

SUPPORTING INFORMATION

A Chemical Space Odyssey of Inhibitors of Histone Deacetylases and Bromodomains

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Table S1. Summary statistics of the distribution of the PCP for the data sets studied in this work

HBA								
	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	Inhibit	Epi-focused
Min	1	1	0	0	0	0	0	0
1st Quart	3	3	4	1	1	2	3	2
Median	4	4	5	1	2	3	4	4
Mean	3.662	4.813	5.357	1	2.919	3.585	4.459	3.416
3rd Quart	4	5	7	2	4	5	5	5
Max	7	37	24	22	18	18	19	8
SD	0.991	2.45	2.5	1.31	2.56	2.37	2.32	1.83

HBD								
	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General	Epi-focused
Min	0	0	0	0	0	0	0	0
1st Quart	1	2	2	0	0	0	1	1
Median	1	3	3	0	1	1	2	2
Mean	1.184	3.265	3.133	0.4666	1.35	1.525	2.298	2.298
3rd Quart	2	4	4	1	2	2	3	3
Max	4	24	12	14	13	13	16	16
SD	0.867	1.511	2.011	1.077	1.559	1.552	1.817	1.294

RTB								
	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General	Epi-focused
Min	1	0	0	0	0	0	0	1
1st Quart	3	6	4	2	2	4	3	4
Median	4	9	5	4	4	6	5	5
Mean	4.734	9.32	5.568	4.291	5.008	6.303	5.433	5.549
3rd Quart	6	11	7	6	7	8	7	7
Max	16	53	27	21	35	35	30	16
SD	2.05	4.03	2.931	3.345	3.779	4.186	3.342	2.84

SLogP								
	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General	Epi-focused
Min	0.183	-14.261	-3.3655	-5.422	-13.6094	-14.04	-8.113	-2.796
1st Quart	3.303	2.559	0.9547	1.519	0.1402	1.089	2.248	1.619
Median	4.161	3.613	2.9577	2.317	1.7112	2.499	3.545	2.588
Mean	4.059	3.594	2.6556	2.317	1.4113	2.274	3.49	2.627
3rd Quart	4.849	4.599	4.2702	3.051	3.0996	3.963	4.729	3.724
Max	6.701	9.65	10.685	8.312	10.4033	11.059	11.059	9.189
SD	1.172	1.69	2.394	1.331	2.91	2.772	1.92	1.715

Table S1. (Continued)

	TPSA							
	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General	Epi-focused
Min	22	26.3	0	0	0	0	0	21.51
1st Quart	67.56	78.43	78.11	18.46	43.37	60.11	61.32	73.25
Median	81.4	101.21	108.65	26.3	71.36	86.19	84.42	89.54
Mean	78.54	110.4	111.7	31.29	82.29	93.97	88.29	91.28
3rd Quart	90.63	129.98	143.87	37.3	105.15	115.15	107.29	105.82
Max	137.07	612.25	358.2	374.13	417.27	391.12	260.04	180.04
SD	19.54	42.897	47.102	24.611	55.74	52.316	40.715	27.983

	MW							
	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General	Epi-focused
Min	213.2	111.1	88.11	18.04	43.33	95.12	136.1	134.2
1st Quart	337.8	349.9	329.83	134.18	254.32	323.15	319.2	321.4
Median	374.4	413.5	403.41	160.25	325.41	407.45	408.4	380.4
Mean	378.7	441.6	402.8	169.9	348.4	418.47	411.6	392.5
3rd Quart	423.5	492.8	462.03	196.29	415.51	501.54	480.1	473.6
Max	552.7	1598.8	1326.78	967.02	988.18	975.64	990.2	720.9
SD	64.229	133.525	118.158	61.857	146.822	148.614	129.24	110.161

Table S2. Distribution of 'actives' compounds and comparison for BRDi compounds

Median	BRD2	BRD3	BRD4	IC ₅₀ (μM)
	4	4	4	1
	4	4	4	10
HBA	4	4	3	≥100
	1	1	1	1
	1	1	1	10
HBD	1	1	1	≥100
	4	4	4	1
	4	4	4	10
RTB	4	4	4	≥100
	4.71362	4.71792	4.54183	1
	4.01176	4.07626	4.2211	10
SLogP	3.782135	3.92028	4.02703	≥100
	81.4	81.4	81.4	1
	81.4	81.4	81.4	10
TPSA	81.4	81.4	78.51	≥100
	415.453	410.441	402.96	1
	376.864	378.493	376.391	10
MW	372.4445	376.864	364.96	≥100

