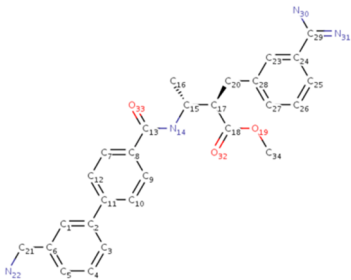
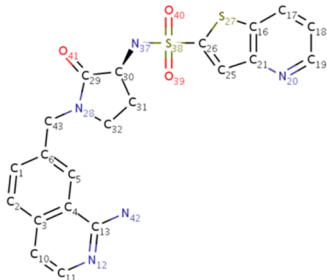
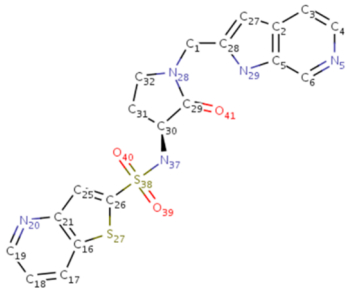
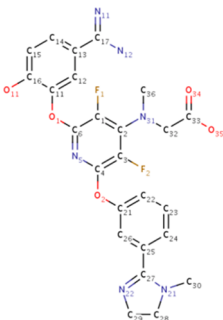
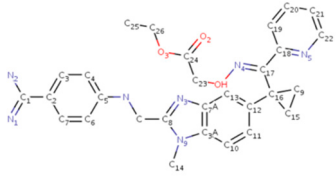
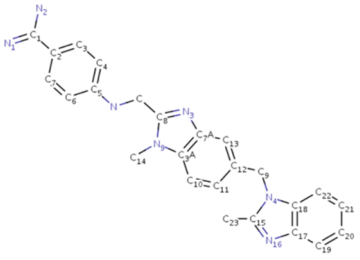
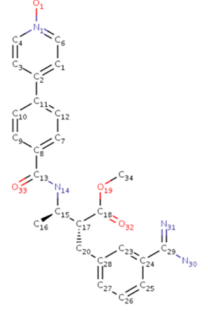
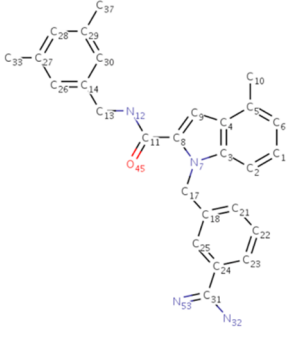
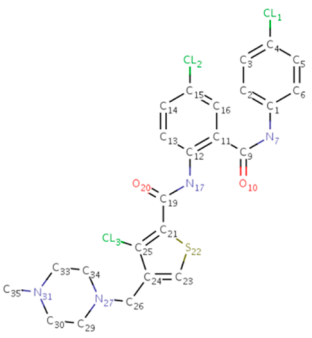


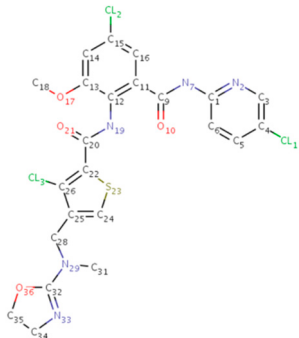
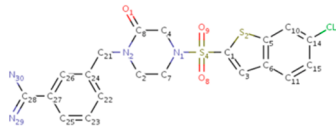
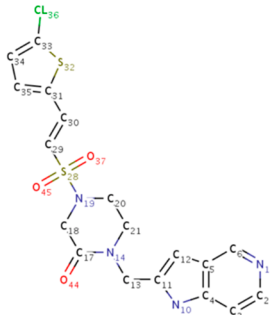
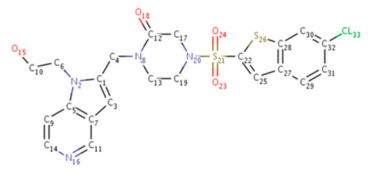
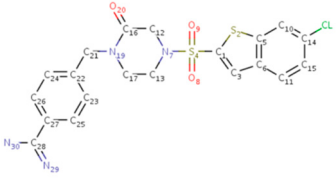
Supplementary Materials: AutoDock-GIST: Incorporating Thermodynamics of Active-Site Water into Scoring Function for Accurate Protein-Ligand Docking

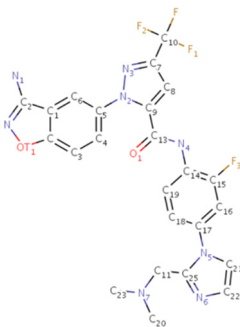
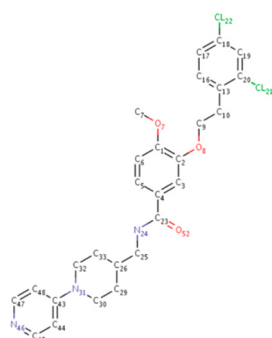
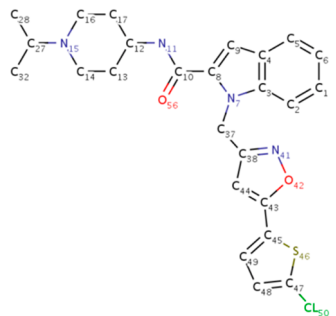
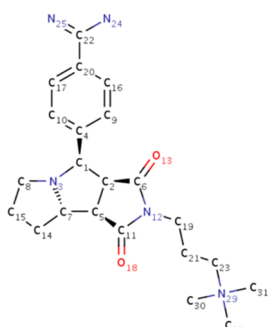
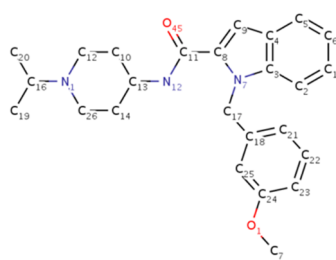
Shota Uehara and Shigenori Tanaka

