inhibitors

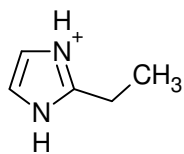


c7



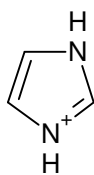
c13

c8



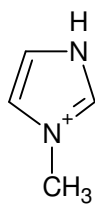
c14

c9



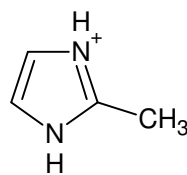
c15

c10



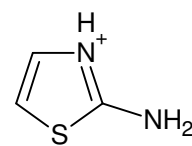
c16

c11



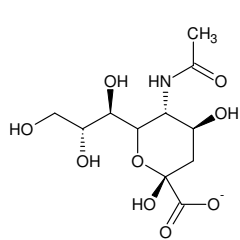
c17

c12

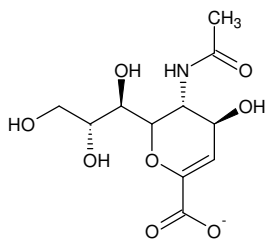


c18

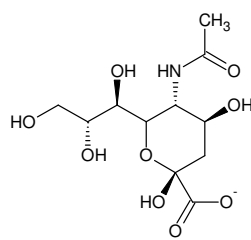
### cytochrome C peroxidase inhibitors



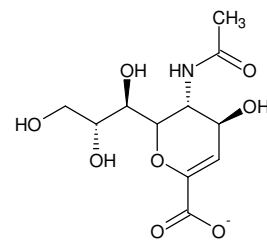
d1



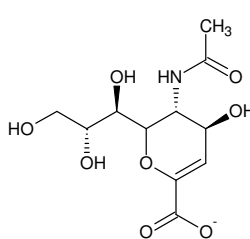
d2



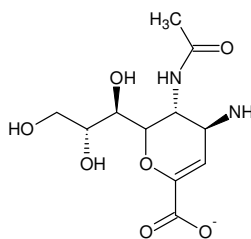
d3



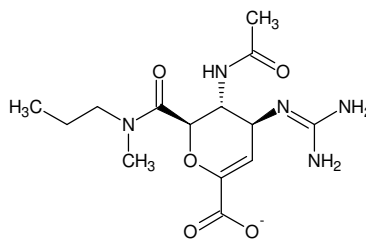
d4



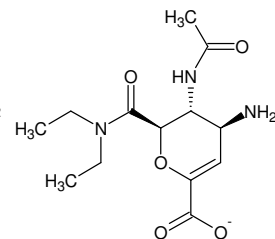
d5



d6

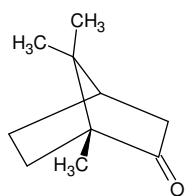


d7

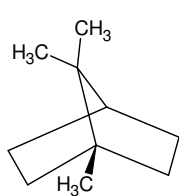


d8

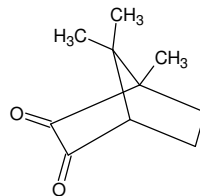
### neuraminidase inhibitors



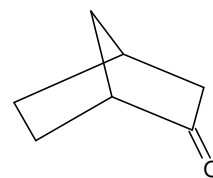
e1



e2

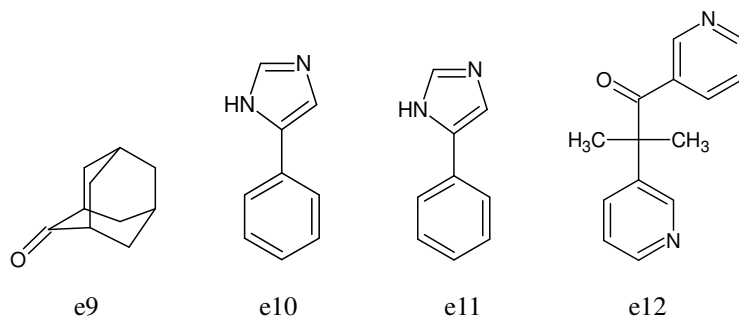
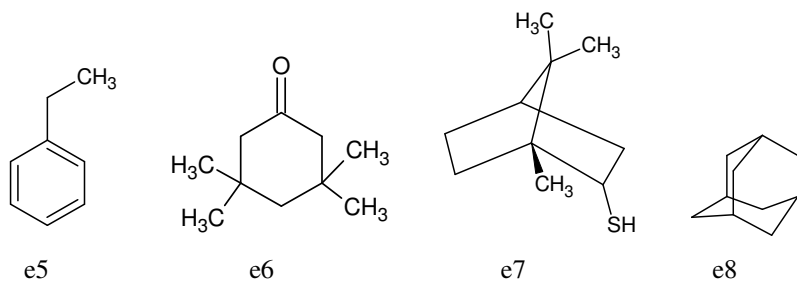