Mean	BRD2	BRD3	BRD4	IC ₅₀ (μM)
	4	4	3.865854	1
	3.671233	3.821918	3.663317	10
HBA	3.684783	3.829268	3.463115	≥100
	1.1333	1.15625	1.268293	1
	1.041096	1.09589	1.201005	10
HBD	1.065217	1.109756	1.17623	≥100
	4.4	4.1875	4.304878	1
	4.260274	4.39726	4.251256	10
RTB	4.206522	4.353659	3.979508	≥100
	4.735897	4.529324	4.354318	1
	3.986589	3.989324	4.101408	10
SLogP	3.767619	3.820534	3.847727	≥100
	84.46933	83.95625	83.88902	1
	78.75466	82.3137	78.82246	10
TPSA	79.08337	82.40805	75.63689	≥100
	405.6323	398.6574	401.2273	1
	369.7882	383.3375	380.6428	10
MW	365.7007	379.908	361.3758	≥100

Table S2. (Continued)

SD	BRD2	BRD3	BRD4	IC ₅₀ (μM)
	0.755929	0.718421	1.015463	1
	0.834246	0.733001	0.990975	10
HBA	0.824479	0.699292	1.127049	≥100
	0.743223	0.677251	0.943448	1
	0.610973	0.581291	0.870272	10
HBD	0.625543	0.588073	0.878228	≥100
	0.736788	0.737804	1.95435	1
	1.302001	1.198839	1.704772	10
RTB	1.338673	1.260726	1.849143	≥100
	0.938424	0.915115	0.988773	1
	1.065457	1.118183	1.149915	10
SLogP	1.144782	1.198718	1.291483	≥100
	12.3812	10.72139	19.0008	1
	15.2	12.97325	19.49417	10
TPSA	15.8889	12.55546	20.81594	≥100
	34.6769	39.93311	60.62833	1
	62.87314	49.93638	63.05745	10
MW	63.4056	48.96492	78.06248	≥100

*Note: Statistical comparison towards “active” compounds of HDACi data set was not made because of the following:

Of the 2000 unique compounds; 1135 show IC₅₀ ≤ 1 μM.

The difference in size makes it difficult to make a good comparison on the same statistic grounds.

It is true that many types of structures have been proposed as HDACs inhibitors; however the most interesting and potent ones are inspired or derived from hydroxamic acid moiety.

That last fact has produced intensive research of those compounds; therefore a high chance of statistical similarities is plausible.

Finally the pharmacophoric features of such compounds are well understood as of today, making good reference for our probe search.

