

Supporting Information

Rational Design and Synthesis of Right-Handed D-Sulfono- γ -AApeptide Helical Foldamers as Potent Inhibitors of Protein-Protein Interactions

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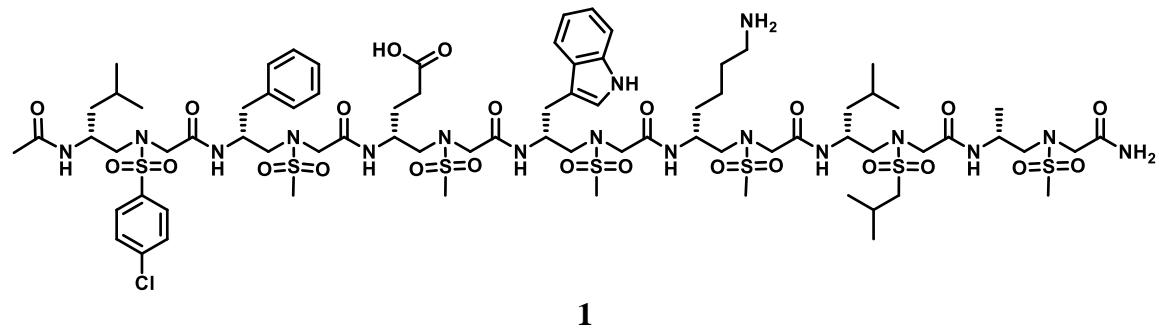
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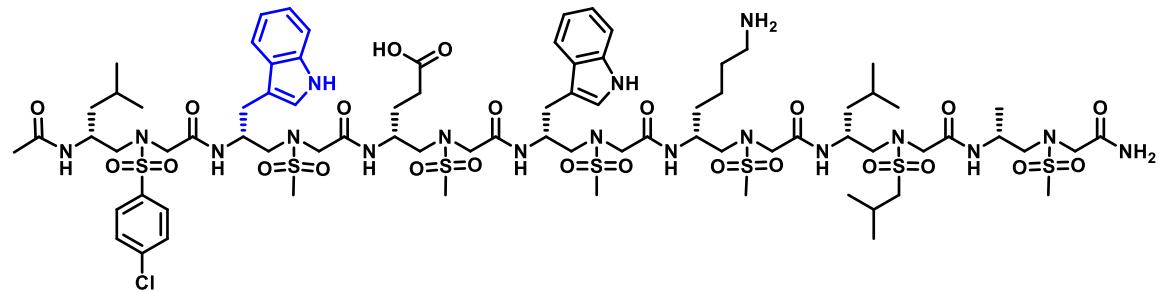
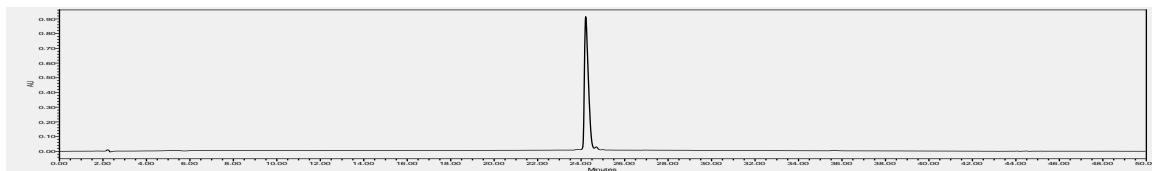
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1. D-Sulfono- γ -AApeptides

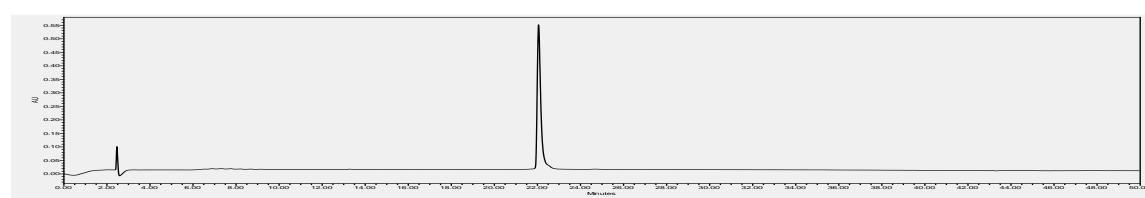
1.1 HPLC Trace

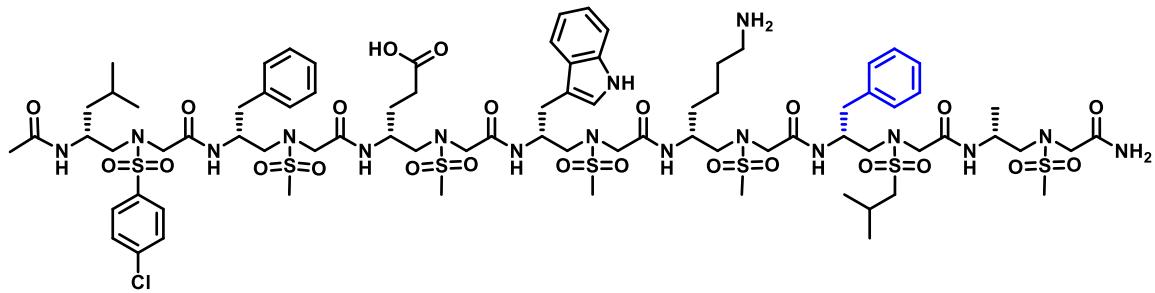


HRMS (ESI) ($[M+H]^+$) Calcd. for $C_{77}H_{127}ClN_{17}O_{24}S_7$: 1932.6973, found: 967.8494
 $[M+2H]^{2+}$, 989.8296 $[M+2Na]^{2+}$.



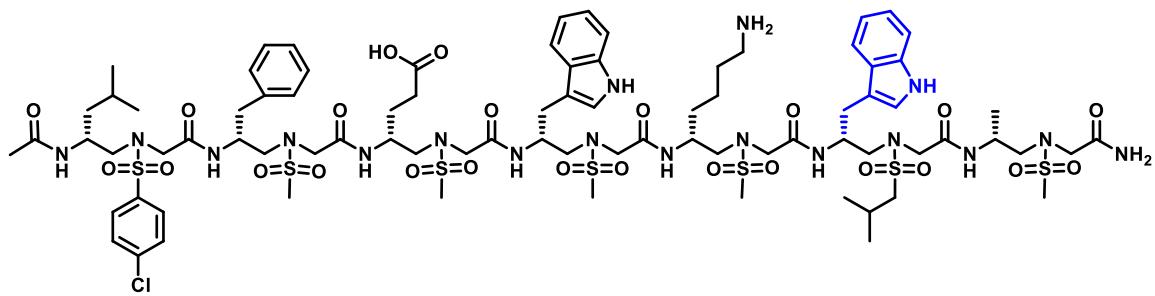
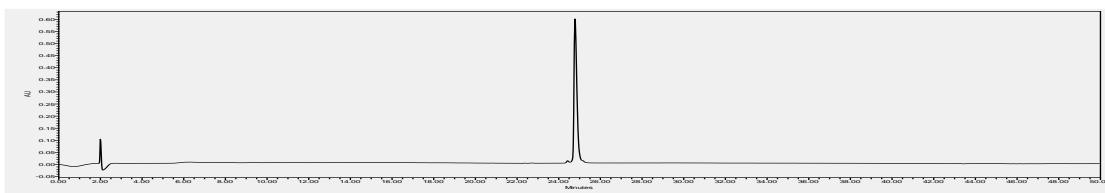
HRMS (ESI) ($[M+H]^+$) Calcd. for $C_{79}H_{128}ClN_{18}O_{24}S_7$: 1971.7082, found: 987.3563
 $[M+2H]^{2+}$, 1008.3370 $[M+2Na]^{2+}$.





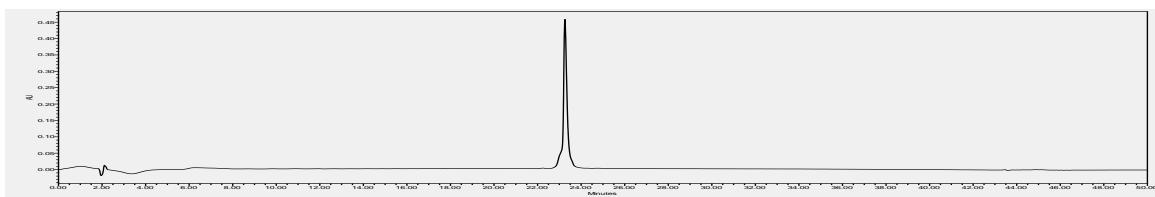
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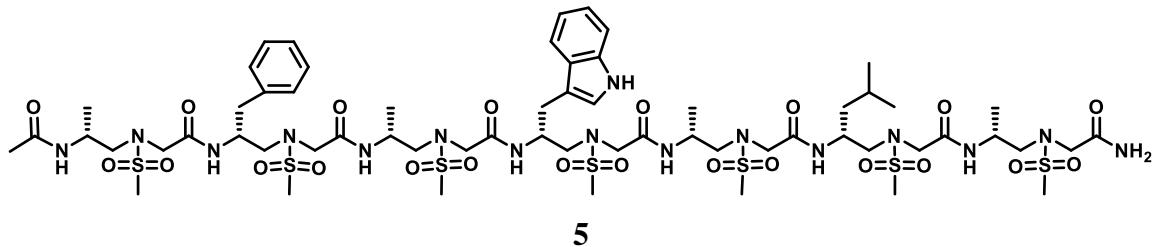
HRMS (ESI) ($[M+H]^+$) Calcd. for $C_{80}H_{125}ClN_{17}O_{24}S_7$: 1966.6817, found: 984.8426
 $[M+2H]^{2+}$, 1006.8241 $[M+2Na]^{2+}$, 1022.7915 $[M+2K]^{2+}$.



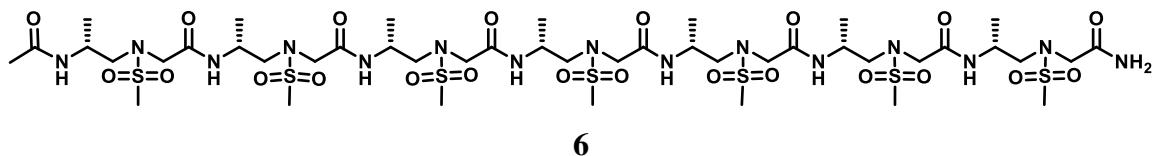
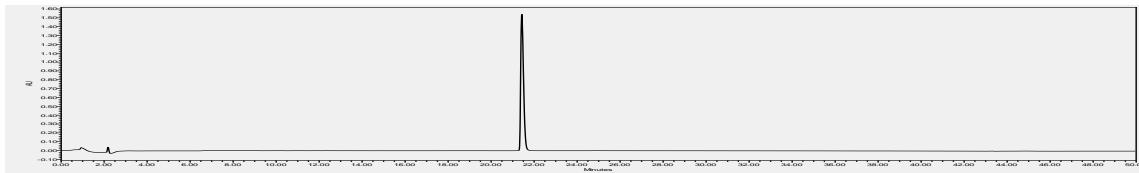
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HRMS (ESI) ($[M+H]^+$) Calcd. for $C_{82}H_{126}ClN_{18}O_{24}S_7$: 2005.6926, found: 1004.3481
 $[M+2H]^{2+}$, 1026.3316 $[M+2Na]^{2+}$.

