#### **Supplementary Information**

#### Electrophilic MiniFrags revealed unprecedented binding sites for covalent HDAC8 inhibitors

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**Supplementary Figure S1.** Thiol reactivity table for heterocyclic electrophiles. Blue dots indicate the position of the warhead. Colouring is in line with the activity, as the darker green colour refers for higher reaction rate and lower half life. Compounds labelled by italic showed parallel reaction with the assay buffer. N.A. stands for "Not available".

					L T	F	Ē			<u> </u>	К	L	N	Р	Q	R
											HZ,Z	HZ,Z	C <sub>N</sub>	<sup>O</sup> <sub>N</sub>	<b>S</b> N	S N
	t <sub>1/2</sub> (h) of M+ fragment (250 uM) with GSH (5 mM)															
1	Cl	>48	>48	>48	>48	>48	>48	>48	>48	>48	N.A.	>48	N.A.	N.A.	N.A.	N.A.
2	Br	35.20	>48	>48	>48	N.A.	>48	>48	>48	>48	>48	>48	N.A.	N.A.	>48	>48
3	1	>48	>48	>48	>48	N.A.	>48	>48	0.16	>48	>48	>48	0.23	>48	N.A.	N.A.
4	CN	>48	40.90	0.79	2.00	40.50	>48	21.90	>48	>48	N.A.	>48	N.A.	N.A.	N.A.	8.20
5	Vinyl	1.00	>48	0.26	>48	N.A.	>48	18.50	N.A.	N.A.	>48	N.A.	N.A.	N.A.	N.A.	N.A.
6	Ethynyl	>48	>48	2.40	43.80	N.A.	>48	>48	N.A.	N.A.	5.70	2.20	N.A.	N.A.	N.A.	N.A.

**Supplementary Figure S2.** Dose-response curves for the determination of  $IC_{50}$ -values of HDAC8 biochemical assay with the heterocyclic MiniFrags.



Supplementary Figure S3. Dose-response curves for the determination of  $IC_{50}$ -values on 10 nM HDAC8 and active site mutants with the heterocyclic MiniFrags.



Supplementary Figure S4. Dose-response curves for the determination of  $IC_{50}$ -values on 100 nM HDAC8 mutants with the heterocyclic MiniFrags.



Supplementary Figure S5. The MSMS spectrum and annotation of C6+, D1+, F2+, D3+, F3+, D6+, G6+ modified HDAC8 peptide [1-34]





G6+



n/z, [a

Supplementary Figure S6. The MSMS spectrum and annotation of C3+ modified HDAC8 peptide [33-83]



Supplementary Figure S7. The MSMS spectrum and annotation of A3+, B6+, C3+, C6+, D1+, D2+, F2+, D3+, F3+, G6+, 3 modified HDAC8 peptide [148-165]



S7











F3+



G6+



# Supplementary Figure S8. The MSMS spectrum and annotation of A3+, C3+, C6+, D1+, D2+, D3+ modified HDAC8 peptide [241-250]















# **Supplementary Figure S9.** The MSMS spectrum and annotation of **C3+**, **F2+**, **D3+**, **F3+**, **G6+** modified HDAC8 peptide [260-287]





G6+

**Supplementary Figure S10.** The MSMS spectrum and annotation of **F3+** modified HDAC8 peptide [327-357]





**Supplementary Figure S11.** Results of the dilution experiments proving the irreversible covalent labeling for compounds **3** and **B6+** 



Reversibility of **B6+** and **3** on HDAC8 WT shown as relative enzyme activity before and after dialysis. Error bars represent mean +/- SD, n = 3



#### Supplementary Figure S12. MS Chromatogram for showing the labelling of GSH with F4+

**Supplementary Figure S13.** Results of time dependent  $IC_{50}$  measurements and K<sub>i</sub>,  $k_{inact}$ ,  $k_{inact}/K_i$  calculations