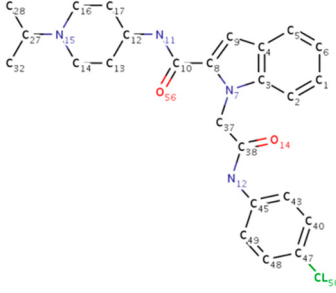
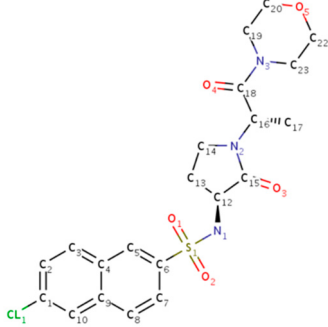
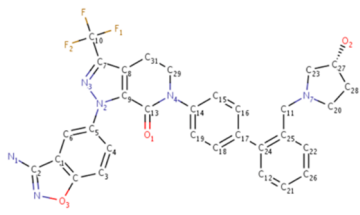
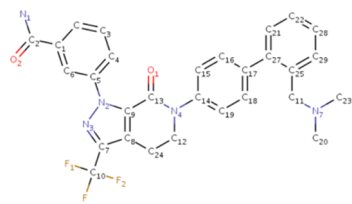
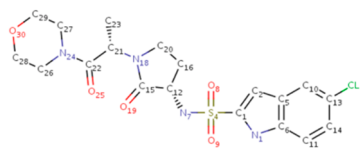
Table S1. 28 ligands of coagulation factor Xa for the training set.

ID	PDB-ID/ res-ID	Chemical Structure	Compound Name	MW [g/mol]	ΔG_{exp} [kcal/mol] (Supplementary Reference)
1	1EZQ/RPR		3-[(3'-Aminomethyl-biphenyl-4-carbonyl)-amino]-2-(3-carbamimidoyl-benzyl)-butyric acid methyl ester	458.5	-12.21 [1]
2	1F0R/815		Thieno[3,2-b]pyridine-2-sulfonic acid [1-(1-amino-isoquinolin-7-ylmethyl)-2-oxo-pyrrolidin-3-yl]-amide	453.5	-10.34 [2]
3	1F0S/PR2		Thieno[3,2-b]pyridine-2-sulfonic acid [2-oxo-1-(1H-pyrrolo[2,3-c]pyridin-2-ylmethyl)-pyrrolidin-3-yl]-amide	427.5	-10.45 [2]
4	1FJS/Z34		N-[2-[5-[Amino(imino)methyl]-2-hydroxyphenoxy]-3,5-difluoro-6-[3-(4,5-dihydro-1-methyl-1H-imidazol-2-yl)phenoxy]pyridin-4-yl]-N-methylglycine	526.5	-13.44 [3]

5	1G2L/T87		[(1-[2-(4-Carbamimidoyl-phenylamino)-methyl]-1-methyl-1H-benzimidazol-5-yl)-cyclopropyl]-pyridin-2-yl-methyleneaminoxy]-acetic acid ethyl ester	525.6	-10.28 [4]
6	1G2M/R11		4-([1-Methyl-5-(2-methyl-benzimidazol-1-ylmethyl)-1H-benzimidazol-2-ylmethyl]-amino)-benzamidine	423.5	-10.49 [4]
7	1KSN/FXV		Methyl-3-(4'-N-oxopyridylphenyl)-3-methyl-2-(m-amidinobenzyl)-propionate	447.5	-12.82 [5]
8	1LQD/CMI		1-(3-Carbamimidoyl-benzyl)-4-methyl-1H-indole-2-carboxylic acid 3,5-dimethylbenzylamide	424.5	-10.97 [6]
9	1MQ5/XLC		3-Chloro-N-[4-chloro-2-[[4-(4-chlorophenyl)amino]carbonyl]phenyl]-4-[(4-methyl-1-piperazinyl)methyl]-2-thiophenecarboxamide	537.9	-12.31 [7]

10	1MQ6/XLD		3-Chloro-N-[(5-chloro-2-pyridinyl)amino]carbonyl-6-methoxyphenyl-4-[[4,5-dihydro-2-oxazolyl)methylamino]methyl]-2-thiophenecarboxamide	568.9	-15.06 [7]
11	1NFU/RRP		3-({4-[(6-Chloro-1-benzothien-2-yl)sulfonyl]-2-oxopiperazin-1-yl)methyl}benzenecarboximidamide	463.0	-10.45 [1]
12	1NFW/RRR		4-[(e)-2-(5-Chlorothien-2-yl)vinyl)sulfonyl]-1-(1h-pyrrolo[3,2-c]pyridin-2-yl)methyl)piperazin-2-one	436.9	-12.09 [1]
13	1NFX/RDR		4-[(6-Chloro-1-benzothien-2-yl)sulfonyl]-1-[[1-(2-hydroxyethyl)-1H-pyrrolo[3,2-c]pyridin-2-yl)methyl]piperazin-2-one	505.0	-11.51 [1]
14	1NFY/RTR		4-({4-[(6-Chloro-1-benzothien-2-yl)sulfonyl]-2-oxopiperazin-1-yl)methyl}benzenecarboximidamide	463.0	-12.00 [1]

15	1Z6E/IK8		1-(3-Amino-1,2-benzisoxazol-5-yl)-N-(4-{2-[(dimethylamino)methyl]-1h-imidazol-1-yl}-2-fluorophenyl)-3-(trifluoromethyl)-1H-pyrazole-5-carboxamide	528.5	-13.12 [8]
16	2BMG/I1H		3-[2-(2,4-Dichlorophenyl)ethoxy]-4-methoxy-N-[(1-pyridin-4-yl)piperidin-4-yl)methyl]benzamide	514.4	-10.56 [9]
17	2BOH/IIA		1-[[5-(5-Chloro-2-thienyl)isoxazol-3-yl)methyl]-N-(1-isopropylpiperidin-4-yl)-1h-indole-2-carboxamide	483.0	-11.62 [10]
18	2BOK/784		[Amino (4-{(3as,4r,8as,8br)-1,3-dioxo-2-[3-(trimethylammonio)propyl]decahydropyrrolo[3,4-a]pyrrolizin-4-yl)phenyl)methylene]ammonium	398.5	-9.173 [11]
19	2BQ7/IIID		N-(1-Isopropylpiperidin-4-yl)-1-(3-methoxybenzyl)-1H-indole-2-carboxamide	405.5	-9.61 [10]