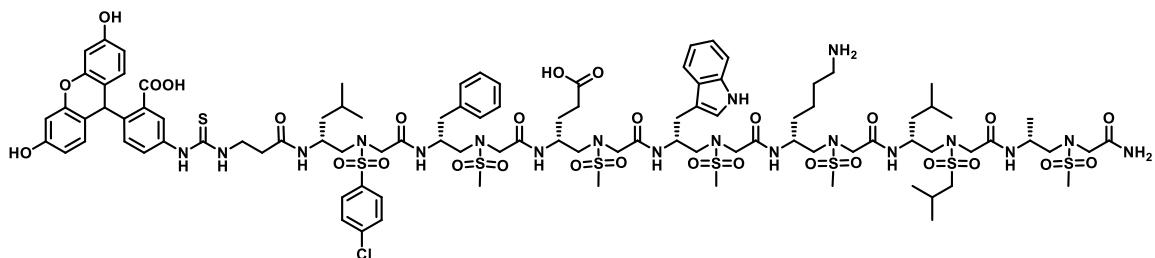
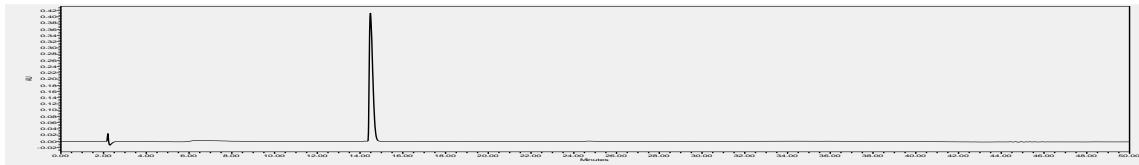




HRMS (ESI) ($[M+H]^+$) Calcd. for $C_{61}H_{105}N_{16}O_{22}S_7$: 1637.5634, found: 819.2843 $[M+2H]^{2+}$.

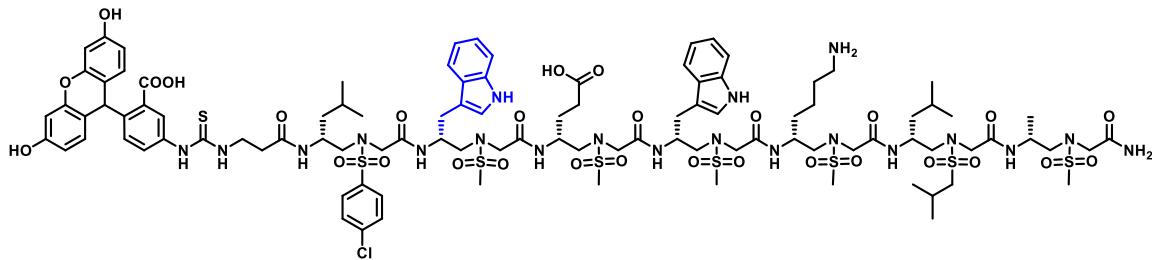
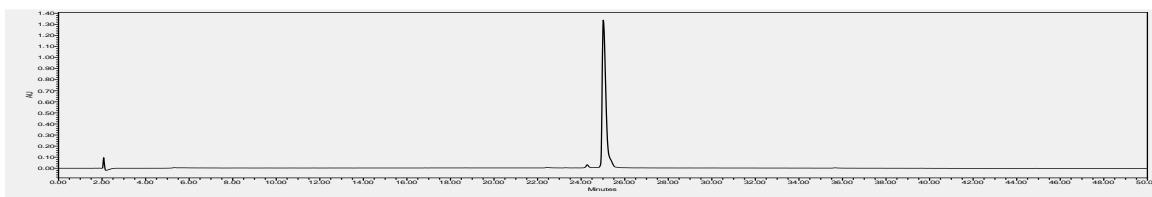


HRMS (ESI) ($[M+H]^+$) Calcd. for $C_{44}H_{90}N_{15}O_{22}S_7$: 1404.4430, found: 702.7242 $[M+2H]^{2+}$, 724.7053 $[M+2Na]^{2+}$.

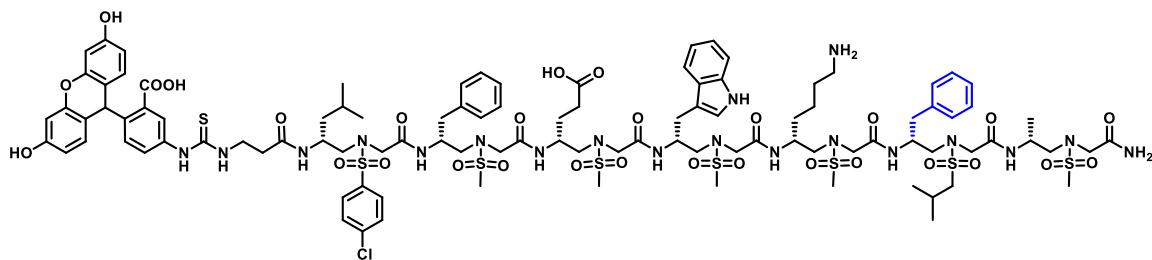
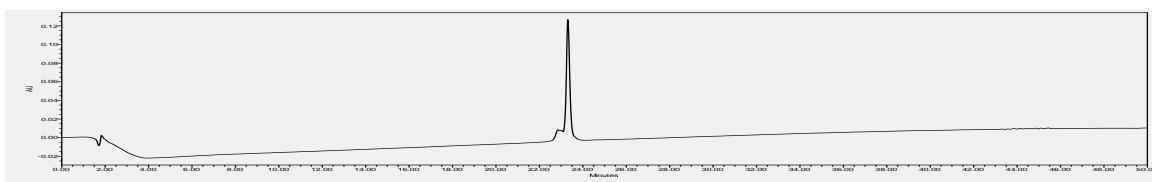


1-FITC

HRMS (ESI) ($[M+H]^+$) Calcd. for $C_{99}H_{143}ClN_{19}O_{29}S_8$: 2352.7753, found: 1176.8810 $[M+2H]^{2+}$.

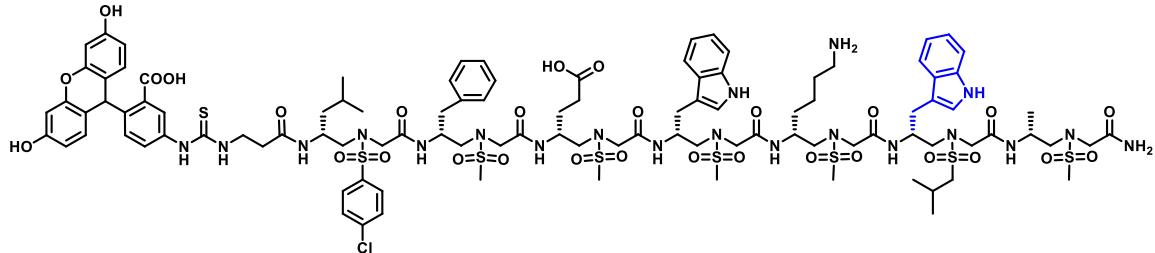
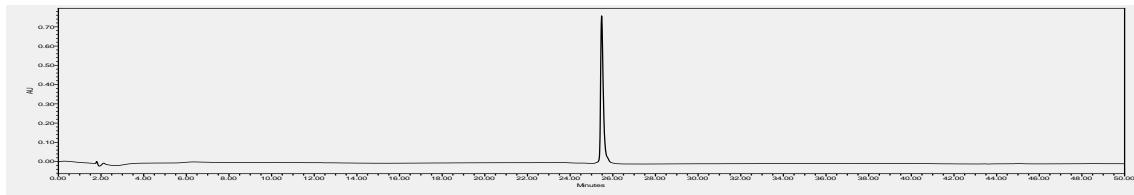


HRMS (ESI) ($[M+H]^+$) Calcd. for $C_{101}H_{144}ClN_{20}O_{29}S_8$: 2391.7862, found: 1196.3865
 $[M+2H]^{2+}$.



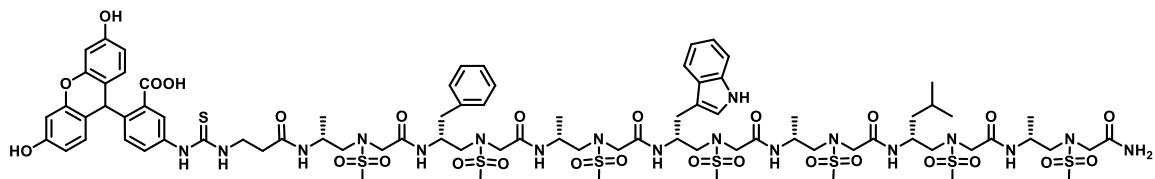
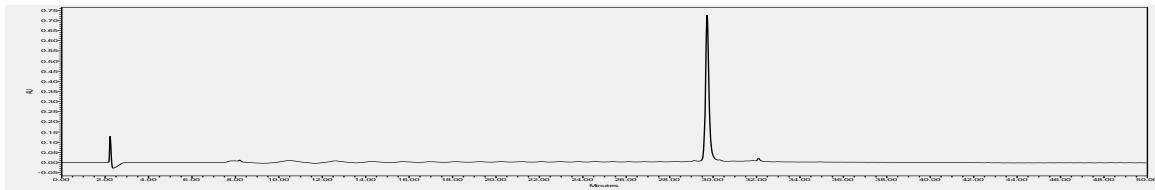
3-FITC

HRMS (ESI) ($[M+H]^+$) Calcd. for $C_{102}H_{141}ClN_{19}O_{29}S_8$: 2386.7597, found: 1193.8731
 $[M+2H]^{2+}$.



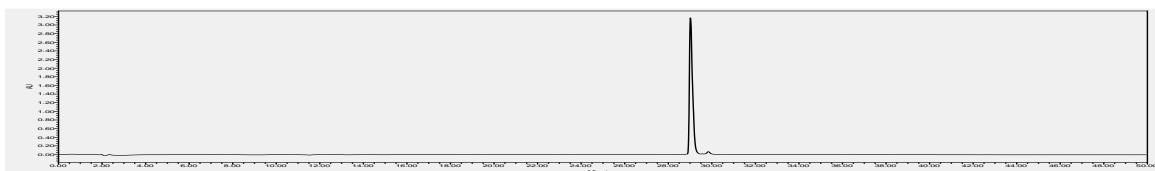
4-FITC

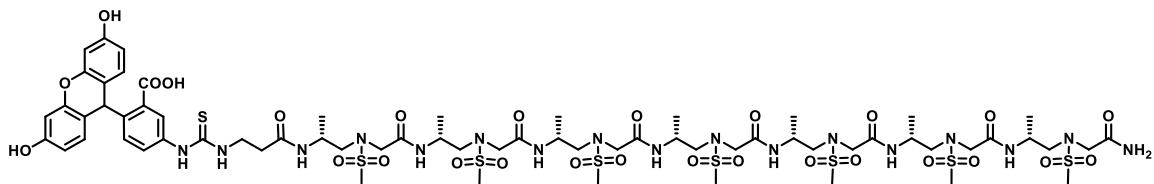
HRMS (ESI) ($[M+H]^+$) Calcd. for $C_{104}H_{142}ClN_{20}O_{29}S_8$: 2425.7706, found: 809.2550 $[M+3H]^{3+}$.



5-FITC

HRMS (ESI) ($[M+H]^+$) Calcd. for $C_{83}H_{121}N_{18}O_{27}S_8$: 2057.6414, found: 1029.3132 $[M+2H]^{2+}$.





6-FITC

HRMS (ESI) ($[M+H]^+$) Calcd. for $C_{66}H_{106}N_{17}O_{27}S_8$: 1824.5210, found: 912.7547
 $[M+2H]^{2+}$, 934.7352 $[M+2Na]^{2+}$.

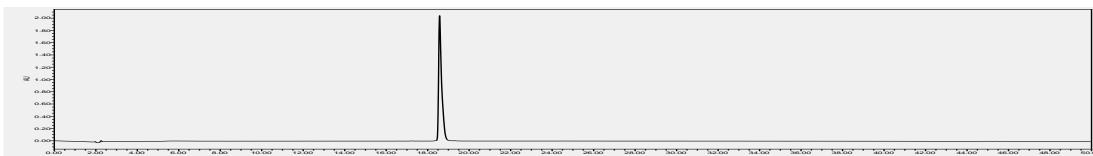


Figure S1. HPLC spectra of D-sulfono- γ -AApeptides

1.2 HPLC Purities and Retention Time of Pure Peptides.

Table S1. HPLC purities and retention time of D-sulfono- γ -AApeptides.^a

Peptide Name	Purity trace after HPLC purification (%)	Retention Time (min)
1	99.56%	24.211
2	100.00%	22.054
3	98.86%	24.784
4	99.10%	23.255
5	100.00%	21.426
6	100.00%	14.484
1-FITC	98.71%	25.021
2-FITC	96.66%	23.312
3-FITC	100.00%	25.476
4-FITC	98.96%	29.723
5-FITC	99.10%	29.021
6-FITC	100.00%	18.391

^a The gradient eluting method of 5% to 100% of solvent B (0.1% TFA in acetonitrile) in A (0.1% TFA in water) over 50 min was performed.