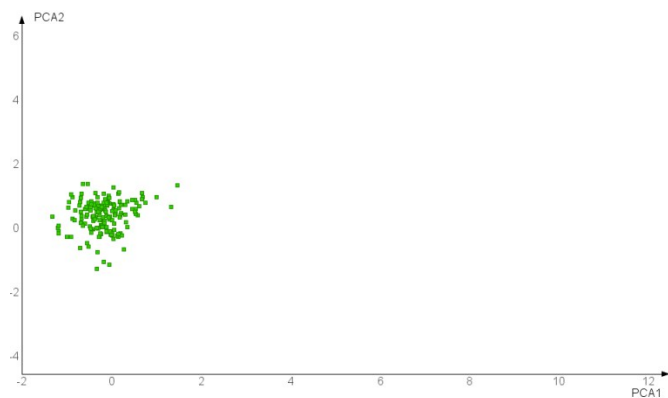
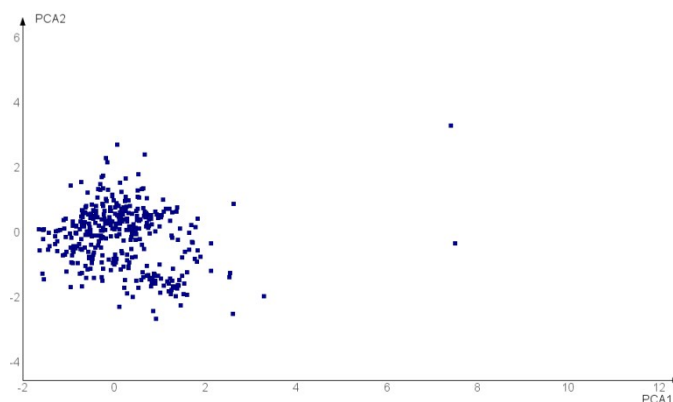
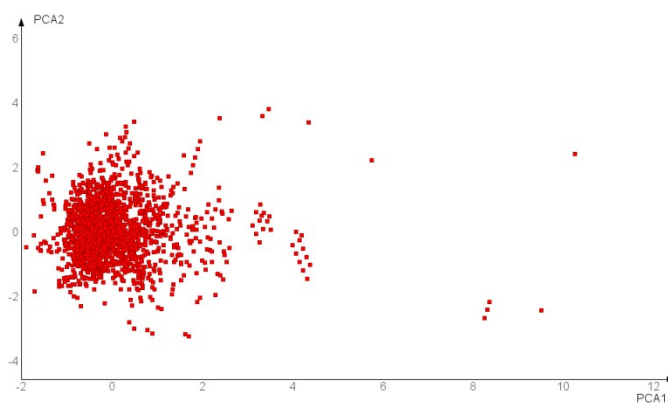
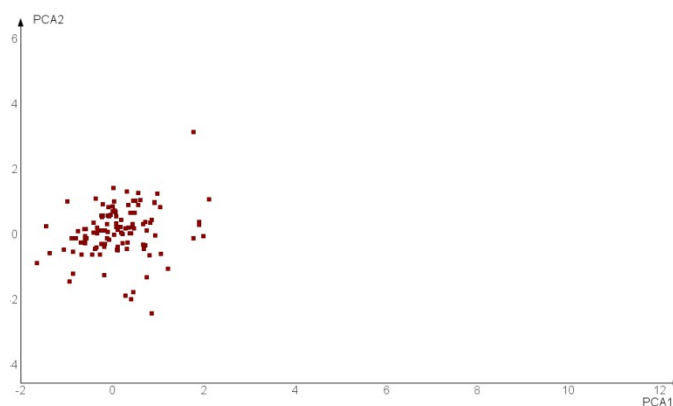
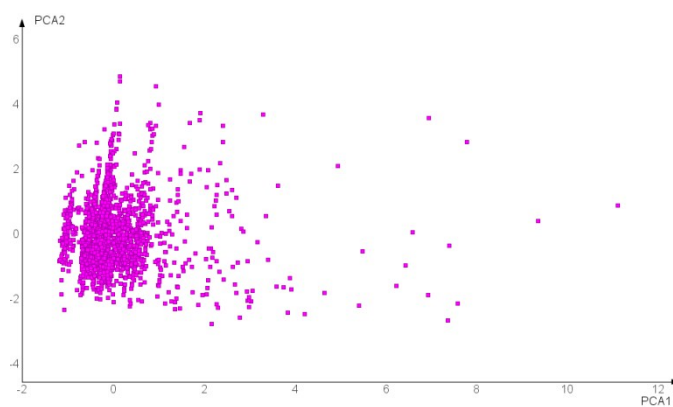
BRDi**DNMT 1****HDACi****'Epi-focused'****GRAS**

Figure S1. Chemical space representation of BRDi, HDACi, DNMT 1, 'Epi-focused', and GRAS. Data sets are in the same reference coordinates. The chemical space was obtained by principal component analysis of six physicochemical properties of pharmaceutical interest. The outliers in the HDACs and GRAS sets are not shown for clarity). See main text for details.

Table S3. Nemenyi test values for the PCPs comparisons of eight data sets (1 BRDi; 2 HDACi; 3 DNMT 1; 4 GRAS; 5 Drugs; 6 Clinic; 7 General; 8 Epi-focused).

SlogP

	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General
2	0.0005	-	-	-	-	-	-
3	9.80E-14	5.20E-14	-	-	-	-	-
4	< 2E-16	< 2E-16	1.10E-07	-	-	-	-
5	< 2E-16	< 2E-16	7.40E-14	< 2E-16	-	-	-
6	< 2E-16	< 2E-16	1.82E-01	< 2E-16	4.50E-05	-	-
7	1.00E-05	0.824	2.90E-13	< 2E-16	< 2E-16	< 2E-16	-
8	2.60E-12	7.50E-07	0.991	0.483	3.00E-04	0.999	0.382

