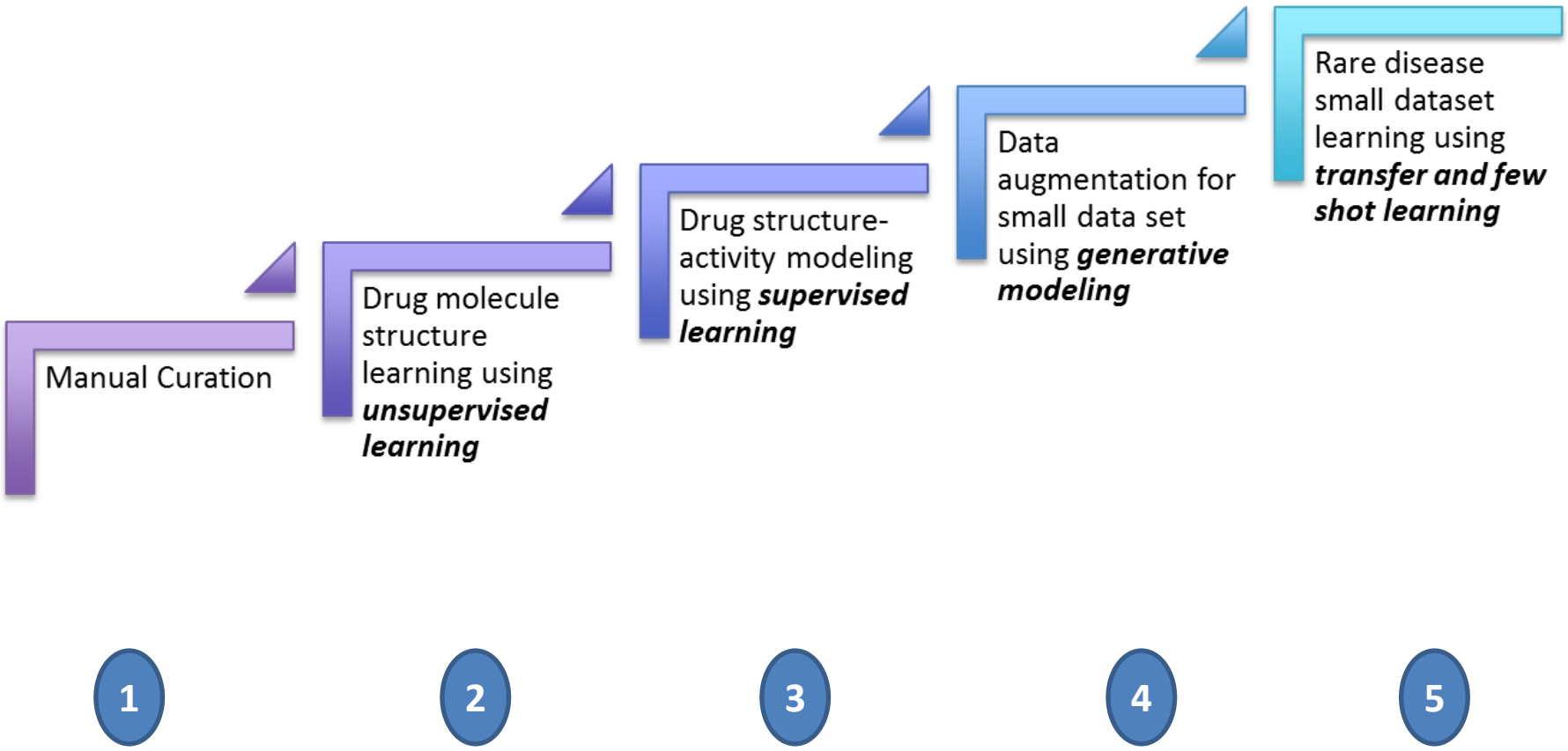
The Technology

Various Components and Subsystems

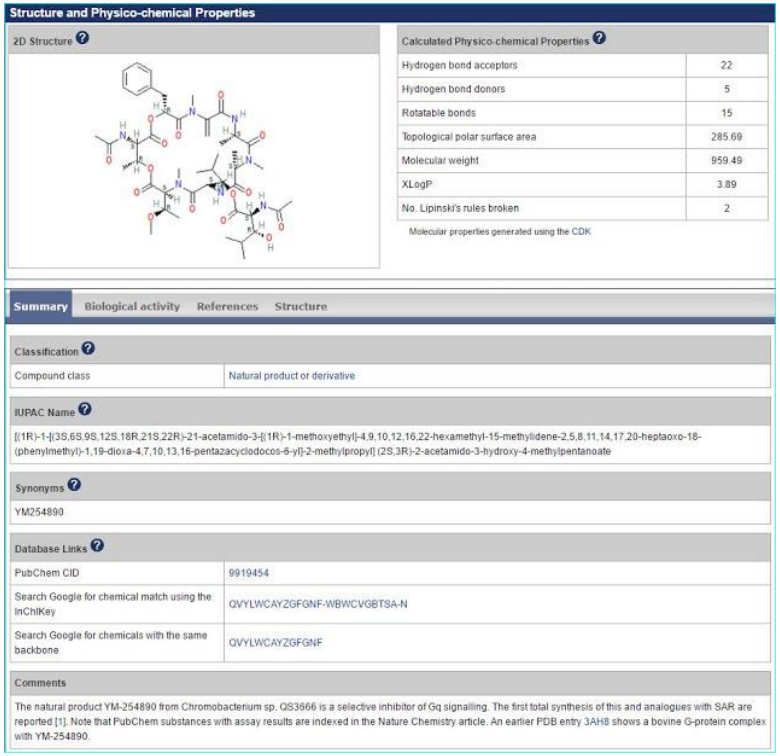


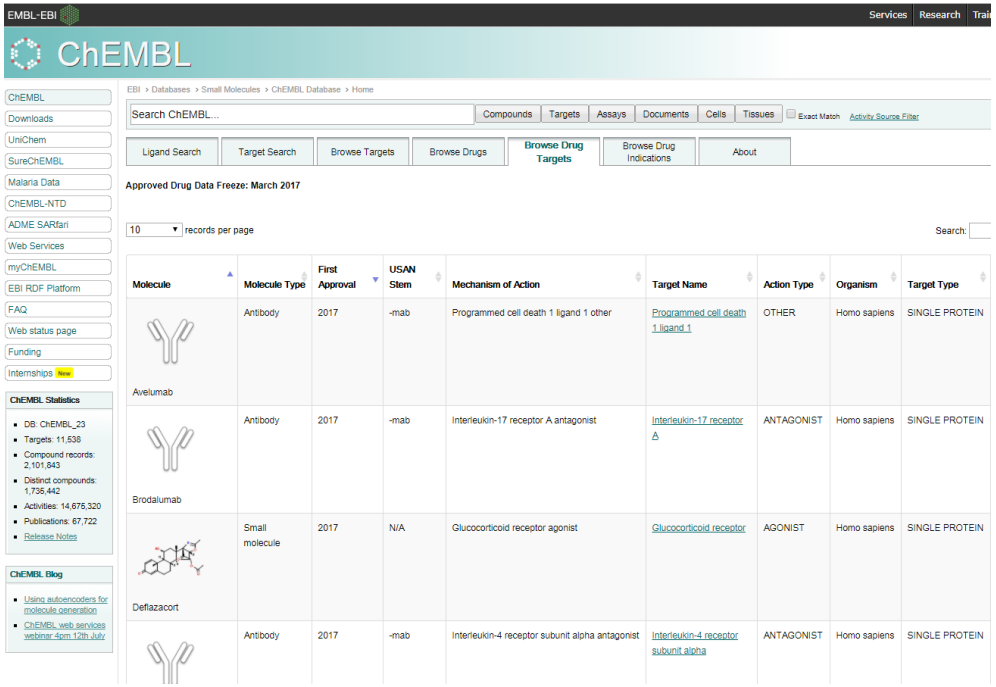
Manual Curation

Extracting biochemical information from unstructured literature (patents/journal article) into structured format (by semi-automation). ChEMBL open source database will be helpful for extracting Protease inhibitors .