2. Fluorescence Polarization Assay

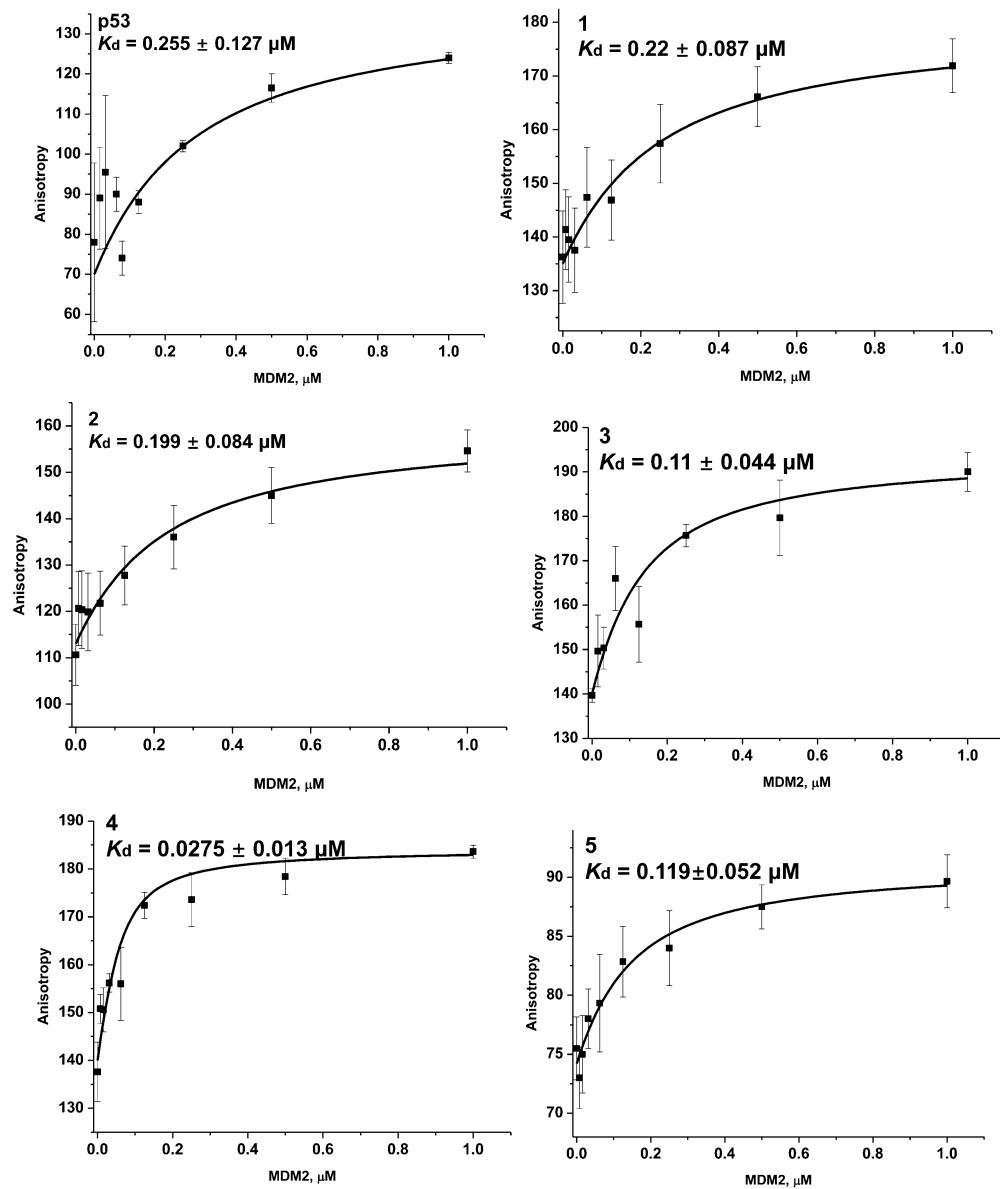


Figure S2. The K_d data of regular p53 and D-sulfono- γ -AApeptides 1-5.

3. ^{15}N - ^1H HSQC NMR of Lead Peptide 4 in Complex with MDM2

Methods

Protein Purification for NMR

^{15}N -labeled human MDM2 residues 17–125 was expressed and purified as described in [2].

NMR Data Collection and Analysis

Experiments for 200 μ M MDM2₁₇₋₁₂₅ in the presence and absence of stoichiometric equivalent of peptide **4** were carried out at 25 °C on a Varian VNMRS 800-MHz spectrometry with a triple resonance pulse field Z-axis gradient cold probe at 30 °C. ^1H - ^{15}N heteronuclear single-quantum coherence spectroscopy experiments were performed on ^{15}N -labeled samples in 90% H₂O/ 10% D₂O. Buffer for peptide **4** and MDM2₁₇₋₁₂₅ experiments was 50 mM NaH₂PO₄, 50 mM NaCl, 1 mM EDTA, 2 mM DTT, 5% DMSO, and 0.02% NaN₃ at pH 6.8. Data were acquired in the ^1H and ^{15}N dimension using 9689.92-Hz (t_2) \times 2430.26-Hz (t_1) sweep widths and 1024 (t_2) \times 128 (t_1) complex data points. Bound spectra were collected in a molar equivalent of the peptide **4**.

Resonance assignments for apo MDM2₁₇₋₁₂₅ were previously made [2]. Bound chemical shifts were inferred based on the overlap of resonances in parallel titrations (Figure S3). Using this method, the bound resonances for 99 residues were assigned. For the nine unassignable residues, four were prolines which are not detected, two residues, S17 and Q18, were near the N-terminus, and the remaining three were D46, L81, and N106 were not assigned in apo MDM2₁₇₋₁₂₅ [2]. In addition, residue Y67 experienced line broadening during titration and the bound peak was not observed.

The combined average chemical shifts were calculated from the formula $\Delta_{\text{ave}} = [((\Delta^1\text{H}^{\text{N}})^2 + (\Delta^{15}\text{N}/5)^2)/2]^{1/2}$. All NMR spectra were processed with NMRFx and analyzed using NMRViewJ software [3, 4].

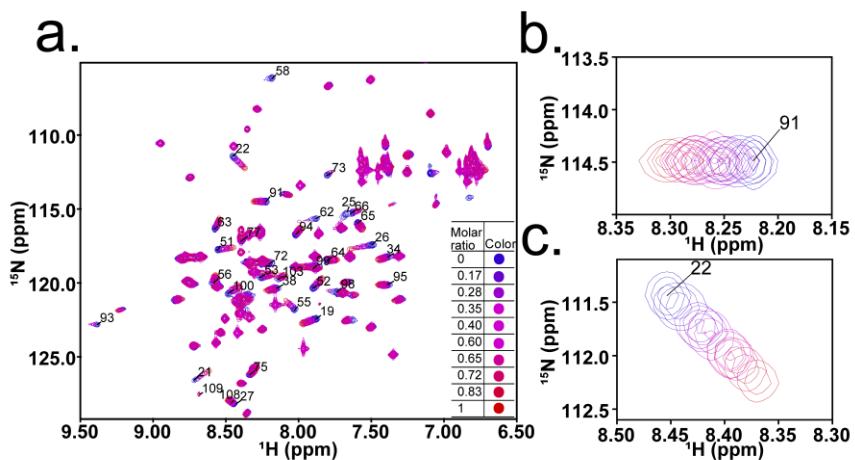


Figure S3. NMR titration results and resonance assignments of MDM2₁₇₋₁₂₅ interacting with **4**. **a.** Overlay of ^1H - ^{15}N HSQC spectra for MDM2₁₇₋₁₂₅ before (blue) and after the addition of increasing concentrations of **4** (see legend for molar ratios of titration points).

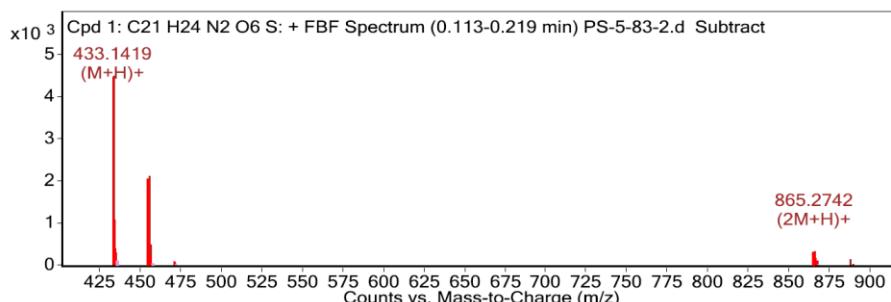
For clarity residue labels are shown for resonances with chemical shift changes greater than 0.032 ppm. **b.** Chemical shifts of F91 during the titration show clear overlap of free and bound resonances. **c.** Chemical shifts of S22 during the titration show clear overlap of free and bound resonances.

4. References

- (1) Sang, P.; Zhang, M.; Shi, Y.; Li, C.; Abdulkadir, S.; Li, Q.; Ji, H.; Cai, J., *Proc. Natl. Acad. Sci. U. S. A.*, **2019**, 116, 10757-10762.
- (2) Borcherds, W.; Theillet, F.-X.; Katzer, A.; Finzel, A.; Mishall, K. M.; Powell, A. T.; Wu, H.; Manieri, W.; Dieterich, C.; Selenko, P.; Loewer, A.; Daughdrill, G. W., *Nat. Chem. Biol.*, **2014**, 10 (12), 1000-1002
- (3) Johnson, B.A., Using NMRView to visualize and analyze the NMR spectra of macromolecules, in *Protein NMR Techniques*. **2004**, Springer. p. 313-352.
- (4) Johnson, B.A.; Blevins, R.A., *J. biomol. NMR*, **1994**, 4(5), 603-614.

5. HRMS Spectra of Building Blocks 1a-j and D-Sulfono- γ -AApeptides

HRMS spectra of 1a

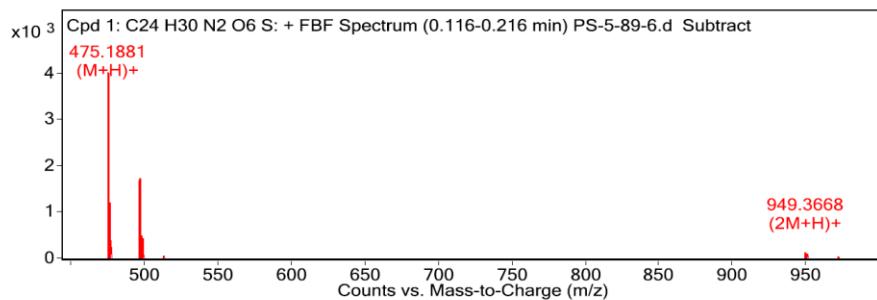


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
433.1419	1	4511.36	C21H25N2O6S	(M+H)+
434.1447	1	1101.61	C21H25N2O6S	(M+H)+
435.1424	1	324.79	C21H25N2O6S	(M+H)+
455.1229	1	2135.01	C21H24N2NaO6S	(M+Na)+
456.1259	1	525.02	C21H24N2NaO6S	(M+Na)+
457.1226	1	52.91	C21H24N2NaO6S	(M+Na)+
471.0901	1	79.55	C21H24KN2O6S	(M+K)+
865.2742	1	359.58	C42H49N4O12S2	(2M+H)+
866.2785	1	129.05	C42H49N4O12S2	(2M+H)+
887.2566	1	164.59	C42H48N4NaO12S2	(2M+Na)+