TPSA

	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General
2	1.10E-09	-	-	-	-	-	-
3	6.90E-09	0.028	-	-	-	-	-
4	< 2E-16	< 2E-16	< 2E-16	-	-	-	-
5	0.0004	< 2E-16	6.50E-14	< 2E-16	-	-	-
6	0.62	0.6	0.999	< 2E-16	7.80E-14	-	-
7	0.6	0.025	0.999	< 2E-16	< 2E-16	1	-
8	0.999	0.303	0.999	< 2E-16	0.003	0.987	0.987

MW

	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General
2	0.006	-	-	-	-	-	-
3	0.879	0.028	-	-	-	-	-
4	< 2E-16	< 2E-16	< 2E-16	-	-	-	-
5	0.0004	< 2E-16	6.50E-14	< 2E-16	-	-	-
6	0.62	0.6	0.999	< 2E-16	7.80E-14	-	-
7	0.6	0.025	0.999	< 2E-16	< 2E-16	1	-
8	0.999	0.303	0.999	< 2E-16	0.003	0.987	0.987

HBA

	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General
2	0.001	-	-	-	-	-	-
3	1.10E-06	0.125	-	-	-	-	-
4	< 2E-16	< 2E-16	< 2E-16	-	-	-	-
5	2.50E-13	< 2E-16	< 2E-16	< 2E-16	-	-	-
6	0.0806	< 2E-16	< 2E-16	< 2E-16	9.10E-14	-	-
7	0.342	0.0211	2.00E-06	< 2E-16	< 2E-16	7.70E-14	-
8	0.598	3.40E-06	4.10E-09	6.40E-14	0.004	1	0.0028

HBD

	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General
2	< 2E-16	-	-	-	-	-	-
3	2.50E-14	0.002	-	-	-	-	-
4	< 2E-16	< 2E-16	< 2E-16	-	-	-	-
5	1	< 2E-16	< 2E-16	< 2E-16	-	-	-
6	0.792	< 2E-16	< 2E-16	< 2E-16	0.04	-	-
7	4.00E-12	< 2E-16	6.10E-10	< 2E-16	< 2E-16	6.00E-14	-
8	0.0002	8.80E-08	0.004	< 2E-16	1.00E-06	0.001	0.382

RTB

	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General
2	< 2E-16	-	-	-	-	-	-
3	0.027	< 2E-16	-	-	-	-	-
4	0.128	< 2E-16	9.50E-14	-	-	-	-
5	1	< 2E-16	2.80E-06	2.10E-07	-	-	-
6	0.0001	< 2E-16	0.694	< 2E-16	1.00E-13	-	-
7	0.367	< 2E-16	0.565	8.80E-14	0.0007	0.0009	-
8	0.39	7.10E-14	1	0.0001	0.164	0.968	0.989

Table S4. Loadings for the first three principal components of the property space of the eight compound data sets.

Principal component	PC1	PC2	PC3
Eigenvalue	1.9185	1.1174	0.7006
Cumulative eigenvalue (%)	61.344	82.157	90.338
SlogP	-0.0125	0.3832	-0.1515
TPSA	0.00904	-0.00414	-0.000591
MW	0.00284	0.00181	-0.000909
HBA	0.1949	-0.0369	-0.1355
HBD	0.2462	-0.0908	-0.1135
RTB	0.0895	0.0743	0.201

Table S5. Summary statistics of the intra-compound set similarity

MACCS					
Dataset	Q1	Mean	Median	Q3	SD
BRDi	0.386	0.48441626	0.463	0.557	0.14001451
HDACi	0.3445	0.4322109	0.4235	0.509	0.12798872
DNMT	0.309	0.41415293	0.4	0.5	0.16258865
GRAS	0.161	0.28927381	0.261	0.387	0.17045788
Drugs	0.22	0.31403743	0.308	0.4	0.1336488
Clinic	0.283	0.37388603	0.375	0.466	0.13127422
General	0.28	0.36957258	0.365	0.453	0.12375299
Epi-Focused	0.313	0.40095354	0.399	0.488	0.12786925

GpiDAPH3					
Dataset	Q1	Mean	Median	Q3	SD
BRDi	0.228	0.32056677	0.303	0.403	0.17481642
HDACi	0.34475	0.43225473	0.42375	0.50975	0.12792421
DNMT	0.161	0.24319609	0.243	0.324	0.18191181
GRAS	0	0.15952901	0	0.342	0.22637638
Drugs	0.2095	0.28546563	0.286	0.364	0.13806921
Clinic	0	0.19669936	0.223	0.3	0.13990382
General	0	0.18843588	0.218	0.303	0.14801609
Epi-Focused	0.194	0.25483202	0.275	0.344	0.13709412