20	2BQW/III		1-[(4-Chlorophenyl)amino]-2-oxoethyl)-N-(1-isopropylpiperidin-4-yl)-1H-indole-2-carboxamide	453.0	-11.62 [10]
21	2CJI/GSK		6-Chloro-N-[(3s)-1-[(1s)-1-methyl-2-(4-morpholinyl)-2-oxoethyl]-2-oxo-3-pyrrolidinyl]-2-naphthalenesulfonamide	466.0	-11.21 [12]
22	2FZZ/5QC		1-(3-Amino-1,2-benzisoxazol-5-yl)-6-(2'-[[[(3r)-3-hydroxypyrrolidin-1-yl]methyl]biphenyl-4-yl]-3-(trifluoromethyl)-1,4,5,6-tetrahydro-7h-pyrazolo[3,4-c]pyridin-7-one	588.6	-14.21 [13]
23	2G00/4QC		3-[6-(2'-[[[(Dimethylamino)methyl]biphenyl-4-yl]-7-oxo-3-(trifluoromethyl)-4,5,6,7-tetrahydro-1h-pyrazolo[3,4-c]pyridin-1-yl]benzamide	533.5	-13.16 [13]
24	2J2U/GSQ		5-Chloro-N-[(3s)-1-[(1s)-1-methyl-2-morpholin-4-yl]-2-5-chloro-n-[(3s)-1-[(1s)-1-methyl-2-morpholin-4-yl]-2-sulfonamide	454.9	-9.61 [14]

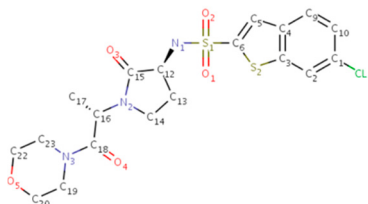
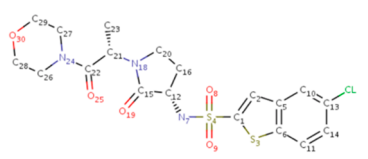
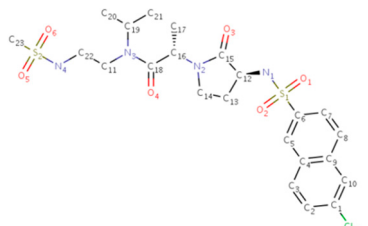
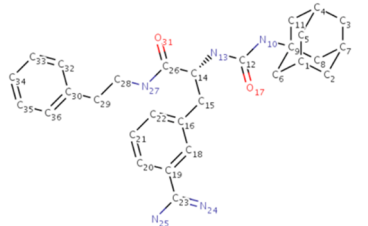
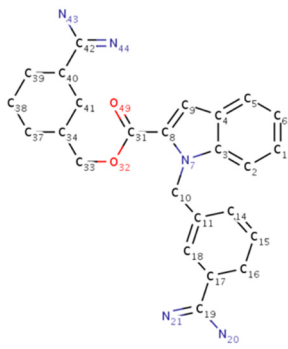
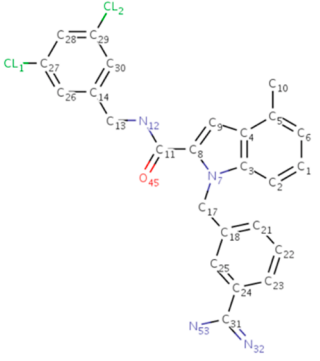
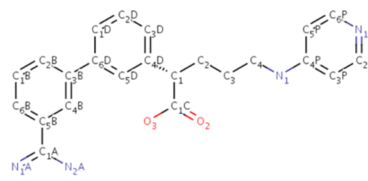
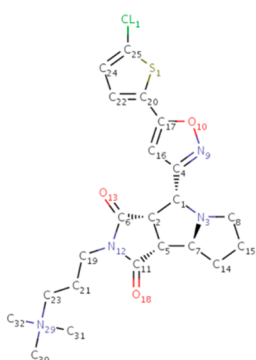
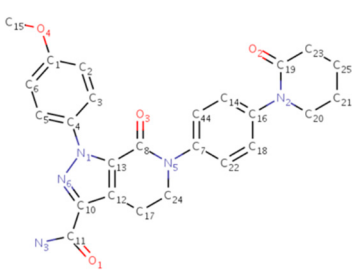
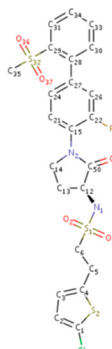
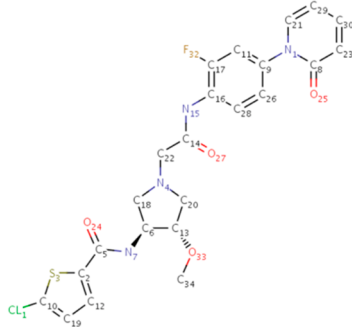
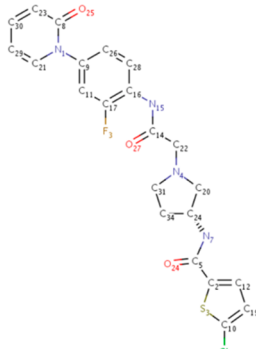
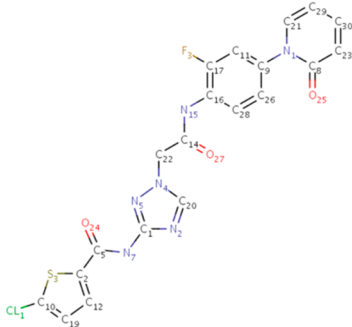
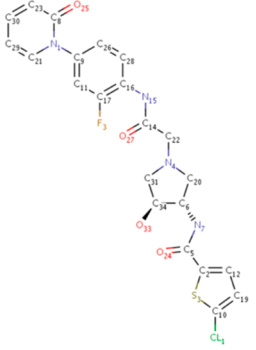
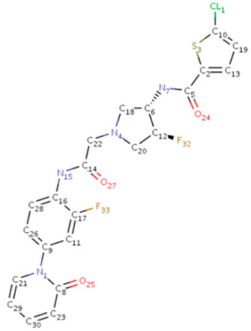
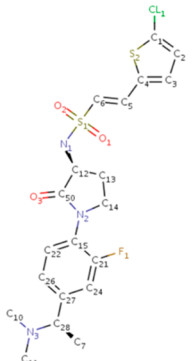
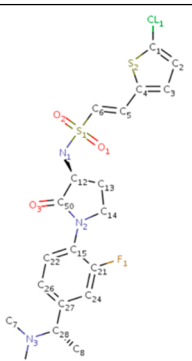
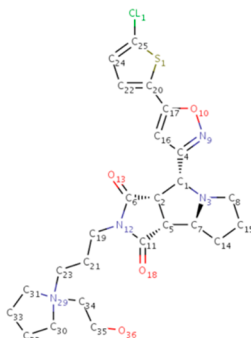
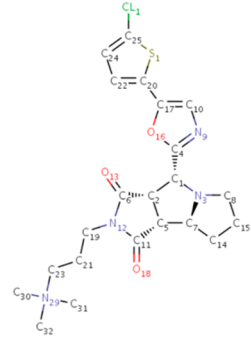
25	2J34/GS6		6-Chloro- <i>N</i> -((3 <i>S</i>)-1-[(1 <i>S</i>)-1-methyl-2-morpholin-4-yl-2-oxoethyl]-2-oxopyrrolidin-3-yl)-1-benzothiophene-2-sulfonamide	472.0	−10.67 [14]
26	2J38/GS5		5-Chloro- <i>N</i> -((3 <i>S</i>)-1-[(1 <i>S</i>)-1-methyl-2-morpholin-4-yl-2-oxoethyl]-2-oxopyrrolidin-3-yl)-1-benzothiophene-2-sulfonamide	472.0	−9.99 [14]
27	2J4I/GSJ		1-Pyrrolidineacetamide, 3-[[[(6-chloro-2-naphthalenyl)sulfonyl]amino]-alpha-methyl-n-(1-methylethyl)- <i>N</i> -[2-[(methylsulfonyl)amino]ethyl]-2-oxo-, (alpha <i>S</i> ,3 <i>S</i>)-	559.1	−12.27 [12]
28	3LIW/RUP		(<i>r</i>)-2-(3-Adamantan-1-yl-ureido)-3-(3-carbamimidoyl-phenyl)- <i>N</i> -phenethyl-propionamide	487.6	−10.37 [15]