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HRMS spectra of 1b

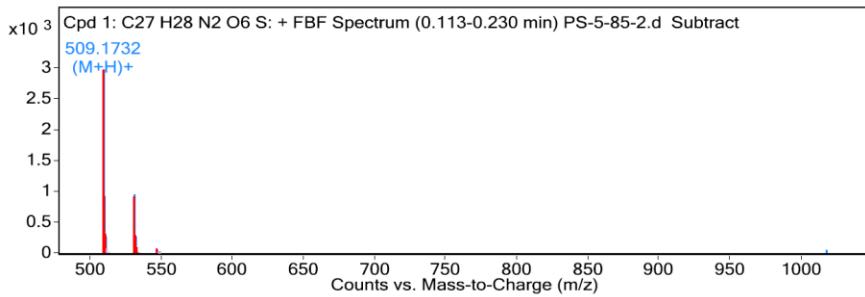


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
475.1881	1	4027.52	C ₂₄ H ₃₁ N ₂ O ₆ S	(M+H) ⁺
476.1914	1	1223.82	C ₂₄ H ₃₁ N ₂ O ₆ S	(M+H) ⁺
477.1893	1	273.12	C ₂₄ H ₃₁ N ₂ O ₆ S	(M+H) ⁺
497.1702	1	1749.32	C ₂₄ H ₃₀ N ₂ NaO ₆ S	(M+Na) ⁺
498.1718	1	450.11	C ₂₄ H ₃₀ N ₂ NaO ₆ S	(M+Na) ⁺
499.1718	1	102.88	C ₂₄ H ₃₀ N ₂ NaO ₆ S	(M+Na) ⁺
513.1378	1	54.12	C ₂₄ H ₃₀ KN ₂ O ₆ S	(M+K) ⁺
949.3668	1	111.59	C ₄₈ H ₆₁ N ₄ O ₁₂ S ₂	(2M+H) ⁺
950.368	1	82	C ₄₈ H ₆₁ N ₄ O ₁₂ S ₂	(2M+H) ⁺
971.3451	1	81.86	C ₄₈ H ₆₀ N ₄ NaO ₁₂ S ₂	(2M+Na) ⁺

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HRMS spectra of 1c

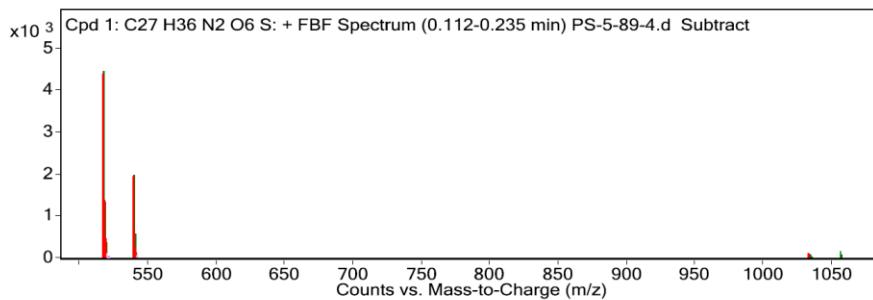


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
509.1732	1	2994.12	C ₂₇ H ₂₉ N ₂ O ₆ S	(M+H) ⁺
510.1754	1	926.66	C ₂₇ H ₂₉ N ₂ O ₆ S	(M+H) ⁺
511.1737	1	290.31	C ₂₇ H ₂₉ N ₂ O ₆ S	(M+H) ⁺
531.1538	1	965.32	C ₂₇ H ₂₈ N ₂ NaO ₆ S	(M+Na) ⁺
532.1577	1	291.12	C ₂₇ H ₂₈ N ₂ NaO ₆ S	(M+Na) ⁺
533.1565	1	30.93	C ₂₇ H ₂₈ N ₂ NaO ₆ S	(M+Na) ⁺
547.1198	1	67.87	C ₂₇ H ₂₈ KN ₂ O ₆ S	(M+K) ⁺
1017.3375	1	79.76	C ₅₄ H ₅₇ N ₄ O ₁₂ S ₂	(2M+H) ⁺

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HRMS spectra of 1d

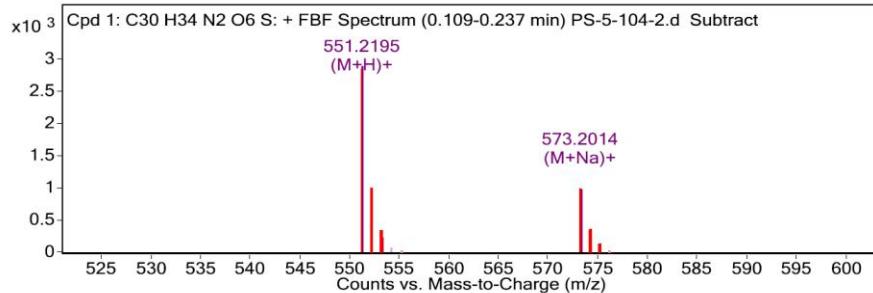


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
517.2355	1	4447.79	C ₂₇ H ₃₇ N ₂ O ₆ S	(M+H) ⁺
518.2377	1	1358.69	C ₂₇ H ₃₇ N ₂ O ₆ S	(M+H) ⁺
519.2376	1	410.22	C ₂₇ H ₃₇ N ₂ O ₆ S	(M+H) ⁺
539.217	1	1988.83	C ₂₇ H ₃₆ N ₂ NaO ₆ S	(M+Na) ⁺
540.2202	1	613.62	C ₂₇ H ₃₆ N ₂ NaO ₆ S	(M+Na) ⁺
541.2178	1	153.67	C ₂₇ H ₃₆ N ₂ NaO ₆ S	(M+Na) ⁺
1033.4607	1	118.26	C ₅₄ H ₇₃ N ₄ O ₁₂ S ₂	(2M+H) ⁺
1034.4656	1	64.81	C ₅₄ H ₇₃ N ₄ O ₁₂ S ₂	(2M+H) ⁺
1055.4442	1	188.55	C ₅₄ H ₇₂ N ₄ NaO ₁₂ S ₂	(2M+Na) ⁺
1056.4472	1	107.43	C ₅₄ H ₇₂ N ₄ NaO ₁₂ S ₂	(2M+Na) ⁺

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HRMS spectra of 1e

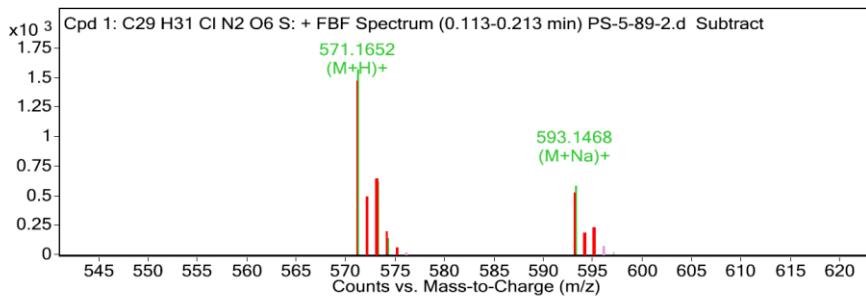


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
551.2195	1	2890.08	C ₃₀ H ₃₅ N ₂ O ₆ S	(M+H) ⁺
552.2227	1	1020.49	C ₃₀ H ₃₅ N ₂ O ₆ S	(M+H) ⁺
553.2221	1	244.34	C ₃₀ H ₃₅ N ₂ O ₆ S	(M+H) ⁺
573.2014	1	1004.49	C ₃₀ H ₃₄ N ₂ NaO ₆ S	(M+Na) ⁺
574.2041	1	372.53	C ₃₀ H ₃₄ N ₂ NaO ₆ S	(M+Na) ⁺
575.2039	1	57.92	C ₃₀ H ₃₄ N ₂ NaO ₆ S	(M+Na) ⁺

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HRMS spectra of 1f

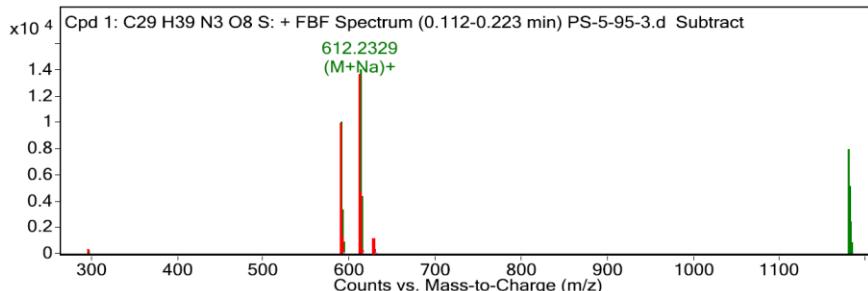


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
571.1652	1	1568.21	C ₂₉ H ₃₂ ClN ₂ O ₆ S	(M+H) ⁺
572.1686	1	474.51	C ₂₉ H ₃₂ ClN ₂ O ₆ S	(M+H) ⁺
573.1635	1	629.57	C ₂₉ H ₃₂ ClN ₂ O ₆ S	(M+H) ⁺
574.1641	1	148.2	C ₂₉ H ₃₂ ClN ₂ O ₆ S	(M+H) ⁺
575.1581	1	22.59	C ₂₉ H ₃₂ ClN ₂ O ₆ S	(M+H) ⁺
593.1468	1	585.31	C ₂₉ H ₃₁ ClN ₂ NaO ₆ S	(M+Na) ⁺
594.1481	1	163.69	C ₂₉ H ₃₁ ClN ₂ NaO ₆ S	(M+Na) ⁺
595.1448	1	182.49	C ₂₉ H ₃₁ ClN ₂ NaO ₆ S	(M+Na) ⁺

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HRMS spectra of 1g

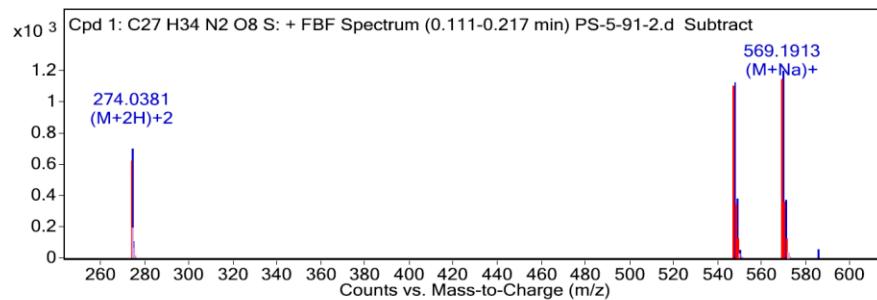


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
295.5567	2	326.82	C ₂₉ H ₄₁ N ₃ O ₈ S	(M+2H) ⁺²
590.2509	1	10094.98	C ₂₉ H ₄₀ N ₃ O ₈ S	(M+H) ⁺
591.254	1	3431.81	C ₂₉ H ₄₀ N ₃ O ₈ S	(M+H) ⁺
612.2329	1	14097.46	C ₂₉ H ₃₉ N ₃ NaO ₈ S	(M+Na) ⁺
613.2357	1	4482.54	C ₂₉ H ₃₉ N ₃ NaO ₈ S	(M+Na) ⁺
614.235	1	1361.39	C ₂₉ H ₃₉ N ₃ NaO ₈ S	(M+Na) ⁺
628.2025	1	1126.39	C ₂₉ H ₃₉ KN ₃ O ₈ S	(M+K) ⁺
1179.495	1	8017.3	C ₅₈ H ₇₉ N ₆ O ₁₆ S ₂	(2M+H) ⁺
1180.498	1	5193.55	C ₅₈ H ₇₉ N ₆ O ₁₆ S ₂	(2M+H) ⁺
1181.4986	1	2517.77	C ₅₈ H ₇₉ N ₆ O ₁₆ S ₂	(2M+H) ⁺

--- End Of Report ---

HRMS spectra of 1h

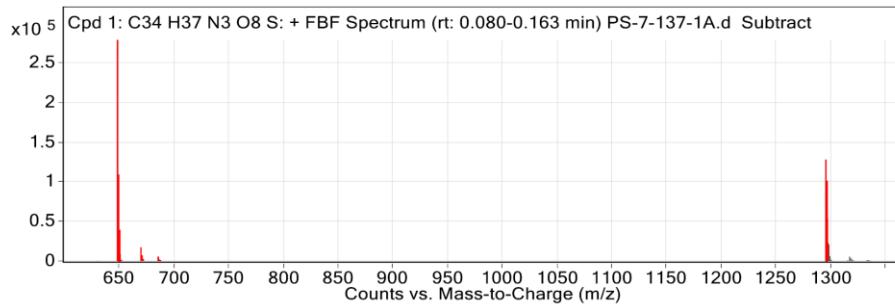


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
274.0381	2	705.22	C27H36N2O8S	(M+2H)+2
274.5394	2	114.17	C27H36N2O8S	(M+2H)+2
547.2094	1	1129.57	C27H35N2O8S	(M+H)+
548.2118	1	384	C27H35N2O8S	(M+H)+
549.2106	1	56.93	C27H35N2O8S	(M+H)+
569.1913	1	1202.22	C27H34N2NaO8S	(M+Na)+
570.195	1	381.85	C27H34N2NaO8S	(M+Na)+
571.1941	1	45.25	C27H34N2NaO8S	(M+Na)+
585.1541	1	65.99	C27H34KN2O8S	(M+K)+