TGD					
Dataset	Q1	Mean	Median	Q3	SD
BRDi	0.678	0.73223165	0.729	0.782	0.08345325
HDACi	0.3965	0.4277573	0.4455	0.4765	0.06328818
DNMT 1	0.526	0.59578227	0.595	0.663	0.1255546
GRAS	0.471	0.60436491	0.607	0.739	0.18054764
Drugs	0.408	0.50042214	0.513	0.603	0.14793078
Clinic	0.485	0.56071604	0.577	0.652	0.13069903
General	0.504	0.57793955	0.587	0.661	0.11752032
Epi-Focused	0.51875	0.58460793	0.599	0.665	0.11791897

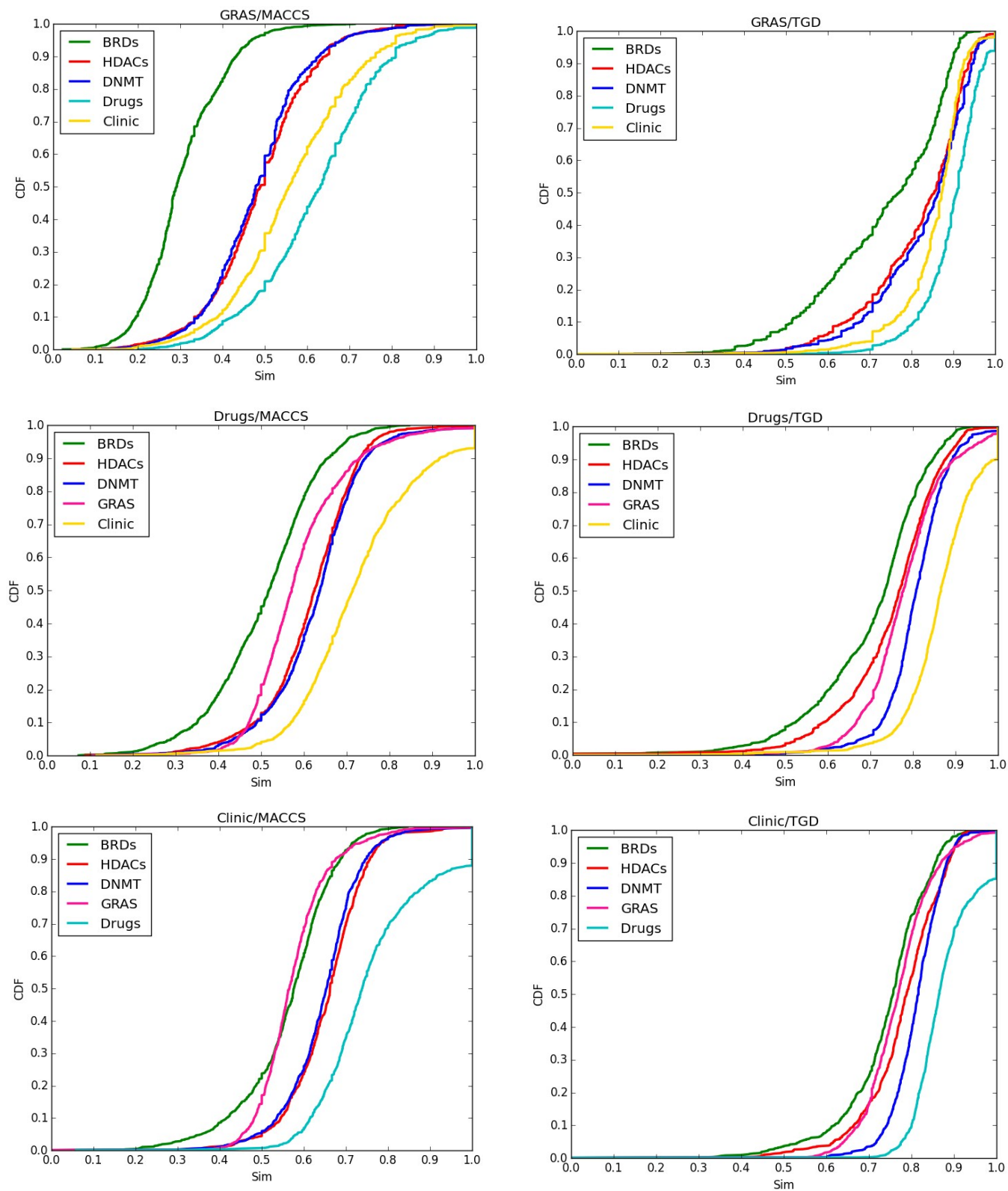


Figure S2. Inter-library similarity: Cumulative distribution function (CDF) of the maximum structure similarity calculated with MACCS keys, TDG and the Tanimoto coefficient comparing different reference and test sets. The reference set is indicated at the top of each graph. Summary statistics of the CDFs are presented in Tables S6 and S7.

Table S6. Summary statistics of the inter-library comparison using MACCS keys; the reference set is shown at the top of each table.

BRDi					
	HDACi	DNMT 1	GRAS	Drugs	Clinic
Min	0.5	0.521	0.431	0.535	0.536
Q1	0.610	0.590	0.511	0.647	0.642
Median	0.661	0.652	0.537	0.684	0.688
Q3	0.700	0.710	0.578	0.733	0.724
Max	1	0.867	0.712	0.846	0.846
SD	0.079	0.073	0.056	0.066	0.068

HDACi					
	BRDi	DNMT 1	GRAS	Drugs	Clinic
Min	0.204	0.392	0.372	0.468	0.395
Q1	0.544	0.586	0.518	0.618	0.636
Median	0.605	0.625	0.559	0.652	0.682
Q3	0.651	0.671	0.611	0.697	0.729
Max	1	1	1	1	1
SD	0.082	0.073	0.073	0.066	0.076

DNMT 1					
	BRDi	HDACi	GRAS	Drugs	Clinic
Min	0.073	0.088	0.153	0.333	0.181
Q1	0.538	0.639	0.531	0.651	0.646
