

Anion Binding Properties of a Cyclic Pseudoheptapeptide Containing 1,5-Disubstituted 1,2,3-Triazole Subunits

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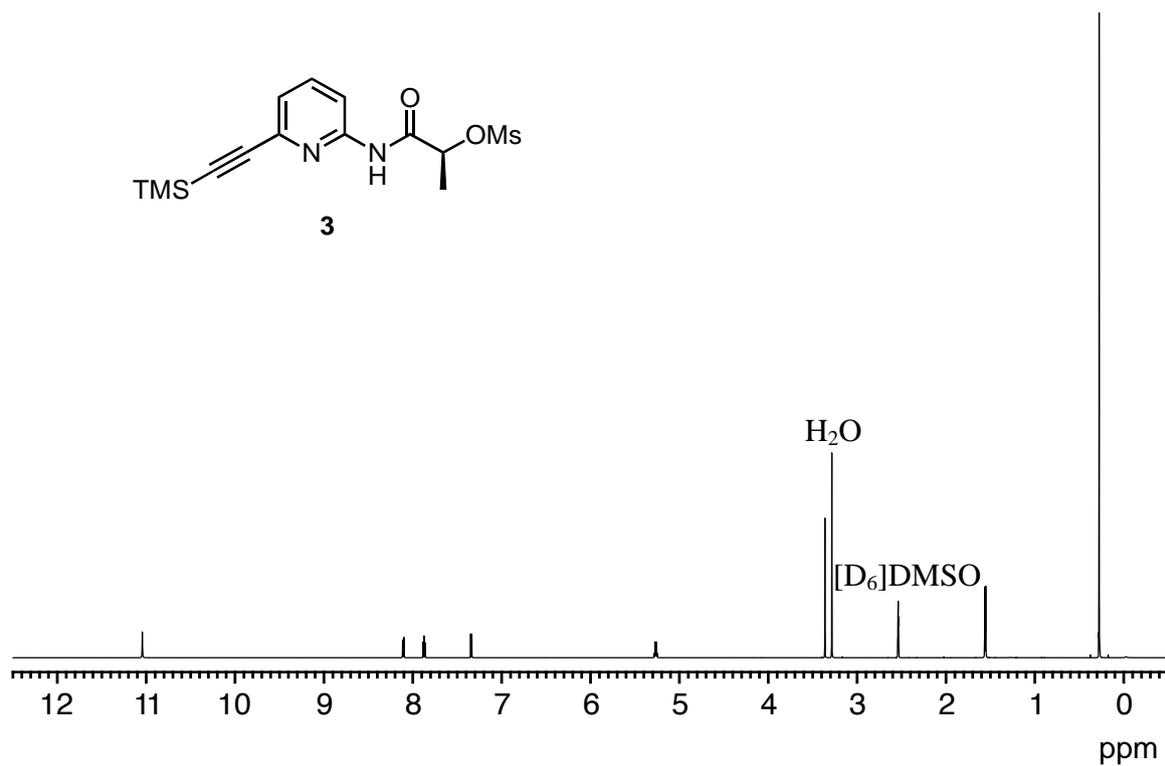
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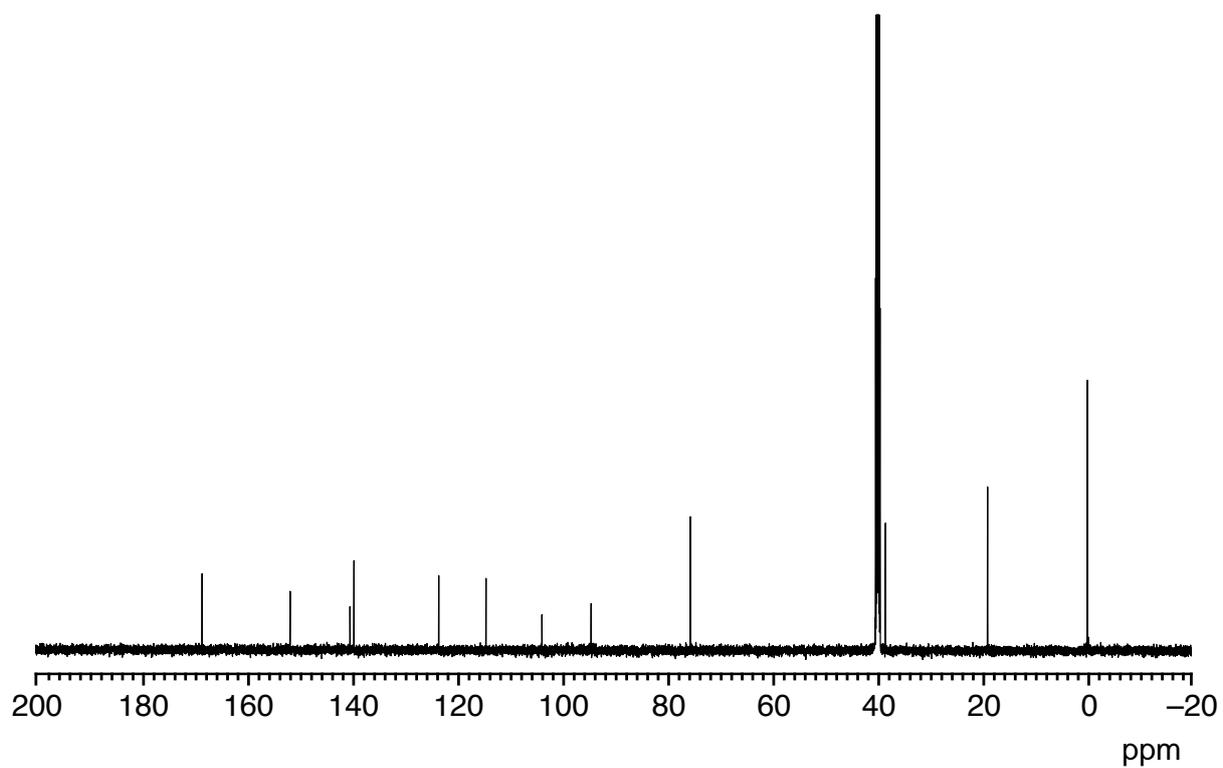
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^1H NMR Spectrum: **3** in $[\text{D}_6]\text{DMSO}$ (600 MHz, 25°C).