Manual Curation

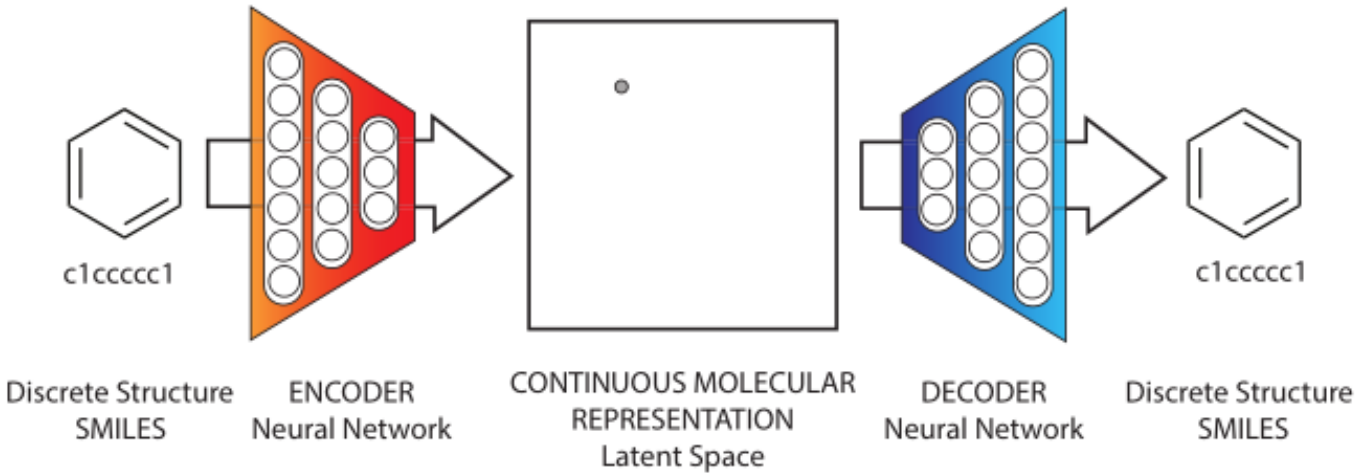
ChEMBL database





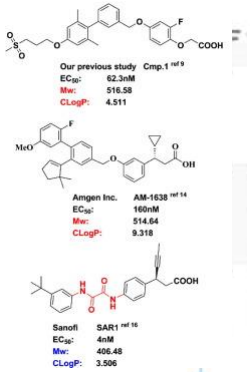
Drug Molecule Structure Learning:

From the curated database, we extract the molecules and encode chemical structures as features. This is how we inject the drug molecules knowledge into Neural Networks



Drug Structure Activity Modelling using Supervised Learning

SAR data
+ Molecular descriptors



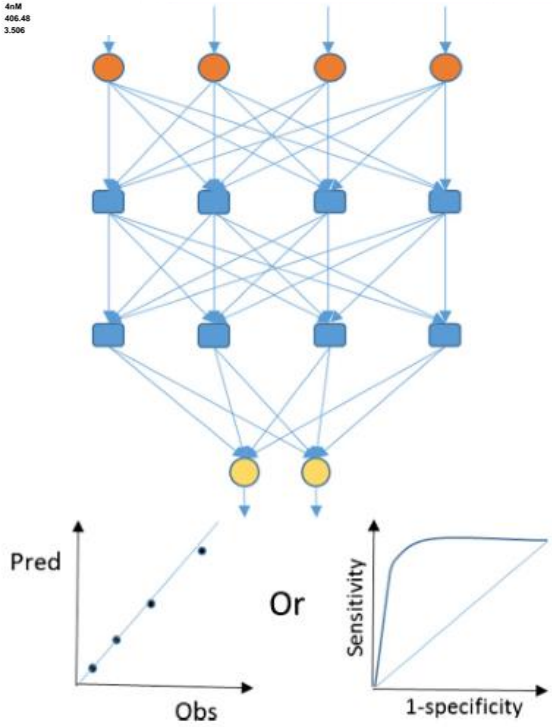
Title	ALogP	ALogp2	AMR	BCUTp-1	BCUTp-1h	fragC	apol	naAromAt	nAtom
Acebutolol	-2.1004	4.41168	67.0727	4.75506	9.76616	2152.06	55.7582	6	52
Amoxicillin	-1.874	3.51188	63.5964	4.78119	12.1043	1516.09	51.0391	6	44
Bendroflumethlazide	-0.8075	0.65206	39.6471	4.24449	12.8541	1147.12	49.7141	12	41
Benzocaine	-0.4883	0.23844	20.5043	4.54796	8.52863	397.03	25.8787	6	23
Benzthlazide	-1.3398	1.79506	50.9651	6.69282	12.9721	1114.11	53.1231	10	40
Clozapine	0.0092	8.46E-05	45.0577	5.92358	11.5903	1519.05	50.9291	12	42
Dibucaine	-0.5602	0.31382	57.7981	5.09923	10.5046	2425.05	59.441	10	54
Diethylstibestrol	-0.2302	0.05299	33.7086	4.85094	11.5338	1301.02	46.6199	12	40
Diflunisal	0.2593	0.06724	11.5289	4.03982	9.84339	423.05	31.7343	12	26
Dipyridamole	-3.3772	11.4055	95.8664	4.30728	11.5724	4981.12	80.9197	10	76
Folic_Acid	-3.3182	11.0105	52.3395	3.83215	9.33155	1817.13	58.6211	12	51
Furosemide	-0.9949	0.98983	33.6078	4.6696	12.2524	669.09	39.7447	11	32
Hydrochlorothiazide	-1.5436	2.3827	35.4327	7.26833	12.8018	404.1	32.1423	6	25
Imipramine	0.091	0.00828	39.2562	5.70631	11.5894	1789.02	51.643	12	45
Indomethacin	0.2922	0.08538	37.8739	4.12177	11.086	1249.06	50.5967	12	41
Ketoprofen	-0.2088	0.0436	21.9194	4.29006	9.91827	814.03	39.9011	12	33
Lidocaine	1.166	1.35956	45.8867	5.09304	9.98654	1249.03	42.3114	6	39
Meclofenamic_acid	1.4704	2.16208	27.2526	4.4468	11.3886	619.05	39.0387	12	30