Median	0.593	0.683	0.593	0.708	0.694
Q3	0.625	0.742	0.655	0.803	0.790
Max	0.867	1	1	1	1
SD	0.095	0.088	0.098	0.105	0.103

GRAS					
	BRDi	HDACi	DNMT 1	Drugs	Clinic
Min	0.022	0.052	0.058	0.2	0.045
Q1	0.351	0.445	0.423	0.411	0.507
Median	0.528	0.632	0.582	0.606	0.670
Q3	0.590	0.683	0.642	0.665	0.733
Max	0.712	1	1	1	1
SD	0.214	0.266	0.244	0.268	0.284

Drugs					
	BRDi	HDACi	DNMT 1	GRAS	Clinic
Min	0.073	0.088	0.108	0.105	0.115
Q1	0.429	0.560	0.571	0.514	0.633
Median	0.518	0.625	0.638	0.571	0.714
Q3	0.589	0.684	0.690	0.641	0.805
Max	0.846	1	1	1	1
SD	0.114	0.099	0.098	0.092	0.128

Table S6. (Continued)

	Clinic				
	BRDi	HDACi	DNMT 1	GRAS	Drugs
Min	0	0.022	0	0	0.058
Q1	0.513	0.604	0.600	0.522	0.675
Median	0.576	0.662	0.654	0.566	0.739
Q3	0.630	0.714	0.700	0.616	0.833
Max	0.846	1	1	1	1
SD	0.110	0.095	0.093	0.087	0.129

Q1: First quartile; Q3: Third quartile; SD: standard deviation

Table S7. Summary statistics of the inter-library comparison using TGD fingerprints; the reference set is shown at the top of each table.

BRDi					
	HDACi	DNMT 1	GRAS	Drugs	Clinic
Min	0.791	0.764	0.725	0.779	0.761
Q1	0.846	0.826	0.809	0.842	0.849
Median	0.862	0.85	0.832	0.862	0.868
Q3	0.885	0.868	0.873	0.885	0.887
Max	1	0.929	0.963	1	0.948
SD	0.035	0.034	0.048	0.032	0.032

HDACi					
	BRDi	DNMT 1	GRAS	Drugs	Clinic
Min	0.506	0.577	0.68	0.693	0.67
Q1	0.769	0.769	0.751	0.793	0.797
Median	0.800	0.800	0.775	0.828	0.830
Q3	0.845	0.845	0.805	0.857	0.867
Max	1	1	1	1	1
SD	0.048	0.048	0.049	0.044	0.045