^{13}C NMR Spectrum: **3** in $[\text{D}_6]\text{DMSO}$ (151 MHz, 25°C).



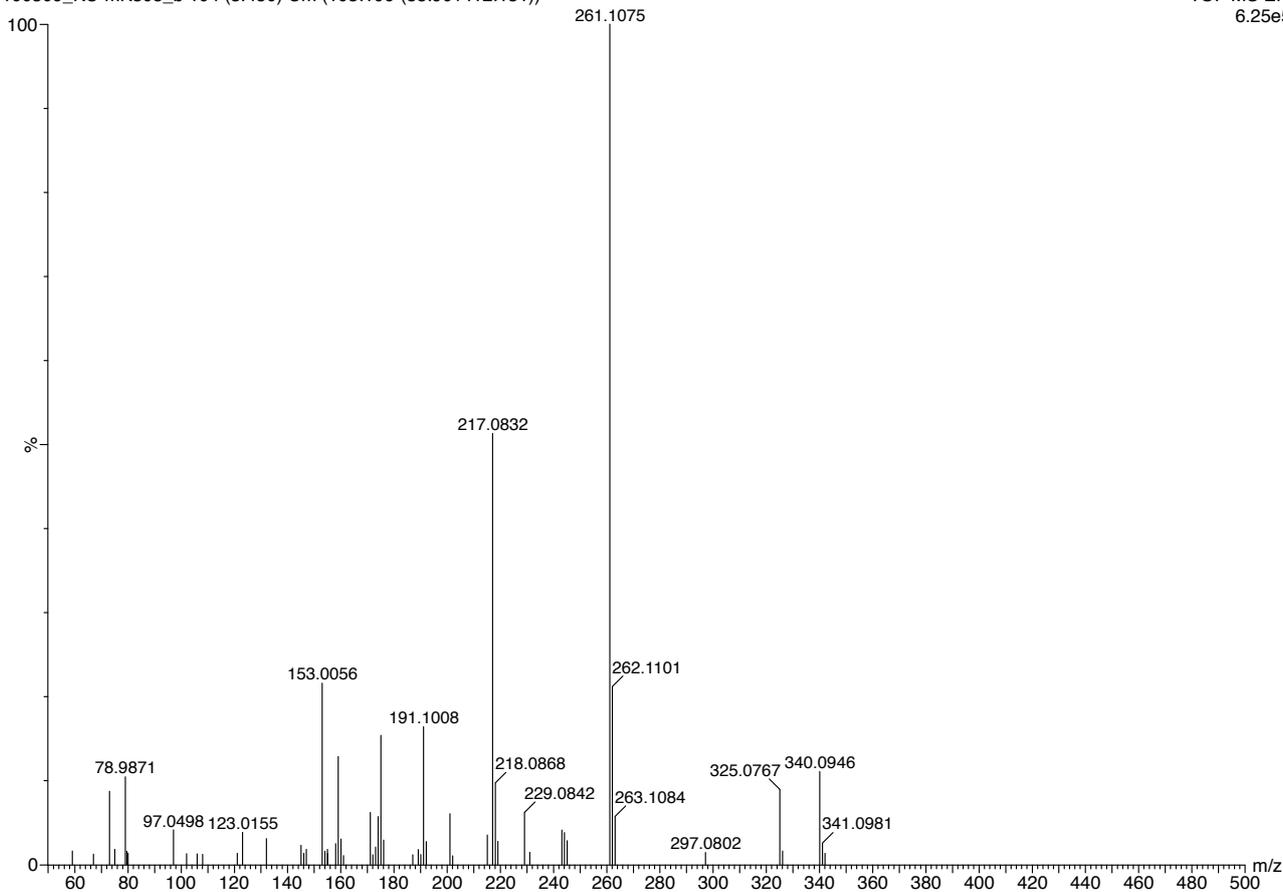
EI MS Spectrum: 3 (positive mode)

09-Mar-2010 11:10:42 EI_Feststoff

09-Mar-2010
GCT Premier CAB163

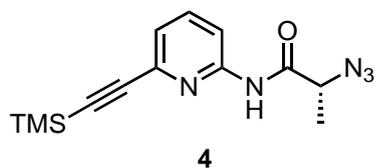
11:10:42
mk305, M 340.47, Ramp2
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6.25e5

100309_KU-mk305_b 104 (3.450) Cm (103:106-(85:90+112:131))