Input Layer

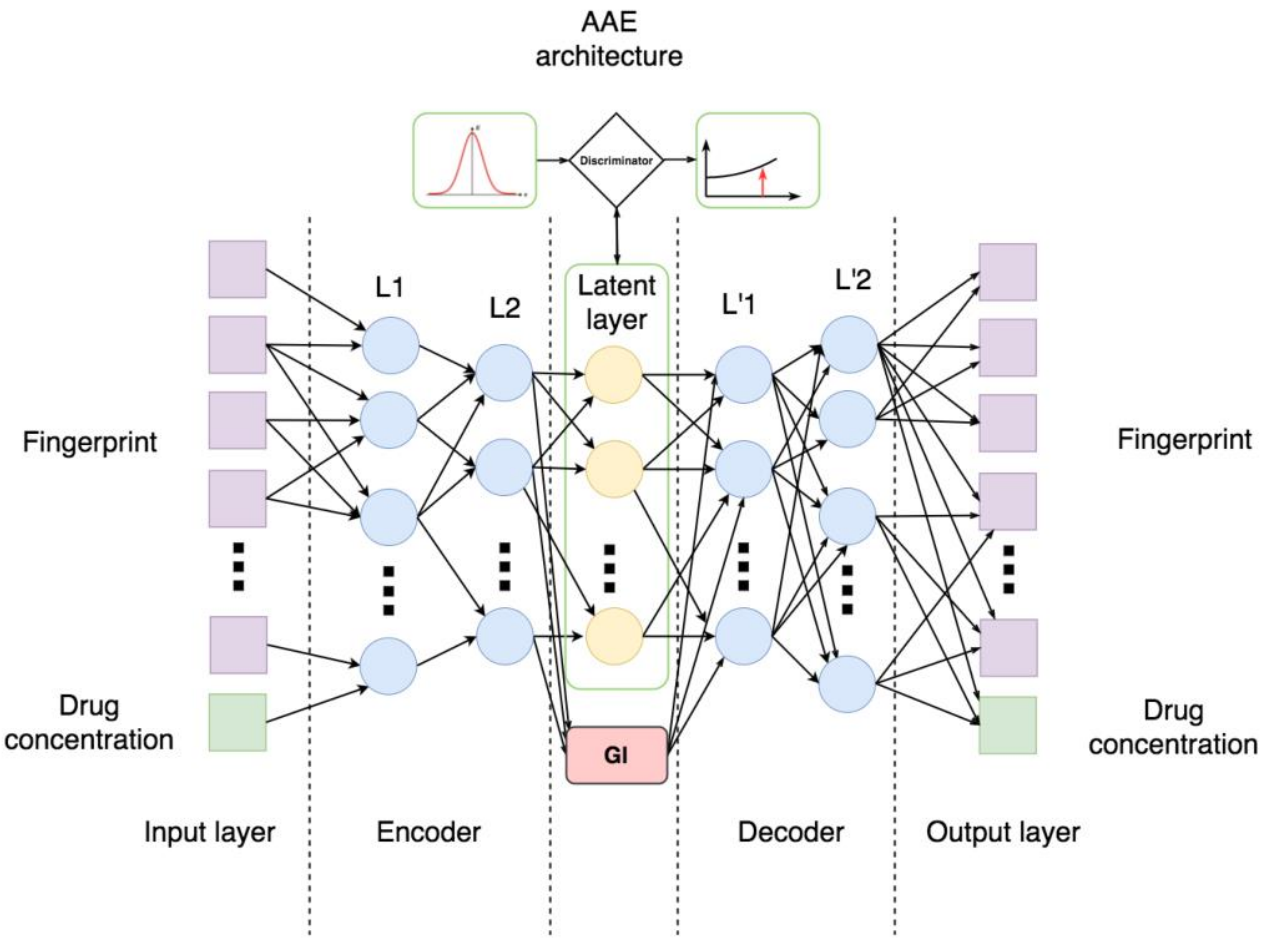
Hidden Layers

Output layer

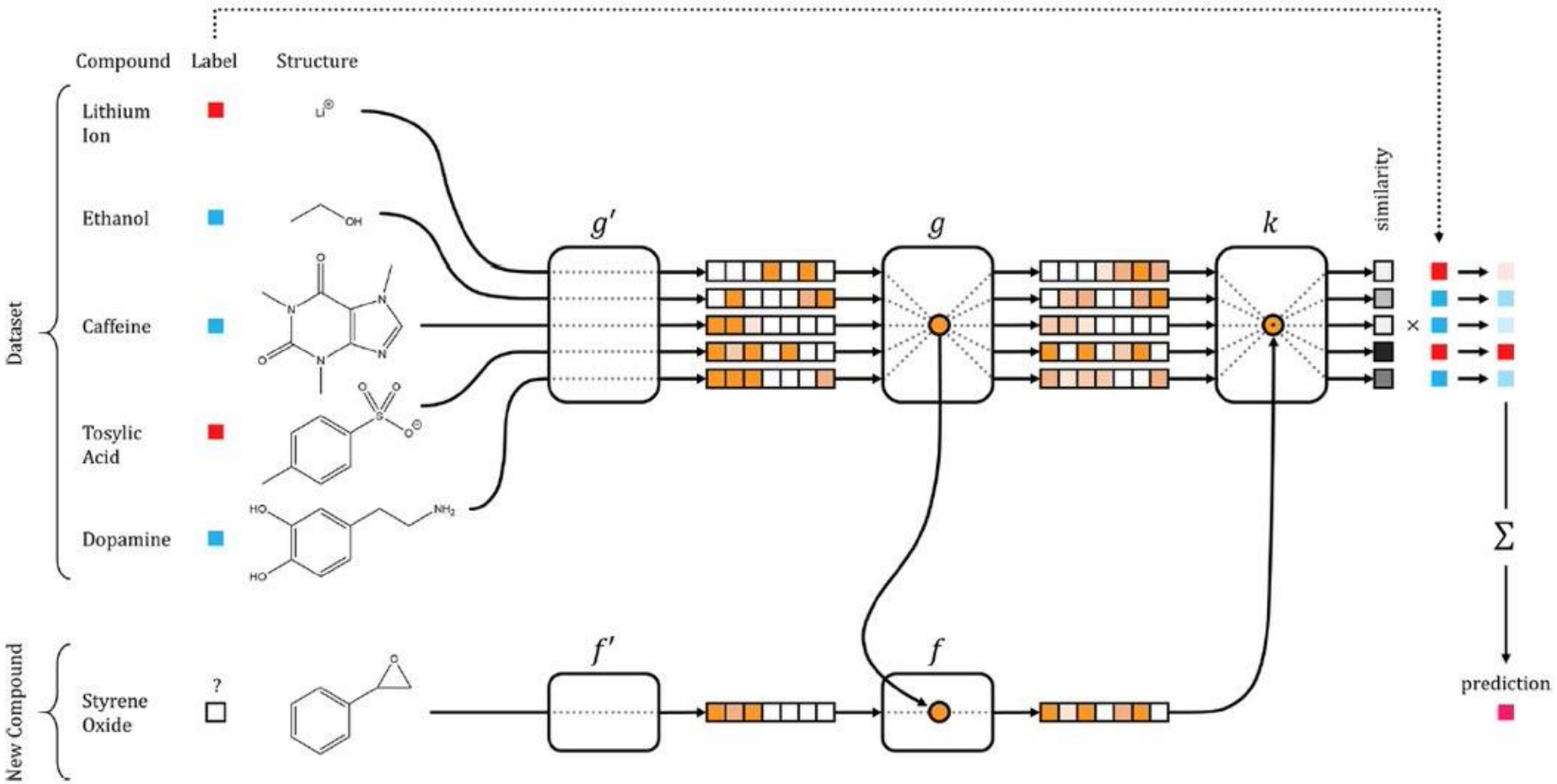
Correlation