--- End Of Report ---

HRMS spectra of 1i

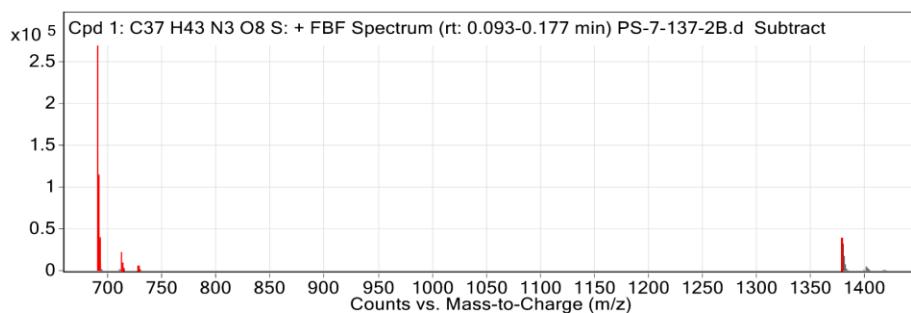


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
630.176	1	179.22	C34H37N3O8S	(M+H)+[−H ₂ O]
648.236	1	278874.63	C34H37N3O8S	(M+H)+
649.2388	1	107343.41	C34H37N3O8S	(M+H)+
670.217	1	17149.75	C34H37N3O8S	(M+Na)+
686.1883	1	5279.58	C34H37N3O8S	(M+K)+
1295.4637	1	128047.95	C34H37N3O8S	(2M+H)+
1296.4672	1	100762.45	C34H37N3O8S	(2M+H)+
1315.4918	1	703.74	C34H37N3O8S	(2M+K)+[−H ₂ O]
1317.4441	1	5706.52	C34H37N3O8S	(2M+Na)+
1333.4162	1	1215.78	C34H37N3O8S	(2M+K)+

--- End Of Report ---

HRMS spectra of 1j

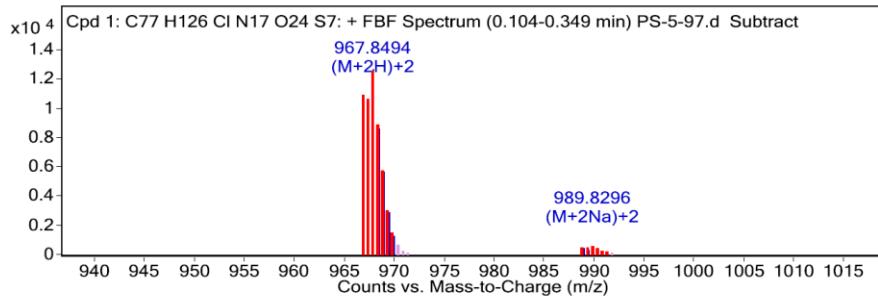


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
690.283	1	269786.13	C37H43N3O8S	(M+H)+
691.2856	1	112971.25	C37H43N3O8S	(M+H)+
692.2854	1	37546.63	C37H43N3O8S	(M+H)+
710.3115	1	1966.81	C37H43N3O8S	(M+K)+[-H2O]
712.2636	1	22394.12	C37H43N3O8S	(M+Na)+
728.2359	1	6018.28	C37H43N3O8S	(M+K)+
1379.5569	1	38415.02	C37H43N3O8S	(2M+H)+
1399.5766	1	806.01	C37H43N3O8S	(2M+K)+[-H2O]
1401.5359	1	5150.48	C37H43N3O8S	(2M+Na)+
1417.511	1	947.69	C37H43N3O8S	(2M+K)+

--- End Of Report ---

HRMS spectra of 1

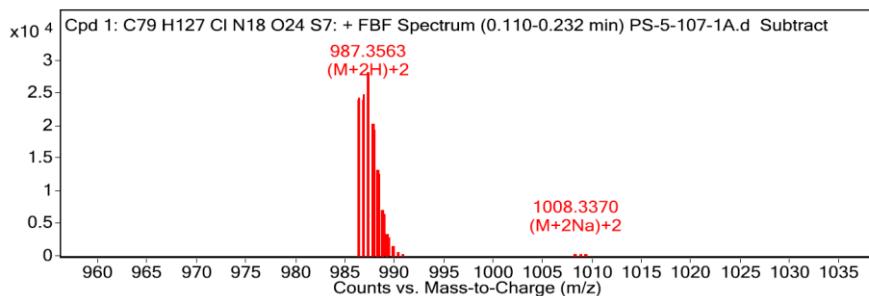


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
966.849	2	10970.96	C77H128ClN17O24S7	(M+2H)+2
967.3504	2	10581.76	C77H128ClN17O24S7	(M+2H)+2
967.8494	2	12732.34	C77H128ClN17O24S7	(M+2H)+2
968.3496	2	8657.85	C77H128ClN17O24S7	(M+2H)+2
968.849	2	5717.75	C77H128ClN17O24S7	(M+2H)+2
969.3486	2	2945.85	C77H128ClN17O24S7	(M+2H)+2
969.8485	2	1355.23	C77H128ClN17O24S7	(M+2H)+2
988.8291	2	502.26	C77H126ClN17Na2O24S7	(M+2Na)+2
989.3307	2	413.52	C77H126ClN17Na2O24S7	(M+2Na)+2
989.8296	2	561.39	C77H126ClN17Na2O24S7	(M+2Na)+2

--- End Of Report ---

HRMS spectra of 2

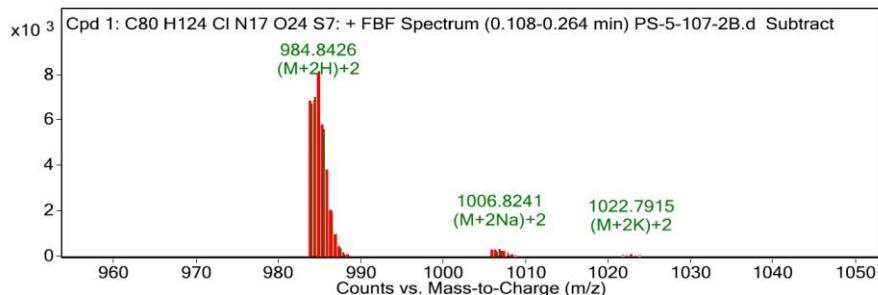


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
986.3557	2	24358.23	C ₇₉ H ₁₂₉ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
986.8571	2	24820.71	C ₇₉ H ₁₂₉ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
987.3563	2	28260.01	C ₇₉ H ₁₂₉ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
987.8564	2	19501.39	C ₇₉ H ₁₂₉ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
988.3555	2	12751.56	C ₇₉ H ₁₂₉ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
988.8556	2	6632.99	C ₇₉ H ₁₂₉ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
989.3553	2	3010.12	C ₇₉ H ₁₂₉ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
989.8553	2	1177.02	C ₇₉ H ₁₂₉ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
990.3543	2	481.65	C ₇₉ H ₁₂₉ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
1008.337	2	140.05	C ₇₉ H ₁₂₇ ClN ₁₈ Na ₂ O ₂₄ S ₇	(M+2Na) ⁺²

--- End Of Report ---

HRMS spectra of 3

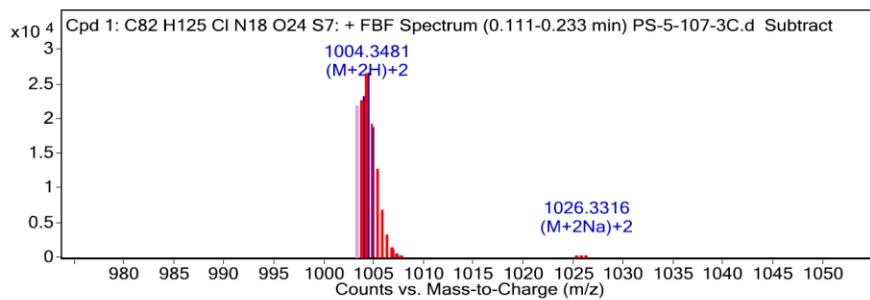


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
983.8422	2	6740.5	C ₈₀ H ₁₂₆ ClN ₁₇ O ₂₄ S ₇	(M+2H) ⁺²
984.3435	2	7053.95	C ₈₀ H ₁₂₆ ClN ₁₇ O ₂₄ S ₇	(M+2H) ⁺²
984.8426	2	8184.98	C ₈₀ H ₁₂₆ ClN ₁₇ O ₂₄ S ₇	(M+2H) ⁺²
985.3428	2	5613.13	C ₈₀ H ₁₂₆ ClN ₁₇ O ₂₄ S ₇	(M+2H) ⁺²
985.8425	2	3663.93	C ₈₀ H ₁₂₆ ClN ₁₇ O ₂₄ S ₇	(M+2H) ⁺²
986.3428	2	1977.39	C ₈₀ H ₁₂₆ ClN ₁₇ O ₂₄ S ₇	(M+2H) ⁺²
986.8415	2	949.15	C ₈₀ H ₁₂₆ ClN ₁₇ O ₂₄ S ₇	(M+2H) ⁺²
987.3431	2	390.48	C ₈₀ H ₁₂₆ ClN ₁₇ O ₂₄ S ₇	(M+2H) ⁺²
1005.8222	2	270.46	C ₈₀ H ₁₂₄ ClN ₁₇ Na ₂ O ₂₄ S ₇	(M+2Na) ⁺²
1022.7915	2	65.45	C ₈₀ H ₁₂₄ ClK ₂ N ₁₇ O ₂₄ S ₇	(M+2K) ⁺²

--- End Of Report ---

HRMS spectra of 4

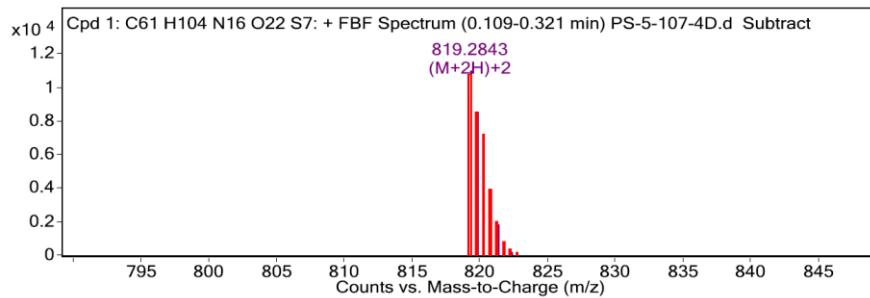


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
1003.8485	2	23340.62	C ₈₂ H ₁₂₇ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
1004.3481	2	26718.61	C ₈₂ H ₁₂₇ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
1004.8482	2	18850.31	C ₈₂ H ₁₂₇ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
1005.3475	2	12182.81	C ₈₂ H ₁₂₇ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
1005.8475	2	6510.62	C ₈₂ H ₁₂₇ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
1006.3471	2	2962.8	C ₈₂ H ₁₂₇ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
1006.8467	2	1308.35	C ₈₂ H ₁₂₇ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
1007.3467	2	481.44	C ₈₂ H ₁₂₇ ClN ₁₈ O ₂₄ S ₇	(M+2H) ⁺²
1025.3265	2	117.69	C ₈₂ H ₁₂₅ ClN ₁₈ Na ₂ O ₂₄ S ₇	(M+2Na) ⁺²
1026.3316	2	145.25	C ₈₂ H ₁₂₅ ClN ₁₈ Na ₂ O ₂₄ S ₇	(M+2Na) ⁺²