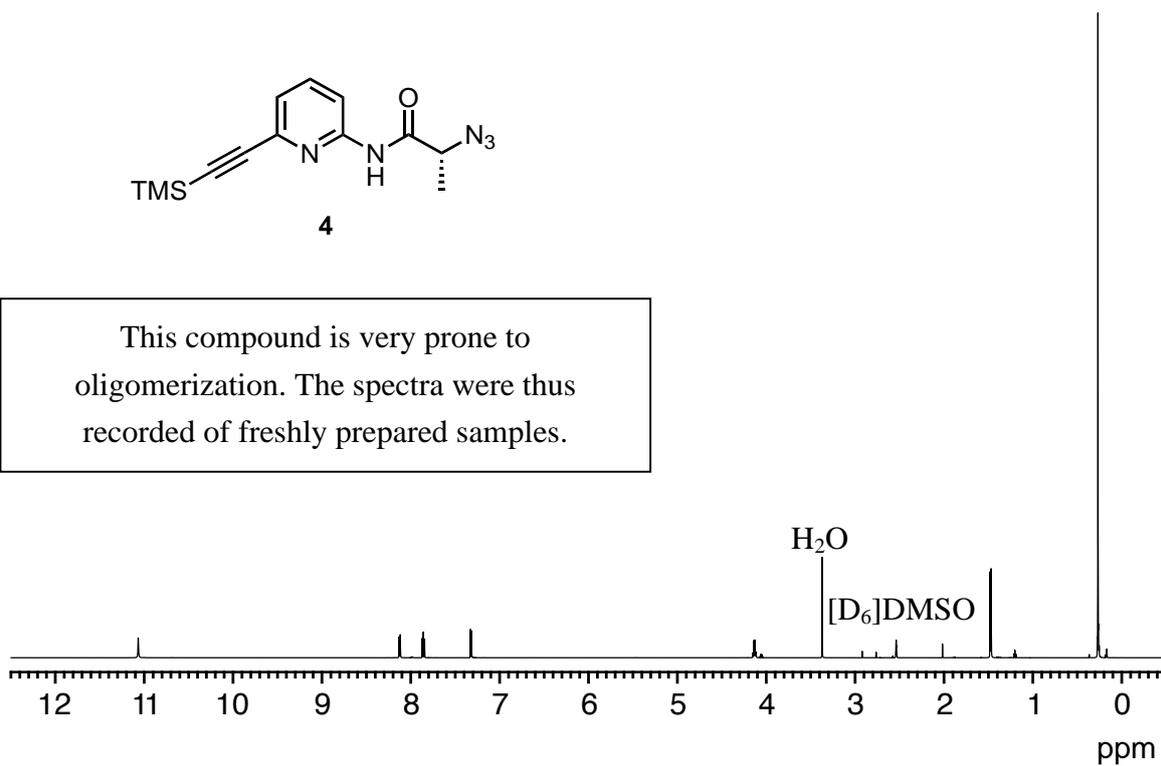


		<i>m/z calcd.</i>	<i>m/z exp.</i>
$3^+ - \text{CH}(\text{CH}_3)\text{OSO}_2\text{CH}_3$	$\text{C}_{14}\text{H}_{20}\text{N}_2\text{O}_4\text{SSi}^+ - \text{CH}(\text{CH}_3)\text{OSO}_2\text{CH}_3$	217.0797	217.0832