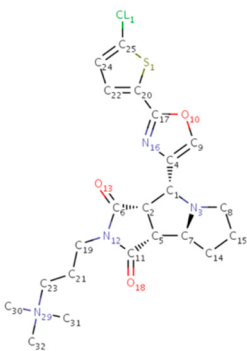
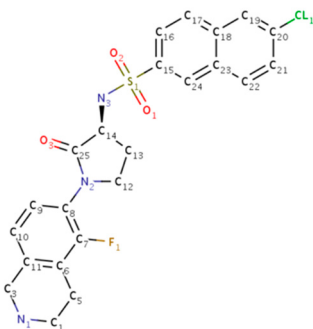
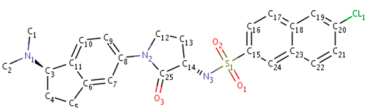
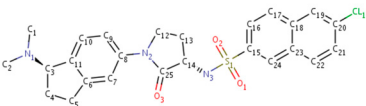
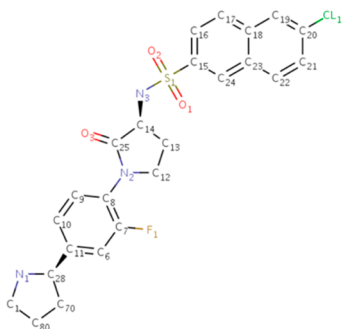
Table S2. 23 ligands of coagulation factor Xa for the test set.

ID	PDB-ID/ res-ID	Chemical Structure	Compound Name	MW [g/mol]	ΔG_{exp} [kcal/mol] (Supplementary Reference)
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1	1LPK/CBB		1-(3-Carbamimidoylbenzyl)-1H-indole-2-carboxylic acid 3-carbamimidoylbenzylester	433.6	-10.20 [6]
2	1LPZ/CMB		1-(3-Carbamimidoylbenzyl)-N-(3,5-dichlorobenzyl)-4-methyl-1H-indole-2-carboxamide	465.4	-10.26 [6]
3	1XKA/4PP		(2s)-(3'-Amidino-3-biphenyl)-5-(4-pyridylamino)pentanoic acid	388.5	-9.29 [16]
4	2JKH/BI7		3-[(3as,4r,8as,8br)-4-[5-(5-chloro-2-thienyl)isoxazol-3-yl]-1,3-dioxooctahydropyrrolo[3,4-a]pyrrolizin-2(3h)-yl]-N,N,N-trimethylpropan-1-aminium	464.0	-10.86 [17]
5	2P16/GG2		1-(4-Methoxyphenyl)-7-oxo-6-[4-(2-oxopiperidin-1-yl)phenyl]-4,5,6,7-tetrahydro-1h-pyrazolo[3,4-c]pyridine-3-carboxamide	459.5	-13.63 [18]

6	2VH6/GSV		2-(5-Chlorothiophen-2-yl)-N-((3s)-1-[3-fluoro-2'-(methylsulfonyl)biphenyl-4-yl]-2-oxopyrrolidin-3-yl)ethanesulfonamide	557.1	-13.09 [19]
7	2VVC/LZF		5-Chloro-N-[(3s,4s)-1-(2-{[2-fluoro-4-(2-oxopyridin-1(2H)-yl)phenyl]amino}-2-oxoethyl)-4-methoxypyrrrolidin-3-yl]thiophene-2-carboxamide	505.0	-11.51 [20]
8	2VVU/H22		5-Chloro-N-[(3r)-1-(2-{[2-fluoro-4-(2-oxopyridin-1(2H)-yl)phenyl]amino}-2-oxoethyl)pyrrolidin-3-yl]thiophene-2-carboxamide	474.9	-10.93 [20]
9	2VVV/H21		5-Chloro-N-[1-(2-{[2-fluoro-4-(2-oxopyridin-1(2H)-yl)phenyl]amino}-2-oxoethyl)-1H-1,2,4-triazol-3-yl]thiophene-2-carboxamide	472.9	-11.10 [20]
10	2VWN/H25		5-Chloro-thiophene-2-carboxylic acid ((3s,4s)-1-[[2-fluoro-4-(2-oxo-2H-pyridin-1-yl)-phenyl]carbonyl]-methyl)-4-hydroxypyrrrolidin-3-yl)-amide	490.9	-10.80 [20]