DNMT 1					
	BRDs	HDACs	GRAS	Drugs	Clinic
Min	0	0	0.426	0.703	0.447
Q1	0.701	0.729	0.712	0.825	0.792
Median	0.752	0.783	0.752	0.857	0.842
Q3	0.812	0.861	0.799	0.889	0.871
Max	0.929	1	1	1	1
SD	0.090	0.086	0.079	0.054	0.059

GRAS					
	BRDi	HDACi	DNMT 1	Drugs	Clinic
Min	0	0	0	0	0
Q1	0.618	0.748	0.761	0.863	0.825
Median	0.771	0.855	0.863	0.907	0.873
Q3	0.863	0.908	0.913	0.943	0.905
Max	0.963	1	1	1	1
SD	0.153	0.121	0.117	0.075	0.085

Drugs					
	BRDs	HDACs	DNMT	GRAS	Clinic
Min	0	0	0	0	0
Q1	0.629	0.692	0.771	0.725	0.821
Median	0.734	0.767	0.808	0.778	0.866
Q3	0.790	0.825	0.850	0.832	0.918
Max	1	1	1	1	1
SD	0.134	0.122	0.081	0.098	0.092

Table S7. (Continued)

	Clinic				
	BRDi	HDACi	DNMT 1	GRAS	Drugs
Min	0	0	0	0	0
Q1	0.701	0.736	0.778	0.717	0.828
Median	0.757	0.789	0.816	0.769	0.865
Q3	0.808	0.840	0.853	0.816	0.919
Max	0.946	1	1	1	1
SD	0.104	0.095	0.065	0.083	0.076

Q1: First quartile; Q3: Third quartile; SD: standard deviation

Table S8. Descriptive statistics and AUC for CSR curves

Set	Q1	FS ₅₀	Q3	AUC	F ₅₀
BRDi	0.661	0.783	0.894	0.739	0.122
HDACi	0.669	0.802	0.901	0.762	0.098
DNMT 1	0.525	0.741	0.87	0.689	0.226
GRAS	0.909	0.948	0.974	0.926	0.004

Q1: First quartile; FS₅₀ fraction of database recovered by 50% of scaffolds; Q3: Third quartile; AUC: area under the curve; F50: fraction of scaffolds to recover 50% of the compound in the data set.

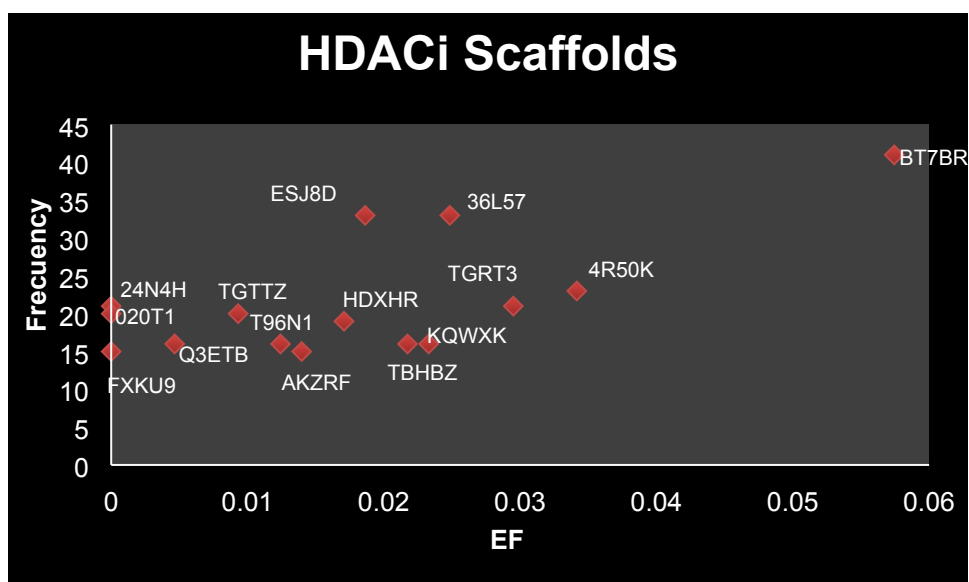
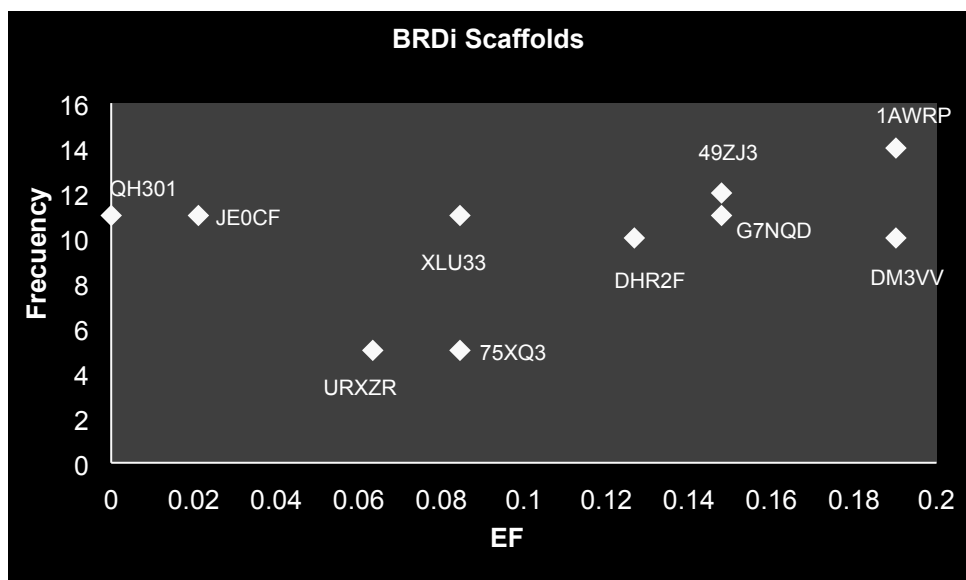