Supplementary Figure S14. Results of the western blot after 72 h treatment of OCI-AML3 cells



### <sup>1</sup>H, <sup>13</sup>C NMR spectra and HPLC-MS chromatogram and spectrum of **10**





Peak#	Ret. Time	Height	Area	Area%	Peak#	Ret. Time	Height	Area	Area%
1	0.725	24741	37777	0.532	1	0.731	1889	1712	0.14
2	1.398	6182	14949	0.21	2	1.481	7918	15921	1.302
3	1.48	7303	13471	0.19	3	1.579	442987	1189213	97.282
4	1.578	2451693	7026877	98.907	4	4.677	8151	15595	1.276
5	4.677	9501	11489	0.162	Total		460945	1222441	100
Total		2499419	7104563	100					

#### <sup>1</sup>H, <sup>13</sup>C NMR spectra and HPLC-MS chromatogram and spectrum of **2**





Peak#	Ret. Time	Height	Area	Area%	Peak#	Ret. Time	Height	Area	Area%
1	0.733	3107	2315	0.143	1	0.729	48007	116486	2.516
2	1.399	1313	5120	0.316	2	1.397	8332	44423	0.959
3	1.618	575510	1595585	98.365	3	1.618	1520747	4449011	96.093
4	2.536	1438	19094	1.177	4	2.428	4781	17338	0.374
Total		581368	1622114	100	5	4.692	7435	2642	0.057
					Total		1589303	4629901	100

# $^1\text{H},\,^{13}\text{C}$ NMR spectra and HPLC-MS chromatogram and spectrum of 12





Peak#	Ret. Time	Height	Area	Area%	Peak#	Ret. Time	Height	Area	Area%
1	1.188	4855	19412	0.632	1	1.188	4855	19412	0.632
2	1.369	6076	32406	1.055	2	1.369	6076	32406	1.055
3	2.623	883032	3009149	97.981	3	2.623	883032	3009149	97.981
4	8.907	10740	10203	0.332	4	8.907	10740	10203	0.332
Total		904703	3071170	100	Total		904703	3071170	100



#### <sup>1</sup>H, <sup>13</sup>C NMR spectra and HPLC-MS chromatogram and spectrum of **4**





Peak#	Ret. Time	Height	Area	Area%	Peak#	Ret. Time	Height	Area	Area%
1	1.412	11222	55150	3.429	1	1.37	165	230	0.149
2	1.851	277	-634	-0.039	2	1.773	546	1103	0.713
3	2.108	4136	13261	0.825	3	2.099	429	1465	0.948
4	2.598	471237	1539452	95.727	4	2.598	48676	151770	98.19
5	7.587	1097	932	0.058	Total		49816	154568	100
Total		487970	1608161	100					



## $^1\text{H},\,^{13}\text{C}$ NMR spectra and HPLC-MS chromatogram and spectrum of 11



Peak#	Ret. Time	Height	Area	Area%	Peak#	Ret. Time	Height	Area	Area%
1	0.714	22833	46626	1.416	1	0.777	3516	7793	1.216
2	1.082	15914	48991	1.488	2	1.309	8676	17482	2.728
3	1.432	1266827	3197446	97.096	3	1.432	294473	615542	96.056
Total		1305574	3293063	100	Total		306664	640818	100



# $^1\text{H},\,^{13}\text{C}$ NMR spectra and HPLC-MS chromatogram and spectrum of 3



Peak#	Ret. Time	Height	Area	Area%	Peak#	Ret. Time	Height	Area	Area%
1	0.566	92037	113344	2.367	1	0.597	648	751	0.116
2	1.065	23731	105671	2.207	2	0.75	5234	11604	1.796
3	1.368	1130888	4558667	95.201	3	1.367	233787	633822	98.088
4	1.666	5801	10776	0.225	Total		239670	646177	100
Total		1252457	4788458	100					

### Supplementary Tables S1, S2, S3