11	2VWO/LZG		5-Chloro-thiophene-2-carboxylic acid ((3s,4s)-4-fluoro-1-([2-fluoro-4-(2-oxo-2h-pyridin-1-yl)-phenylcarbamoyl]-methyl)-pyrrolidin-3-yl)-amide	492.9	-10.14 [20]
12	2WYG/461		(e)-2-(5-Chlorothiophen-2-yl)-N-[(3s)-1-[4-[(1r)-1-(dimethylamino)ethyl]-2-fluorophenyl]-2-oxopyrrolidin-3-yl]ethanesulfonamide	472.0	-11.74 [21]
13	2WYJ/898		(e)-2-(5-Chlorothiophen-2-yl)-N-[(3s)-1-[4-[(1s)-1-(dimethylamino)ethyl]-2-fluorophenyl]-2-oxopyrrolidin-3-yl]ethanesulfonamide	472.0	-12.15 [21]
14	2Y5F/XWG		(3as,4r,5s,8as,8br)-4-[5-(5-Chlorothiophen-2-yl)-1,2-oxazol-3-yl]-2-[3-[1-(2-hydroxyethyl)pyrrolidin-1-ium-1-yl]propyl]-4,6,7,8,8a,8b-hexahydro-3ah-pyrrolo[3,4-a]pyrrolizine-1,3-dione	520.1	-11.74 [22]
15	2Y5G/FJD		3-[(3as,4r,5s,8as,8br)-4-[5-(5-Chlorothiophen-2-yl)-1,3-oxazol-2-yl]-1,3-dioxo-4,6,7,8,8a,8b-hexahydro-3ah-pyrrolo[3,4-a]pyrrolizin-2-yl]propyl-trimethyl-azanium	464.0	-9.23 [22]

16	2Y5H/Y5H		3-[(3as,4r,5s,8as,8br)-4-[2-(5-Chlorothiophen-2-yl)-1,3-oxazol-4-yl]-1,3-dioxo-4,6,7,8,8a,8b-hexahydro-3ah-pyrrolo[3,4-a]pyrrolizin-2-yl]propyl-trimethyl-azanium	464.0	-7.82 [22]
17	2Y7X/MZA		6-Chloro-N-[(3s)-1-(5-fluoro-1,2,3,4-tetrahydroisoquinolin-6-yl)-2-oxo-pyrrolidin-3-yl]naphthalene-2-sulfonamide	474.0	-12.00 [23]
18	2Y7Z/C0Z		6-Chloro-N-[(3s)-1-[(1s)-1-dimethylamino-2,3-dihydro-1h-inden-5-yl]-2-oxo-pyrrolidin-3-yl]naphthalene-2-sulfonamide	484.0	-11.74 [24]
19	2Y80/439		6-Chloro-N-[(3s)-1-[(1s)-1-(dimethylamino)-2,3-dihydro-1H-inden-5-yl]-2-oxo-3-pyrrolidinyl]-2-naphthalenesulfonamide	484.0	-10.86 [24]
20	2Y81/931		6-Chloro-N-[(3s)-2-oxo-1-[4-[(2r)-2--pyrrolidinyl] phenyl]-3-pyrrolidinyl]-2-naphthalenesulfonamide	488.0	-11.74 [24]

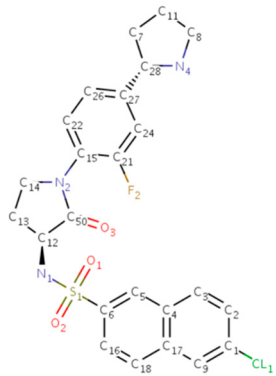
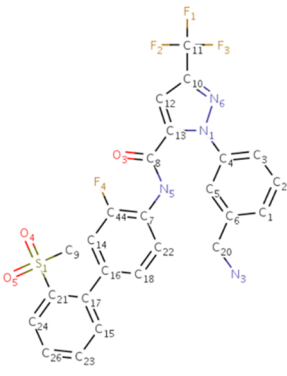
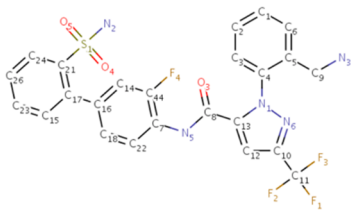
21	2Y82/930		6-Chloro- <i>N</i> -((3 <i>s</i>)-2-oxo-1-[4-((2 <i>s</i>)-2-pyrrolidinyl)phenyl]-3-pyrrolidinyl)-2-naphthalenesulfonamide	488.0	−11.34 [24]
22	3M36/M35		1-[3-(Aminomethyl)phenyl]- <i>N</i> -[3-fluoro-2'-(methylsulfonyl)biphenyl-4-yl]-3-(trifluoromethyl)-1 <i>H</i> -pyrazole-5-carboxamide	532.5	−13.26 [8]
23	3M37/M37		1-[2-(Aminomethyl)phenyl]- <i>N</i> -(3-fluoro-2'-sulfamoylbiphenyl-4-yl)-3-(trifluoromethyl)-1 <i>H</i> -pyrazole-5-carboxamide	533.5	−12.21 [25]

Table S3. Experimentally measured binding free energies and calculated scores for the training set ¹.