$3^+ - \text{SO}_2\text{CH}_3$	$\text{C}_{14}\text{H}_{20}\text{N}_2\text{O}_4\text{SSi}^+ - \text{SO}_2\text{CH}_3$	261.1059	261.1075
3^+	$\text{C}_{14}\text{H}_{20}\text{N}_2\text{O}_4\text{SSi}^+$	340.0913	340.0946

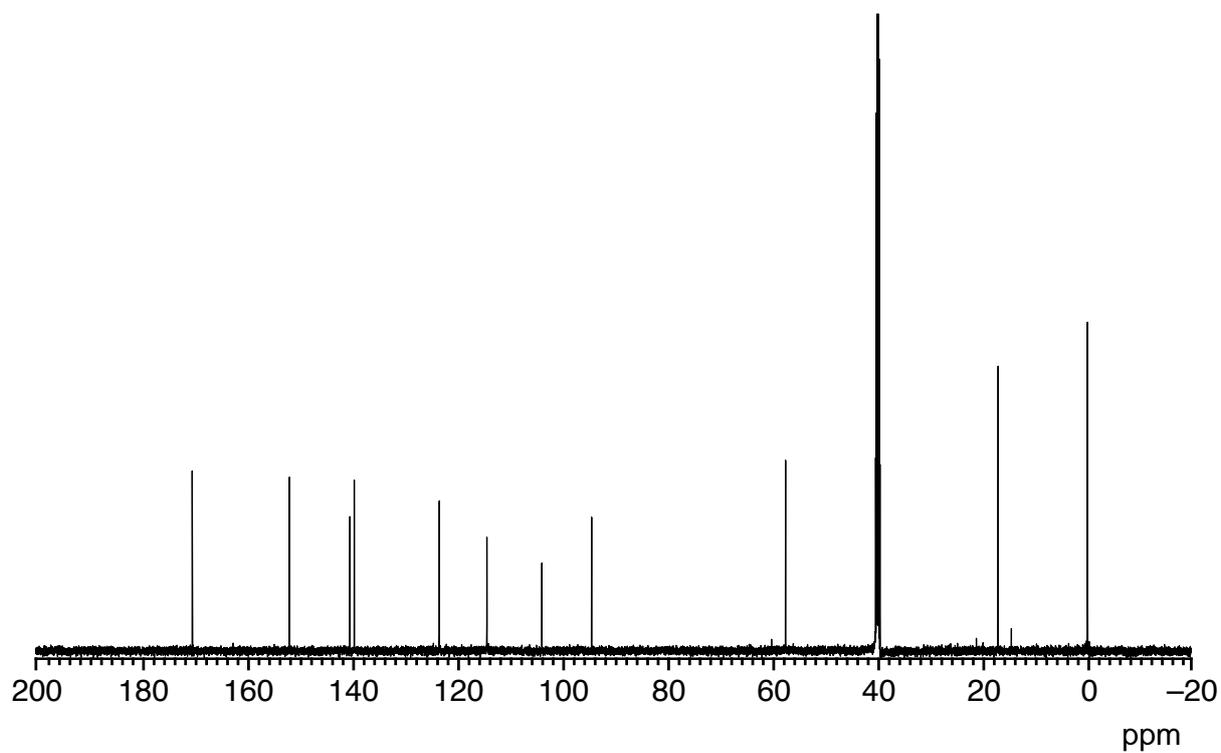
^1H NMR Spectrum: **4** in $[\text{D}_6]\text{DMSO}$ (600 MHz, 25°C).