--- End Of Report ---

HRMS spectra of 5

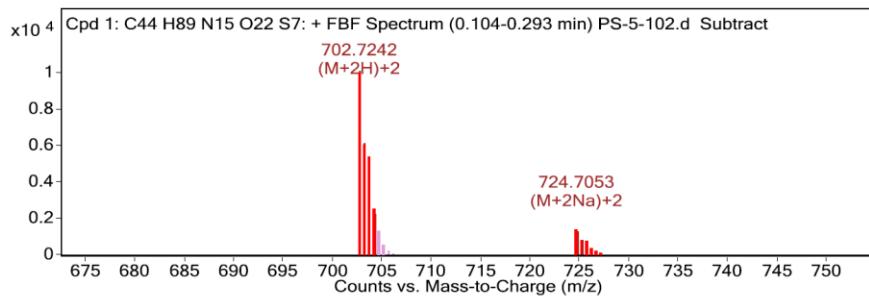


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
819.2843	2	11062.34	C ₆₁ H ₁₀₆ N ₁₆ O ₂₂ S ₇	(M+2H) ⁺²
819.7855	2	8579.07	C ₆₁ H ₁₀₆ N ₁₆ O ₂₂ S ₇	(M+2H) ⁺²
820.2846	2	6986.26	C ₆₁ H ₁₀₆ N ₁₆ O ₂₂ S ₇	(M+2H) ⁺²
820.7845	2	3902.19	C ₆₁ H ₁₀₆ N ₁₆ O ₂₂ S ₇	(M+2H) ⁺²
821.2841	2	1947.75	C ₆₁ H ₁₀₆ N ₁₆ O ₂₂ S ₇	(M+2H) ⁺²
821.7839	2	760.61	C ₆₁ H ₁₀₆ N ₁₆ O ₂₂ S ₇	(M+2H) ⁺²
822.2832	2	284.88	C ₆₁ H ₁₀₆ N ₁₆ O ₂₂ S ₇	(M+2H) ⁺²
822.782	2	79.1	C ₆₁ H ₁₀₆ N ₁₆ O ₂₂ S ₇	(M+2H) ⁺²

--- End Of Report ---

HRMS spectra of 6

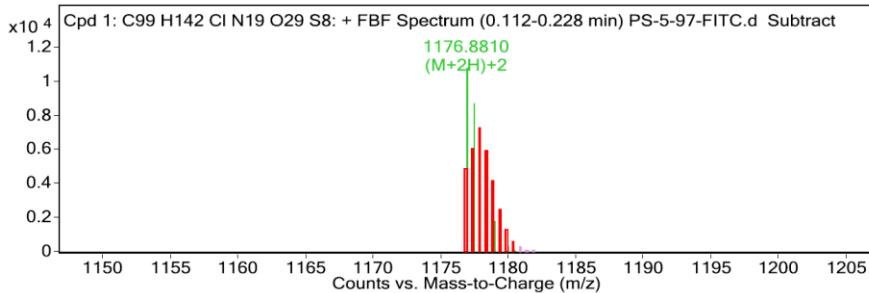


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
702.7242	2	10162.73	C44H91N15O22S7	(M+2H) ⁺²
703.2252	2	6169.12	C44H91N15O22S7	(M+2H) ⁺²
703.7242	2	5215.91	C44H91N15O22S7	(M+2H) ⁺²
704.2244	2	2309.95	C44H91N15O22S7	(M+2H) ⁺²
724.7053	2	1342.12	C44H89N15Na2O22S7	(M+2Na) ⁺²
725.2065	2	849.6	C44H89N15Na2O22S7	(M+2Na) ⁺²
725.7046	2	728.89	C44H89N15Na2O22S7	(M+2Na) ⁺²
726.2048	2	294.39	C44H89N15Na2O22S7	(M+2Na) ⁺²
726.7051	2	127.69	C44H89N15Na2O22S7	(M+2Na) ⁺²
727.2028	2	29.15	C44H89N15Na2O22S7	(M+2Na) ⁺²

--- End Of Report ---

HRMS spectra of 1-FITC

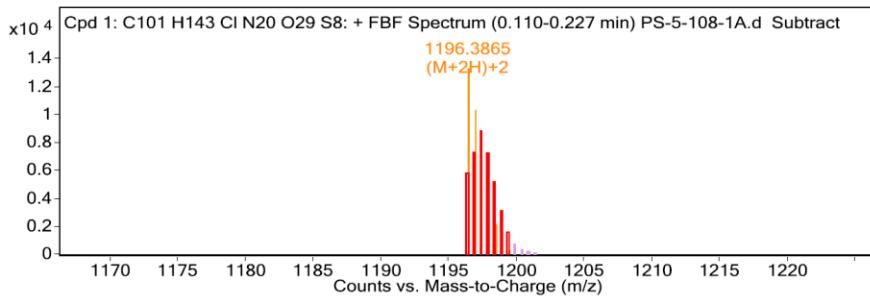


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
1176.881	2	10861.65	C99H144ClN19O29S8	(M+2H) ⁺²
1177.3815	2	8735.57	C99H144ClN19O29S8	(M+2H) ⁺²
1177.8808	2	6133.46	C99H144ClN19O29S8	(M+2H) ⁺²
1178.3807	2	3570.78	C99H144ClN19O29S8	(M+2H) ⁺²
1178.8809	2	1910.52	C99H144ClN19O29S8	(M+2H) ⁺²
1179.3798	2	856.76	C99H144ClN19O29S8	(M+2H) ⁺²
1179.8785	2	366.92	C99H144ClN19O29S8	(M+2H) ⁺²
1180.3788	2	134.07	C99H144ClN19O29S8	(M+2H) ⁺²

--- End Of Report ---

HRMS spectra of 2-FITC

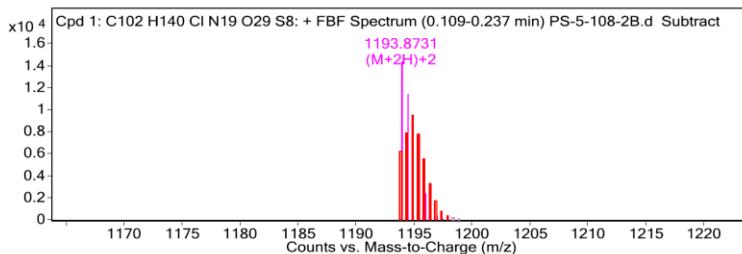


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
1196.3865	2	13316.92	C101H145ClN20O29S8	(M+2H)+2
1196.8869	2	10396.94	C101H145ClN20O29S8	(M+2H)+2
1197.387	2	7434.07	C101H145ClN20O29S8	(M+2H)+2
1197.8868	2	4156.46	C101H145ClN20O29S8	(M+2H)+2
1198.386	2	2234.87	C101H145ClN20O29S8	(M+2H)+2
1198.885	2	1041.17	C101H145ClN20O29S8	(M+2H)+2
1199.3857	2	432.68	C101H145ClN20O29S8	(M+2H)+2

--- End Of Report ---

HRMS spectra of 3-FITC

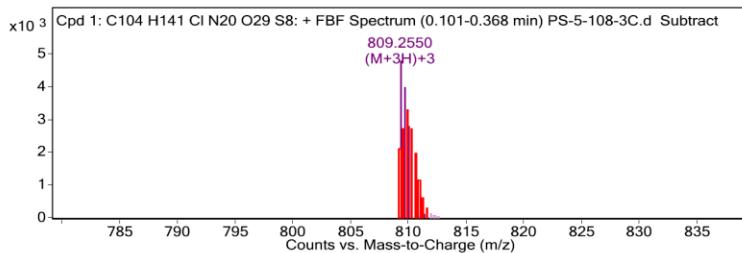


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
1193.8731	2	14409.95	C102H142ClN19O29S8	(M+2H)+2
1194.3732	2	11427.2	C102H142ClN19O29S8	(M+2H)+2
1194.8729	2	7915.61	C102H142ClN19O29S8	(M+2H)+2
1195.3725	2	4890.26	C102H142ClN19O29S8	(M+2H)+2
1195.8723	2	2459.77	C102H142ClN19O29S8	(M+2H)+2
1196.372	2	1077.29	C102H142ClN19O29S8	(M+2H)+2
1196.8731	2	524.97	C102H142ClN19O29S8	(M+2H)+2
1197.3721	2	153.16	C102H142ClN19O29S8	(M+2H)+2
1197.8741	2	27.78	C102H142ClN19O29S8	(M+2H)+2

--- End Of Report ---

HRMS spectra of 4-FITC

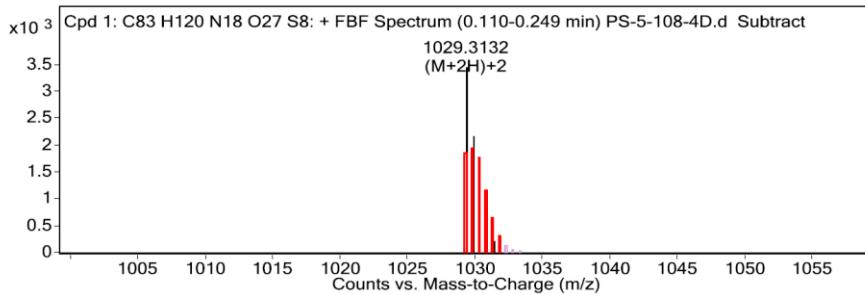


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
809.255	3	4817.22	C104H144ClN20O29S8	(M+3H) ⁺³
809.5885	3	4011.91	C104H144ClN20O29S8	(M+3H) ⁺³
809.9217	3	2813.06	C104H144ClN20O29S8	(M+3H) ⁺³
810.2548	3	1670.02	C104H144ClN20O29S8	(M+3H) ⁺³
810.5885	3	876.2	C104H144ClN20O29S8	(M+3H) ⁺³
810.9207	3	422.02	C104H144ClN20O29S8	(M+3H) ⁺³
811.2533	3	143.89	C104H144ClN20O29S8	(M+3H) ⁺³
811.5848	3	54.35	C104H144ClN20O29S8	(M+3H) ⁺³

--- End Of Report ---

HRMS spectra of 5-FITC

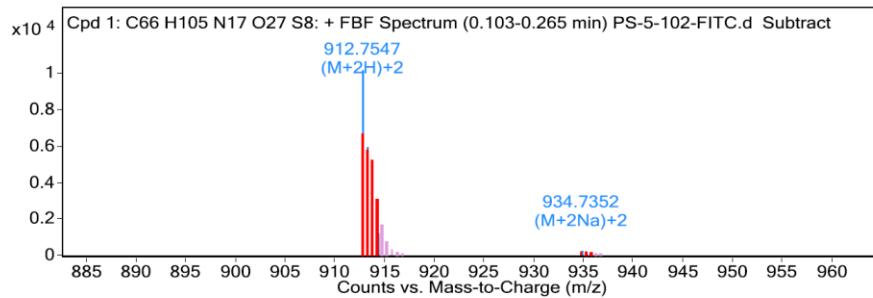


MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
1029.3132	2	3449.52	C83H122N18O27S8	(M+2H) ⁺²
1029.8132	2	2189.28	C83H122N18O27S8	(M+2H) ⁺²
1030.3134	2	1153.83	C83H122N18O27S8	(M+2H) ⁺²
1030.8125	2	554.84	C83H122N18O27S8	(M+2H) ⁺²
1031.3132	2	232.76	C83H122N18O27S8	(M+2H) ⁺²
1031.809	2	74.75	C83H122N18O27S8	(M+2H) ⁺²