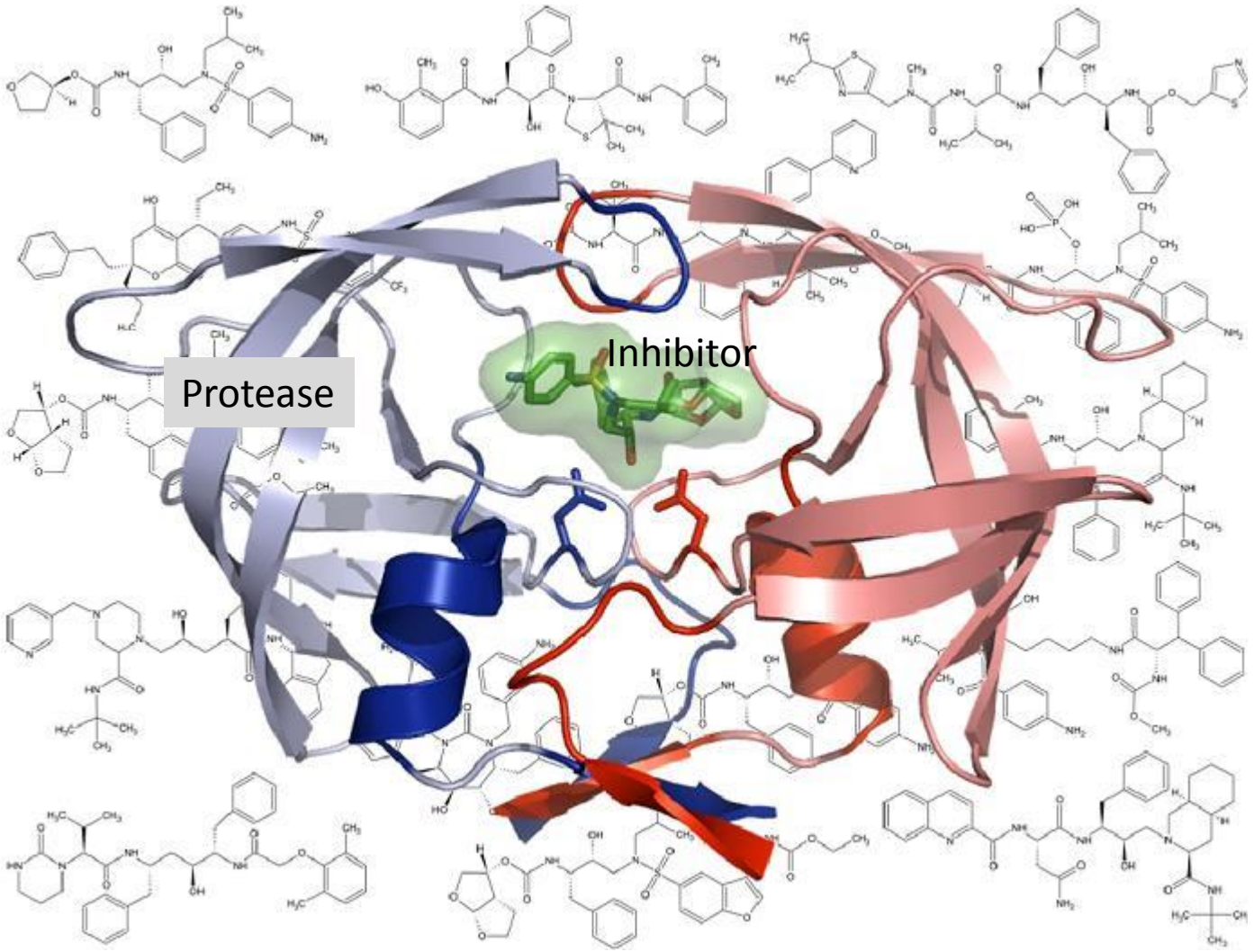
Data Augumentation using Generative Adversarial Networks



Network Architecture for One-Shot Learning in Drug Discovery



HIV Protease Inhibitors are Powerful anti-AIDS drugs



Quality Inhibitors for Quality Results

VINGYANI