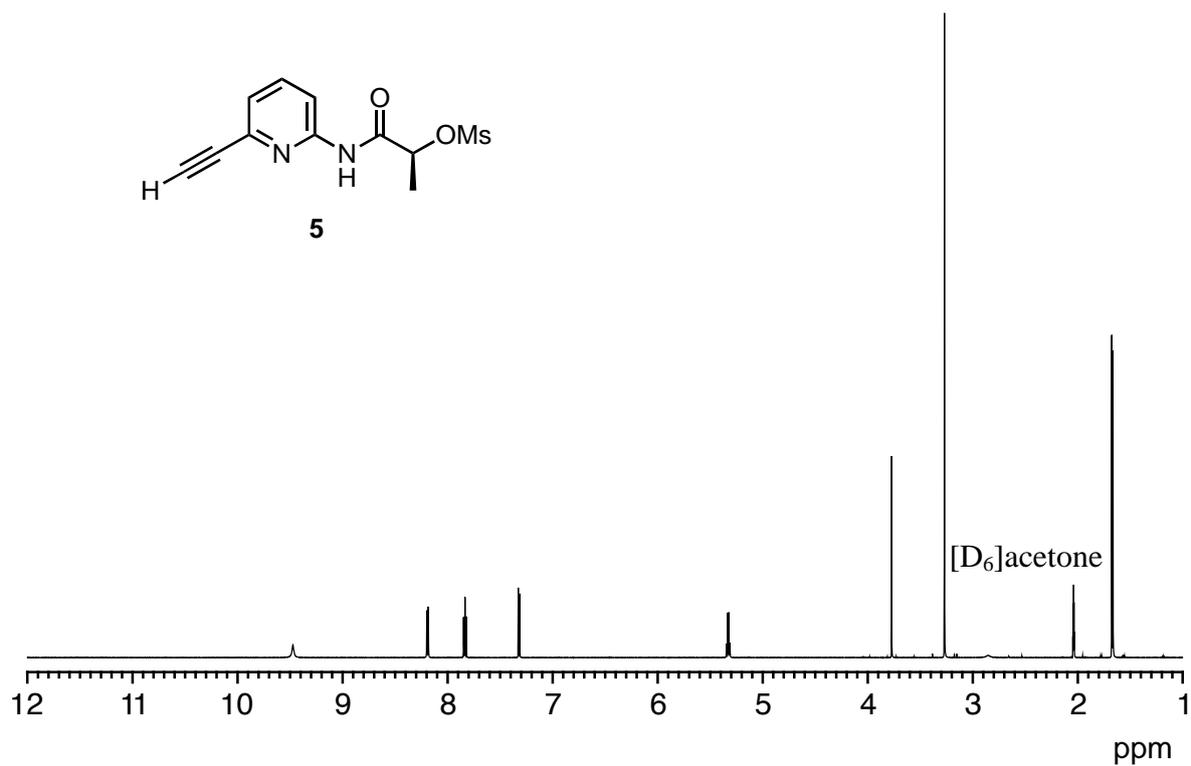
This compound is very prone to oligomerization. The spectra were thus recorded of freshly prepared samples.