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HRMS spectra of 6-FITC

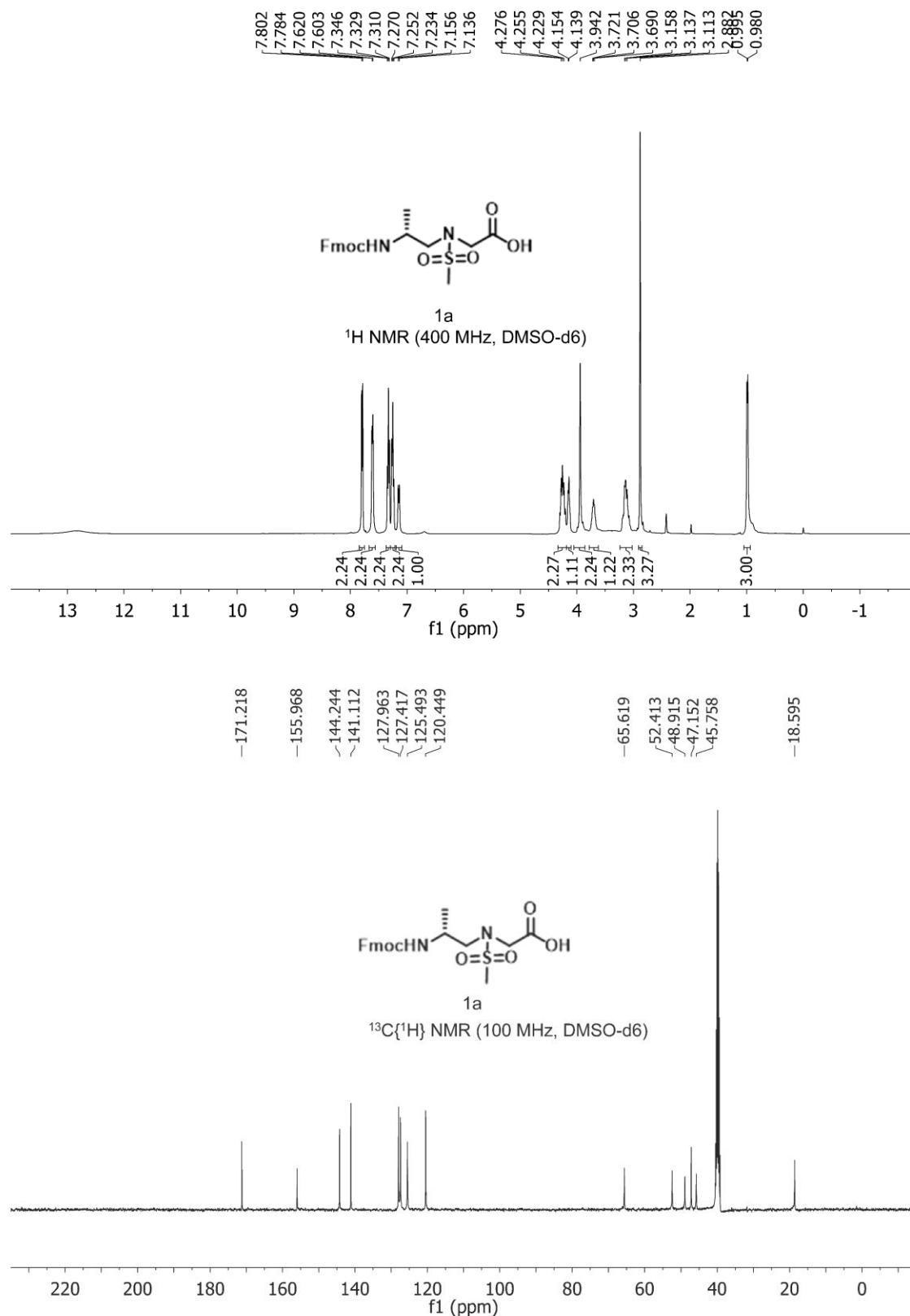


MS Spectrum Peak List

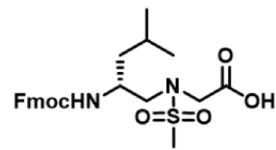
m/z	z	Abund	Formula	Ion
912.7547	2	10222.49	C ₆₆ H ₁₀₇ N ₁₇ O ₂₇ S ₈	(M+2H) ⁺²
913.2544	2	5983.9	C ₆₆ H ₁₀₇ N ₁₇ O ₂₇ S ₈	(M+2H) ⁺²
913.7537	2	3049.32	C ₆₆ H ₁₀₇ N ₁₇ O ₂₇ S ₈	(M+2H) ⁺²
914.2537	2	1313.44	C ₆₆ H ₁₀₇ N ₁₇ O ₂₇ S ₈	(M+2H) ⁺²
934.7352	2	278.5	C ₆₆ H ₁₀₅ N ₁₇ Na ₂ O ₂₇ S ₈	(M+2Na) ⁺²
935.2348	2	139.58	C ₆₆ H ₁₀₅ N ₁₇ Na ₂ O ₂₇ S ₈	(M+2Na) ⁺²
935.7305	2	51.55	C ₆₆ H ₁₀₅ N ₁₇ Na ₂ O ₂₇ S ₈	(M+2Na) ⁺²

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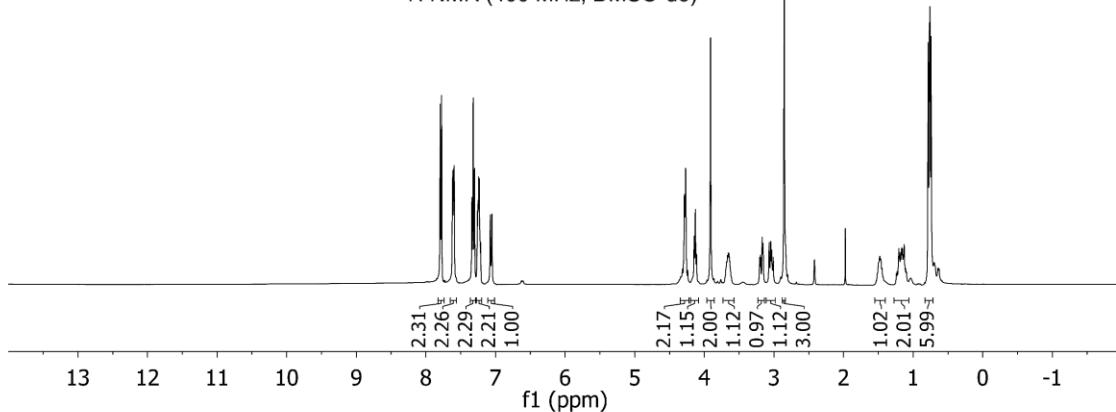
6. The ^1H and ^{13}C NMR Spectra of D-Sulfono- γ -AApeptide Building Blocks 1a-j



7.794
7.775
7.612
7.598
7.594
7.339
7.321
7.302
7.260
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7.242
7.232
7.225
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1.195
1.182
1.164
1.152
1.140
1.129
0.785
0.768
0.758
0.742



¹H NMR (400 MHz, DMSO-d6)



-171.144

-156.268

144.272

141.127

127.920

127.365

125.480

120.411

65.434

51.862

48.772

48.107

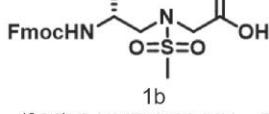
47.242

41.223

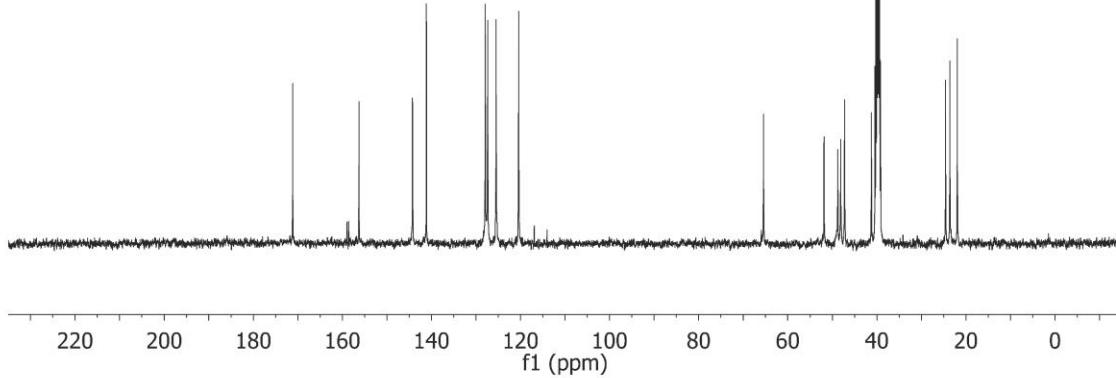
24.571

23.601

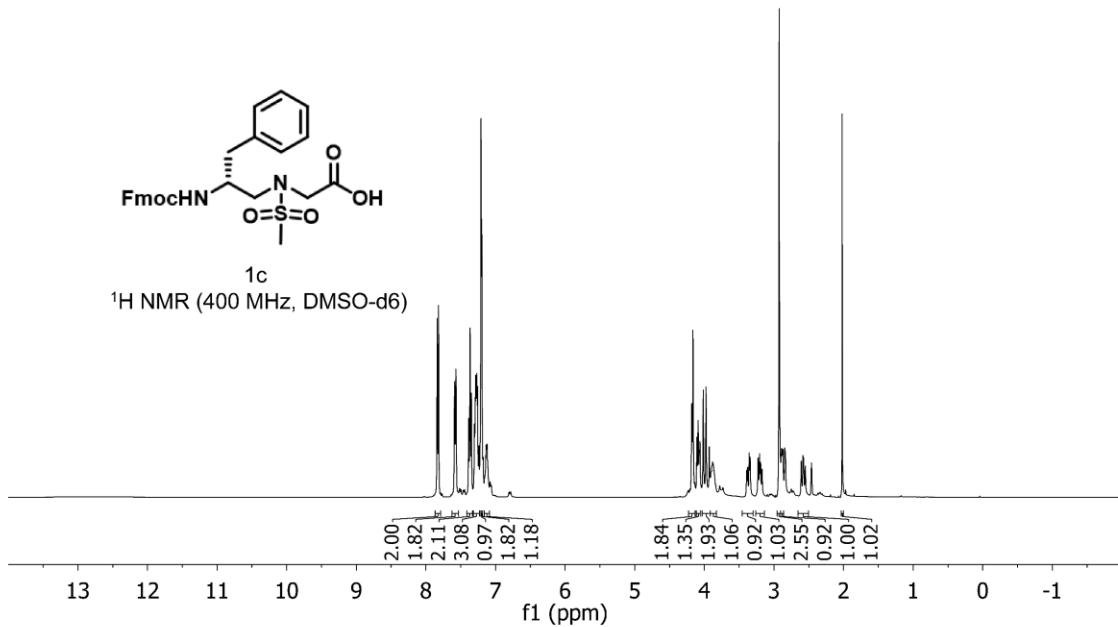
21.944



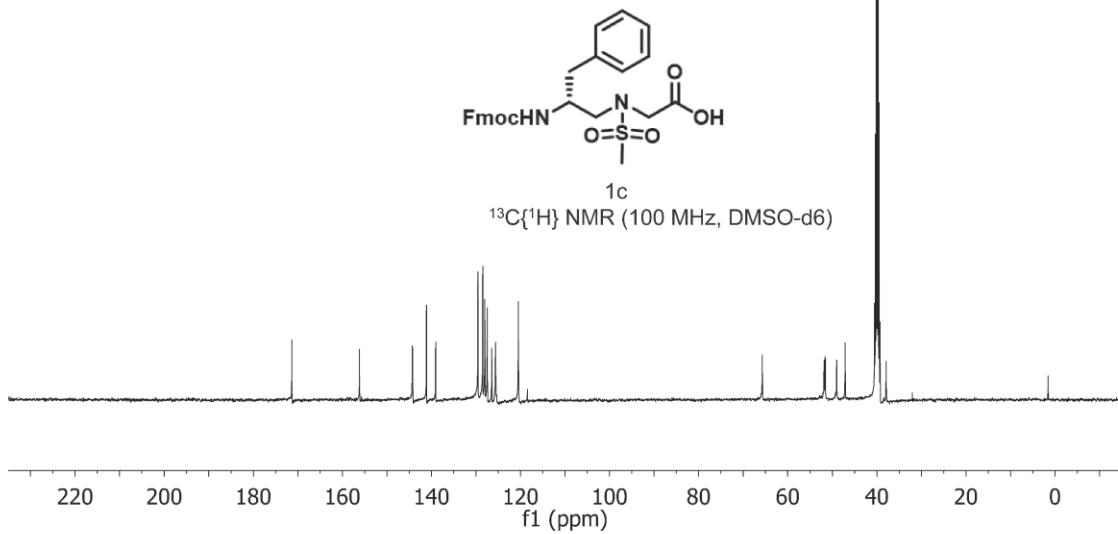
¹³C{¹H} NMR (100 MHz, DMSO-d6)