PDB-ID	ΔG_{exp} [kcal/mol]	AutoDock4 Score	AutoDock-GIST Score ² (Affinity Parameter Set)	AutoDock-GIST Score ² (Pose Parameter Set)
1EZQ	-12.21	-10.50	-27.50	-18.50
1F0R	-10.34	-9.59	-22.09	-15.84
1F0S	-10.45	-8.46	-21.46	-15.46
1FJS	-13.44	-10.51	-28.01	-18.51
1G2L	-10.28	-9.82	-25.32	-17.57
1G2M	-10.49	-10.49	-25.99	-18.24
1KSN	-12.82	-11.00	-27.50	-18.50
1LQD	-10.97	-9.54	-24.54	-17.04
1MQ5	-12.31	-11.80	-27.80	-19.05
1MQ6	-15.06	-11.10	-27.60	-18.35
1NFU	-10.45	-9.69	-23.69	-16.19
1NFW	-12.09	-9.86	-23.86	-16.36
1NFX	-11.51	-10.33	-25.83	-17.08
1NFY	-12.00	-9.89	-24.89	-16.14
1Z6E	-13.12	-11.57	-30.07	-20.57
2BMG	-10.56	-11.67	-26.67	-19.17
2BOH	-11.62	-10.01	-26.01	-18.01
2BOK	-9.173	-7.59	-21.09	-14.34
2BQ7	-9.61	-9.12	-23.62	-16.62
2BQW	-11.62	-10.88	-26.38	-18.38
2CJI	-11.21	-10.89	-25.89	-18.39
2FZZ	-14.21	-13.57	-32.07	-22.32
2G00	-13.16	-13.52	-32.02	-22.52
2J2U	-9.61	-10.76	-24.76	-17.51
2J34	-10.67	-11.05	-25.05	-17.80
2J38	-9.99	-10.78	-24.78	-17.78
2J4I	-12.27	-9.64	-26.14	-17.14
3LIW	-10.37	-8.53	-24.03	-15.28

¹ Crystal structures of 28 binding ligands for coagulation factor Xa. ² The absolute values of AutoDock-GIST scores are not scaled to those of experimentally measured binding free energies.

Table S4. Experimentally measured binding free energies and calculated scores for the test set ¹.

PDB-ID	ΔG_{exp} [kcal/mol]	AutoDock4 Score	AutoDock-GIST Score ² (Affinity Parameter Set)	AutoDock-GIST Score ² (Pose Parameter Set)
1LPK	-10.20	-8.30	-23.80	-16.30
1LPZ	-10.26	-9.71	-25.21	-17.21
1XKA	-9.29	-7.62	-20.62	-14.12
2JKH	-10.86	-8.70	-23.70	-15.70
2P16	-13.63	-11.87	-27.87	-19.87
2VH6	-13.09	-11.24	-27.74	-19.24
2VVC	-11.51	-10.73	-25.23	-18.48
2VVU	-10.93	-11.56	-26.06	-18.81
2VVV	-11.10	-10.72	-25.22	-18.22
2VWN	-10.80	-10.81	-25.31	-18.06
2VWO	-10.14	-10.89	-25.89	-18.39
2WYG	-11.74	-9.27	-23.27	-16.27
2WYJ	-12.15	-9.89	-24.39	-16.89
2Y5F	-11.74	-7.60	-23.60	-14.85
2Y5G	-9.23	-7.72	-23.22	-14.97
2Y5H	-7.82	-6.31	-20.81	-13.56
2Y7X	-12.00	-8.51	-24.01	-15.76
2Y7Z	-11.74	-10.75	-26.75	-18.50
2Y80	-10.86	-9.79	-25.29	-17.54
2Y81	-11.74	-11.84	-26.84	-19.34
2Y82	-11.34	-11.59	-27.09	-19.34
3M36	-13.26	-13.06	-30.06	-21.06
3M37	-12.21	-11.58	-28.08	-19.58

¹ Crystal structures of 23 binding ligands for coagulation factor Xa. ² The absolute values of AutoDock-GIST scores are not scaled to those of experimentally measured binding free energies.

Table S5. RMSDs between crystal structures and docking poses for the training set.

PDB-ID	AutoDock4 [Å]	AutoDock-GIST (Affinity Parameter Set) [Å]	AutoDock-GIST (Pose Parameter Set) [Å]
1EZQ	1.47	1.78	1.15
1F0R	1.60	1.67	1.76
1F0S	1.71	1.66	1.83
1FJS	0.89	0.68	1.11
1G2L	1.05	1.06	1.17
1G2M	0.91	0.90	0.92
1KSN	1.64	0.87	0.70
1LQD	2.61	3.15	2.47
1MQ5	0.86	2.25	0.78
1MQ6	3.77	3.28	1.18
1NFU	5.71	5.50	0.77
1NFW	0.75	1.65	0.92
1NFX	1.24	6.18	1.42
1NFY	0.68	0.94	0.77
1Z6E	0.57	0.67	0.55
2BMG	3.70	1.24	3.02
2BOH	1.02	0.62	0.55
2BOK	0.58	1.34	0.51
2BQ7	2.90	0.52	1.17
2BQW	1.78	3.49	4.52
2CJI	1.09	2.78	1.03
2FZZ	1.47	0.55	1.25
2G00	0.63	2.89	0.31
2J2U	1.54	1.58	0.22
2J34	0.45	1.07	0.25
2J38	0.78	1.39	0.30
2J4I	3.72	0.36	1.85
3LIW	2.18	0.24	1.19

Table S6. RMSDs between crystal structures and docking poses for the test set.

PDB-ID	AutoDock4 [Å]	AutoDock-GIST (Affinity Parameter Set) [Å]	AutoDock-GIST (Pose Parameter Set) [Å]
1LPK	4.43	0.19	2.87
1LPZ	3.71	3.72	1.28
1XKA	3.06	0.66	0.96
2JXH	0.89	0.26	0.44
2P16	0.24	0.36	0.24
2VH6	1.81	1.38	0.90
2VVC	1.18	1.13	0.48
2VVU	1.02	1.03	0.39
2VVV	0.78	1.04	0.80
2VWN	1.33	0.97	0.39
2VWO	1.72	0.23	0.43
2WYG	1.33	1.22	1.55
2WYJ	1.05	0.95	1.18
2Y5F	1.22	0.62	1.14
2Y5G	0.61	2.73	0.55
2Y5H	2.75	1.11	0.84
2Y7X	1.20	0.30	1.15
2Y7Z	0.64	0.90	0.25
2Y80	0.71	0.39	0.65
2Y81	0.33	0.33	0.13
2Y82	0.43	1.53	0.31
3M36	0.77	0.58	0.61
3M37	1.86	1.79	1.90