Figure S3. Chemotype enrichment plots for BRDi and HDACi data sets.

Table S9. Descriptive statistics for F-chiral and F-sp³ distributions

F-chiral								
	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General	Epi-focused
Min	0	0	0	0	0	0	0	0
1st Quart	0	0	0	0	0	0	0	0
Median	0	0	0	0	0.058	0.032	0	0
Mean	0.024	0.034	0.091	0.057	0.107	0.076	0.048	0.034
3rd Quart	0.045	0.043	0.142	0.1	0.173	0.105	0.058	0.038
Max	0.166	0.7	0.667	0.833	0.833	0.833	0.55	0.5

F-sp³								
	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General	Epi-focused
Min	1	1	0	0	0	0	0	0
1st Quart	3	3	0.08	0.4	0.29	0.24	0.19	0.13
Median	4	4	0.2	0.67	0.43	0.38	0.3	0.28
Mean	3.662	4.813	0.264	0.622	0.458	0.401	0.325	0.271
3rd Quart	4	5	0.42	0.86	0.61	0.53	0.44	0.37
Max	7	37	0.91	1	1	1	1	0.68

Table S10. Nemenyi pairing values for F-chiral and F-sp3 distributions (1 BRDi; 2 HDACi; 3 DNMT 1; 4 GRAS; 5 Drugs; 6 Clinic; 7 General; 8 Epi-focused).

F-chiral

	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General
2	0.9991	-	-	-	-	-	-
3	0.04151	9.90E-08	-	-	-	-	-
4	0.99064	0.0603	0.0013	-	-	-	-
5	2.60E-11	<2E-16	5.00E-07	< 2E-16	-	-	-
6	0.0044	2.40E-13	0.996	4.10E-07	1.00E-06	-	-
7	0.971	0.08	0.017	0.999	< 2E-16	9.00E-05	-
8	0.912	0.927	0.002	0.372	1.10E-10	0.0003	0.305

F-sp³

	BRDi	HDACi	DNMT 1	GRAS	Drugs	Clinic	General
2	0.021	-	-	-	-	-	-
3	0.951	9.90E-08	-	-	-	-	-
4	< 2E-16	< 2E-16	< 2E-16	-	-	-	-
5	< 2E-16	< 2E-16	< 2E-16	< 2E-16	-	-	-
6	0.0044	2.40E-13	7.60E-14	< 2E-16	4.50E-05	-	-
7	0.971	0.08	1.40E-05	< 2E-16	< 2E-16	4.20E-09	-
8	0.912	0.927	1	< 2E-16	6.80E-12	1.20E-05	0.382