e3

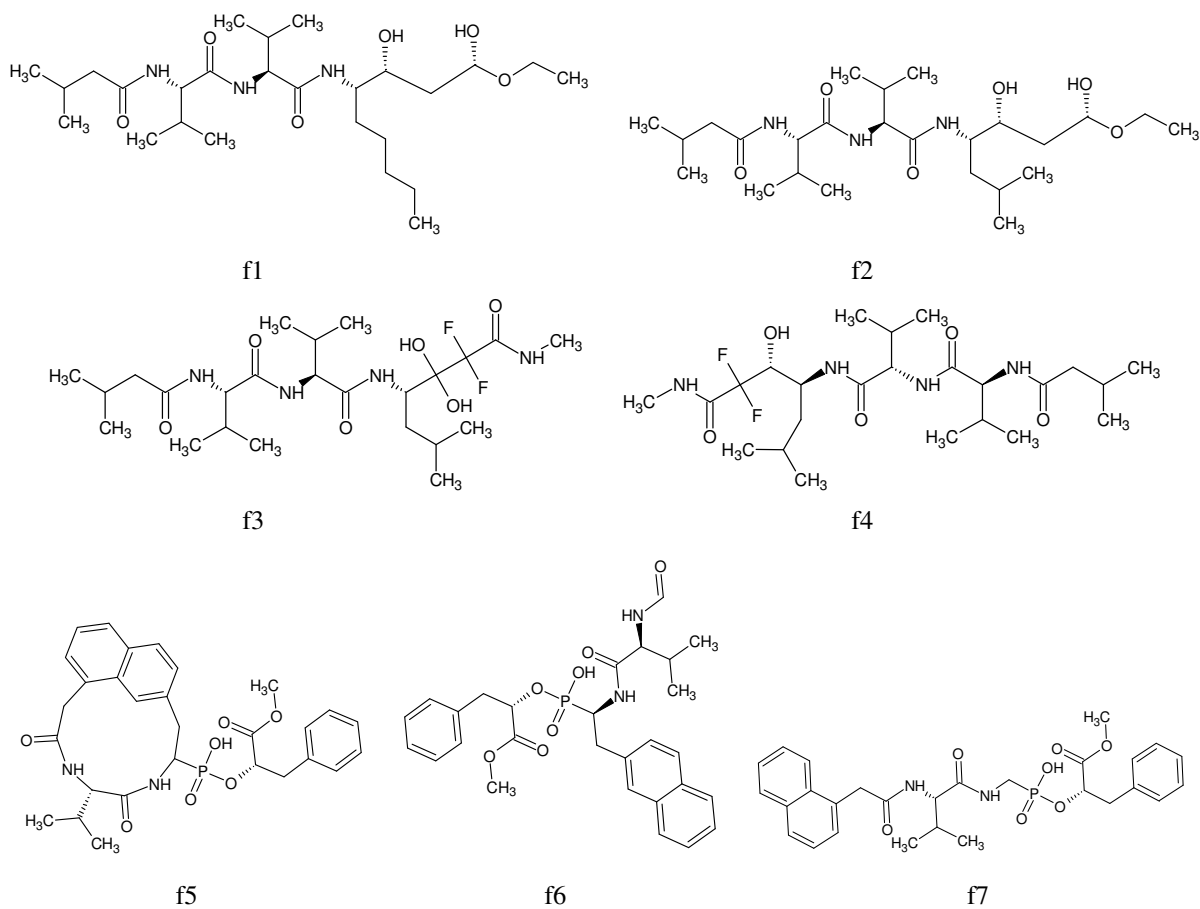


e4





P450cam inhibitors



penicillopepsin inhibitors

**Figure S1.** Structures of the inhibitors studied: (a1~a7) al-thrombin inhibitors, (b1~b7) avidin inhibitors, (c1~c18) cytochrome C peroxidase inhibitors, (d1~d8) neuraminidase inhibitors, (e1~e12) P450cam inhibitors, (f1~f7) penicillopepsin inhibitors. The name of the inhibitors can be found in Table S1 in the supplementary materials.

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