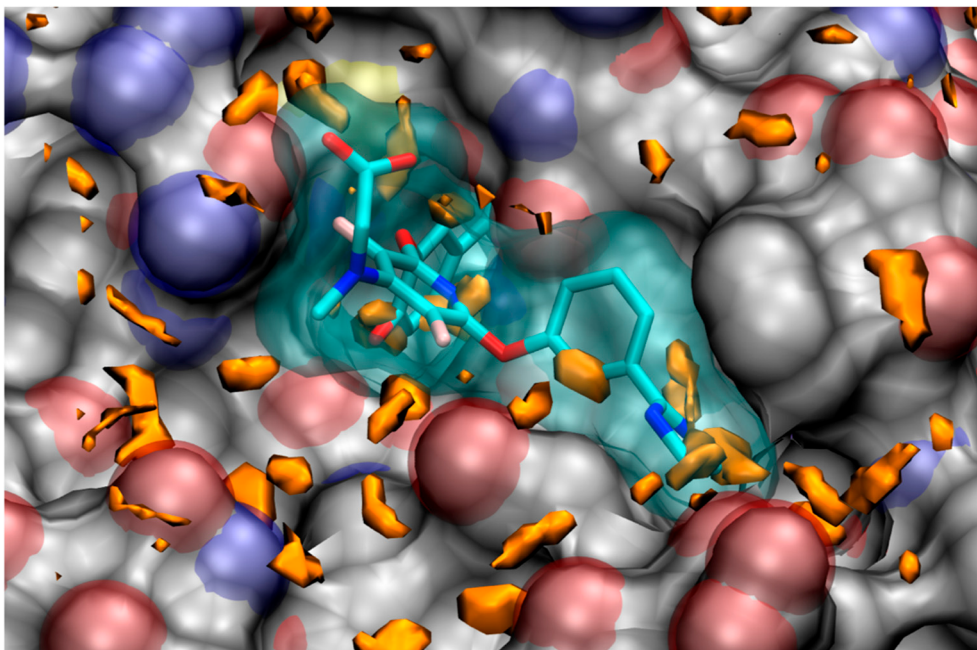


Figure S1. The structure of 1FJS ligand (cyan stick and transparent surface) bound to the active site of FXa (gray). Oxygen, nitrogen, sulfur and fluorine are represented in red, blue yellow and pink, respectively. The unfavorable water distribution of affinity parameter set is shown in orange.

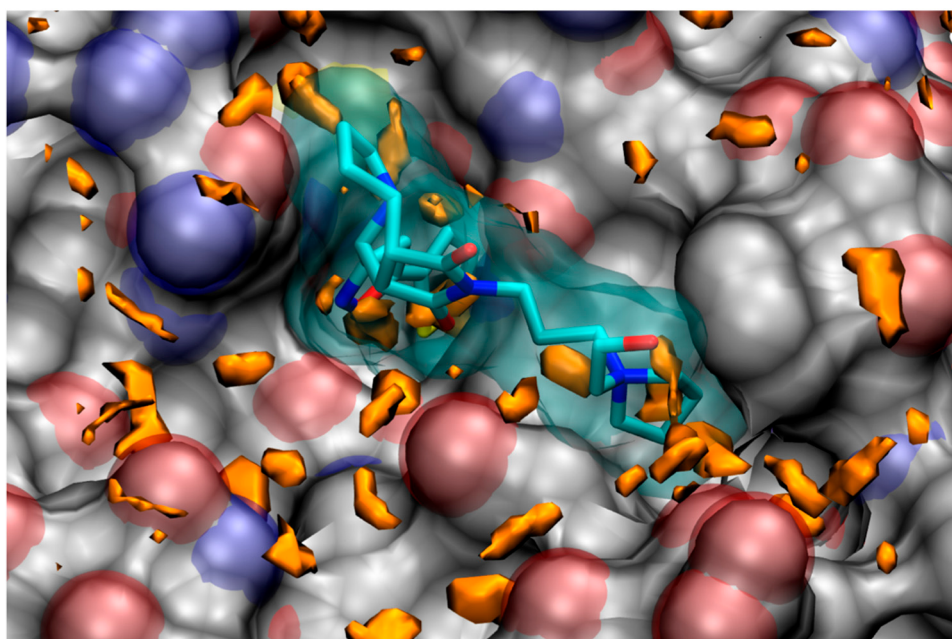


Figure S2. The structure of 2Y5F ligand (cyan stick and transparent surface) bound to the active site of FXa (gray). Oxygen, nitrogen and sulfur are represented in red, blue and yellow, respectively. The unfavorable water distribution of affinity parameter set is shown in orange.