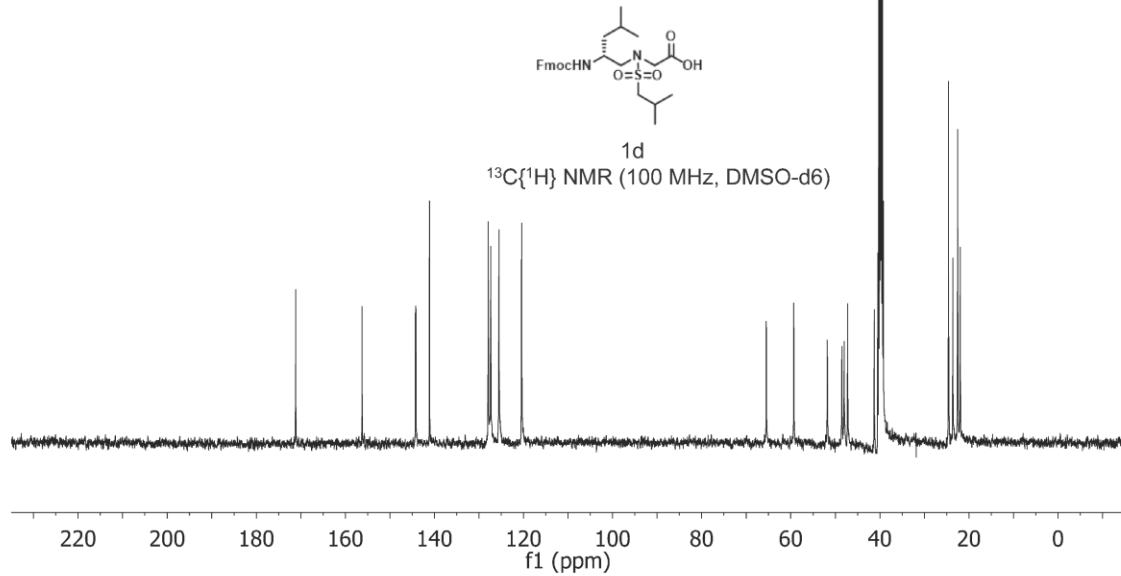
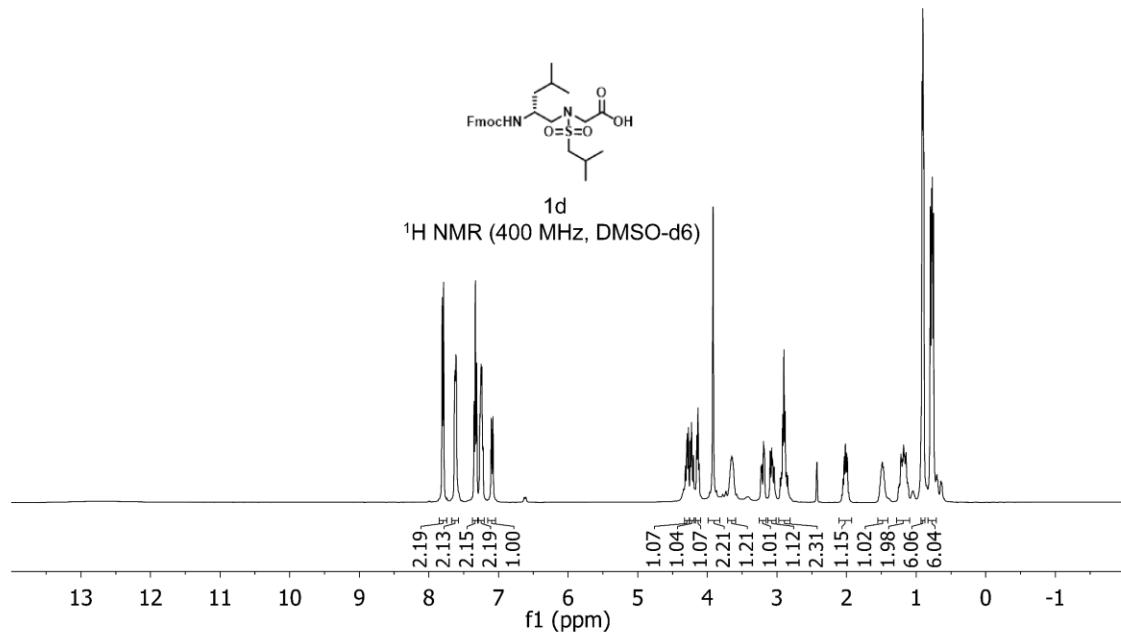
7.838
 7.819
 7.587
 7.569
 7.385
 7.366
 7.348
 7.302
 7.292
 7.284
 7.277
 7.269
 7.259
 7.209
 7.200
 7.141
 7.134
 7.126
 7.120
 7.113
 4.180
 4.163
 4.107
 4.090
 4.071
 4.063
 4.016
 3.976
 3.929
 3.378
 3.356
 3.342
 3.227
 3.206
 3.191
 2.925
 2.892
 2.880
 2.870
 2.608
 2.583
 2.574
 2.549
 2.019

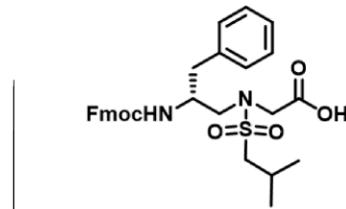
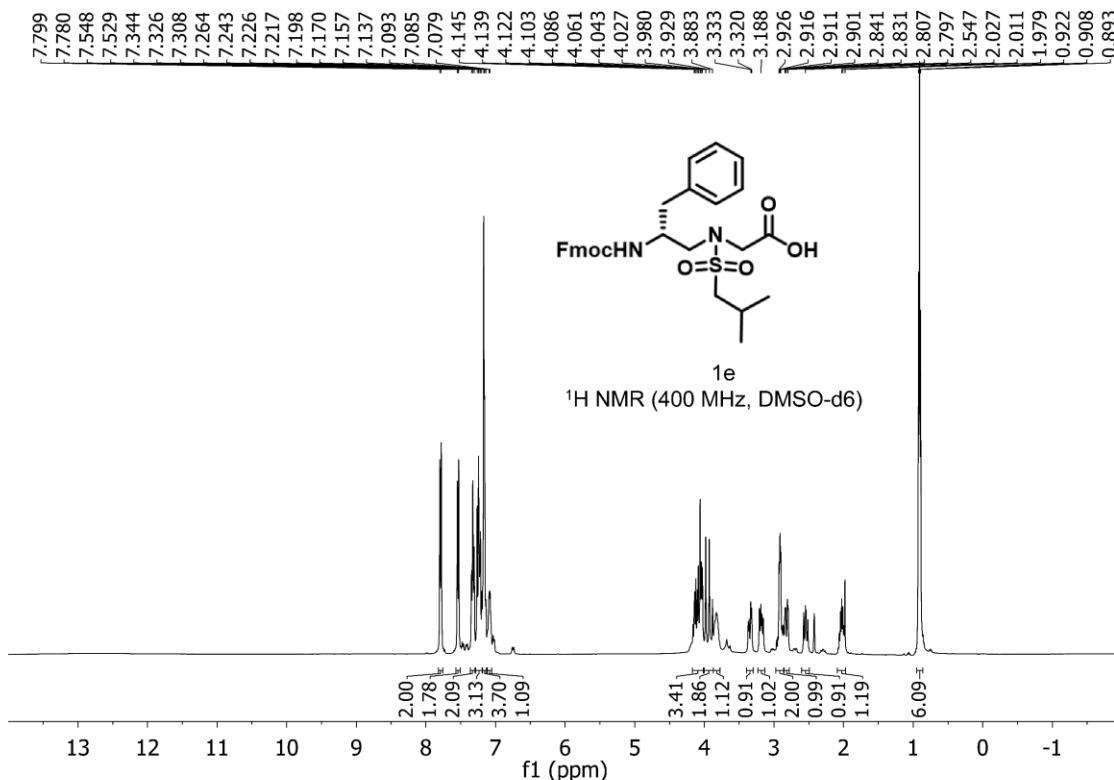


-171.337
 -156.145
 -144.278
 -141.117
 -141.194
 -139.052
 -129.561
 -128.475
 -128.014
 -127.456
 -126.446
 -125.617
 -120.485

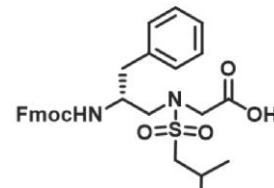
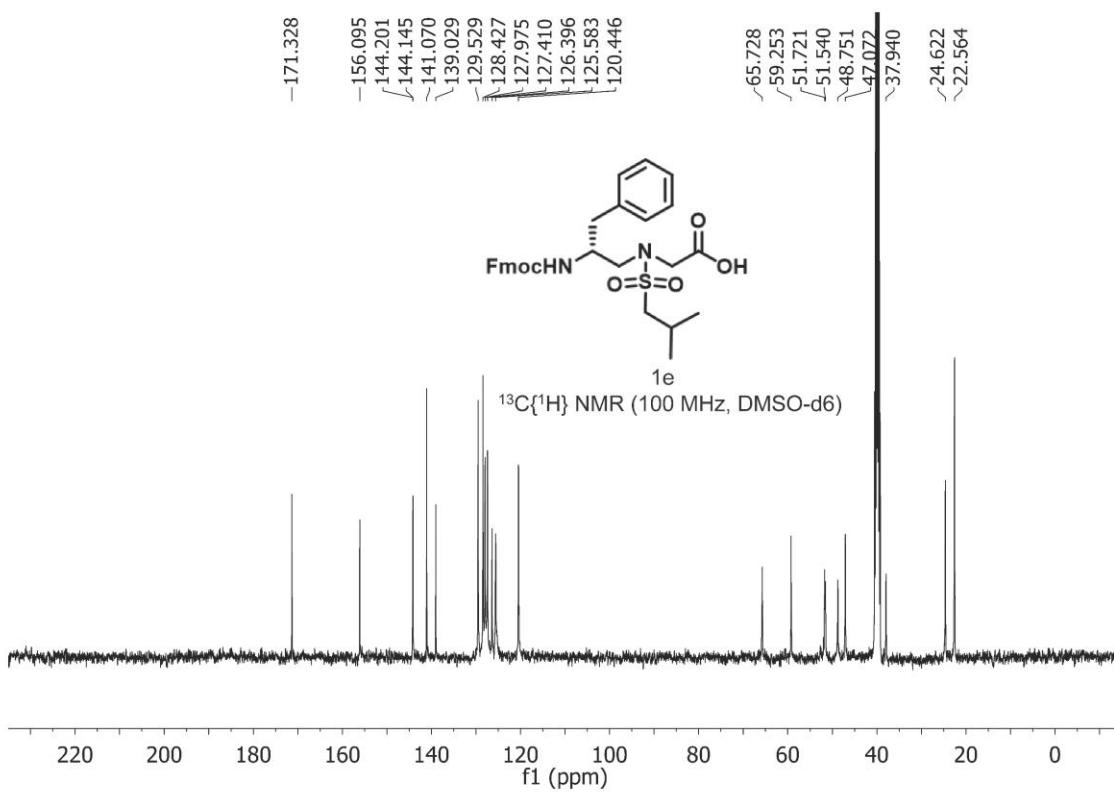


7.808
 7.790
 7.624
 7.612
 7.351
 7.333
 7.314
 7.267
 7.251
 7.240
 7.222
 7.103
 7.080
 4.289
 4.271
 4.244
 4.227
 4.200
 4.151
 4.134
 3.918
 3.650
 3.640
 3.192
 3.181
 3.097
 3.075
 3.061
 2.917
 2.900
 2.882
 2.033
 2.017
 2.000
 1.987
 1.218
 1.194
 1.180
 1.168
 1.144
 0.914
 0.903
 0.889
 0.795
 0.779
 0.767
 0.751





¹H NMR (400 MHz, DMSO-d₆)



¹³C{¹H} NMR (100 MHz, DMSO-d₆)