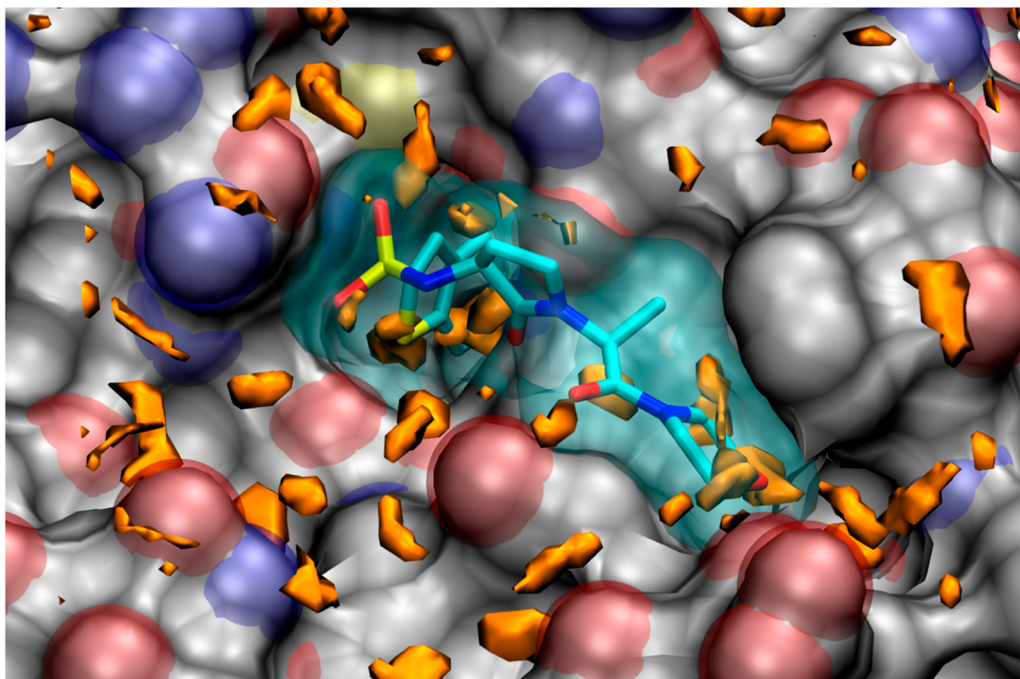


Figure S3. The structure of 2J34 ligand (cyan stick and transparent surface) bound to the active site of FXa (gray). Oxygen, nitrogen and sulfur are represented in red, blue and yellow, respectively. The unfavorable water distribution of affinity parameter set is shown in orange.

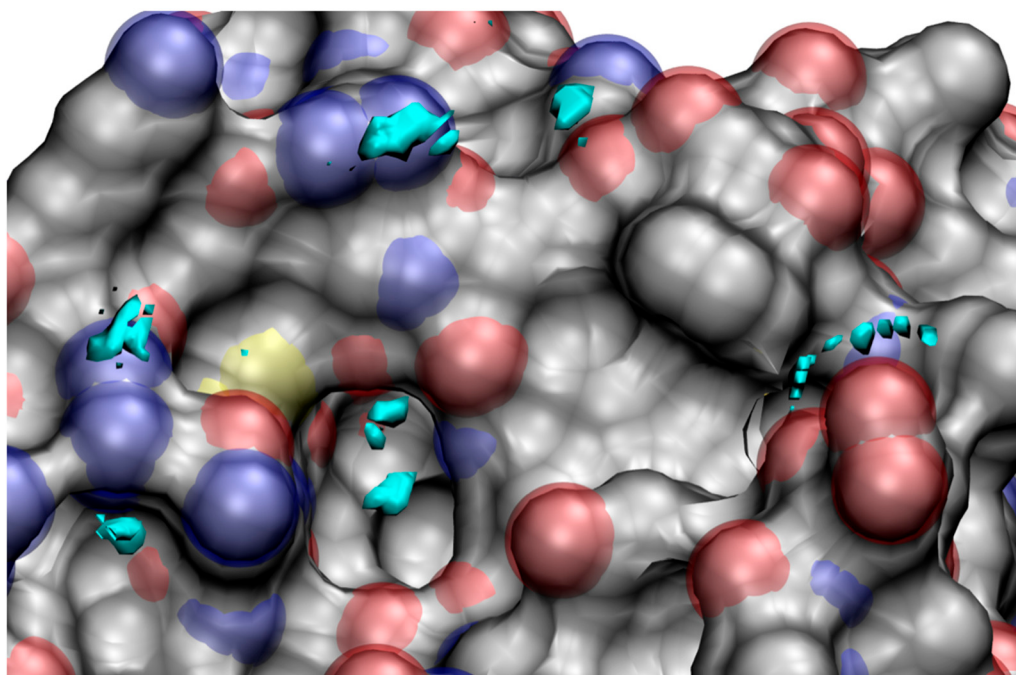


Figure S4. The high-occupancy and favorable water distribution (cyan) in FXa active site (gray) calculated by GIST. Oxygen, nitrogen and sulfur are represented in red, blue and yellow, respectively.

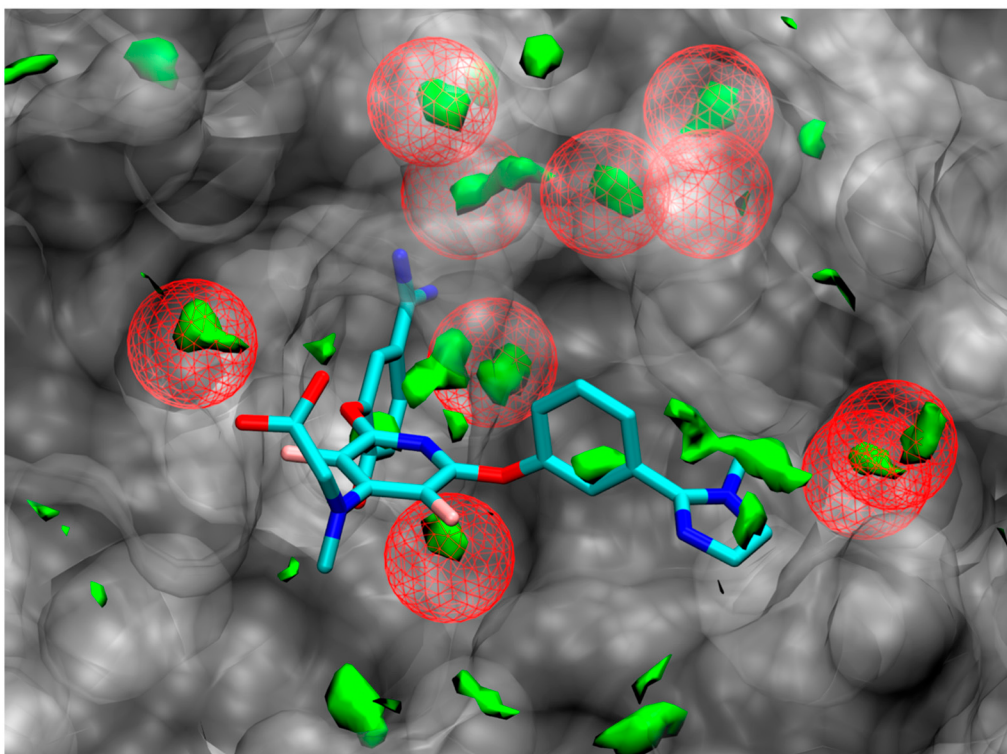


Figure S5. Co-crystal structure of water molecules (red sphere) in the active site of FXa (transparent surface) with a ligand (cyan stick) of PDB entry 1FJS. Green surface represents high-occupancy water region calculated by GIST.

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