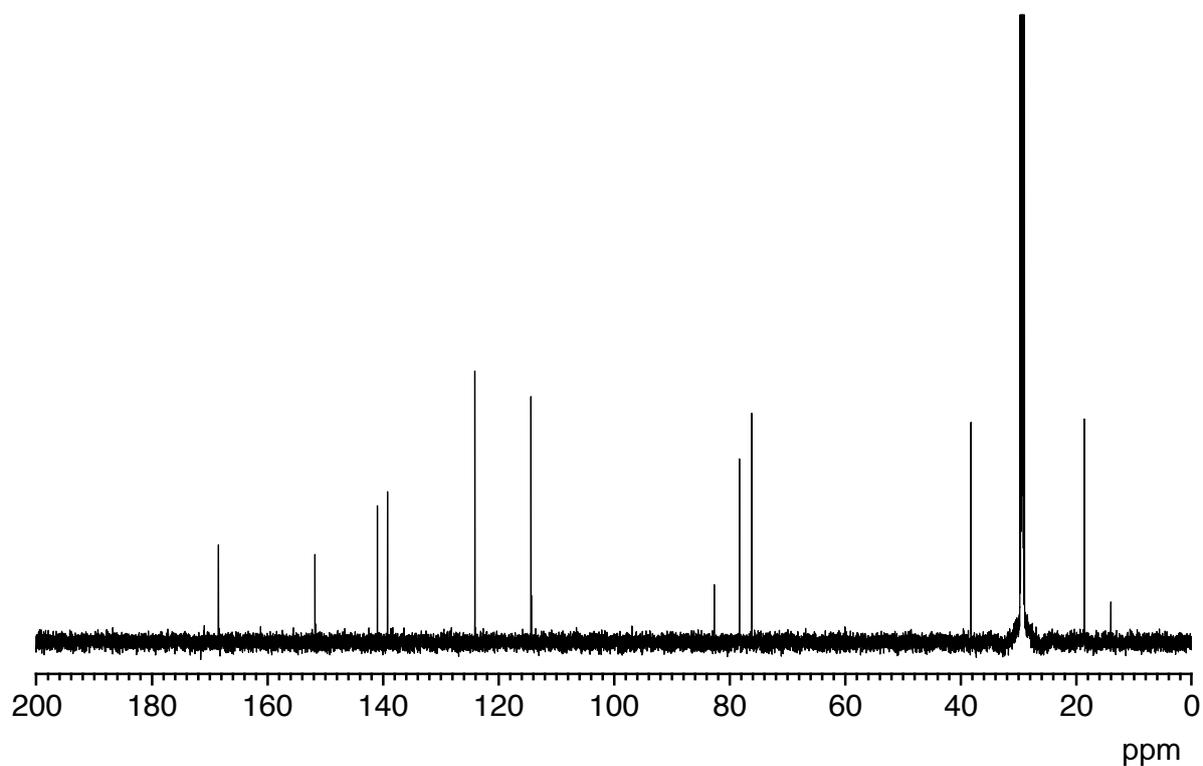
^{13}C NMR Spectrum: **4** in $[\text{D}_6]\text{DMSO}$ (151 MHz, 25°C).



^1H NMR Spectrum: **5** in $[\text{D}_6]$ acetone (600 MHz, 25°C).



^{13}C NMR Spectrum: **5** in $[\text{D}_6]$ acetone (151 MHz, 25°C).



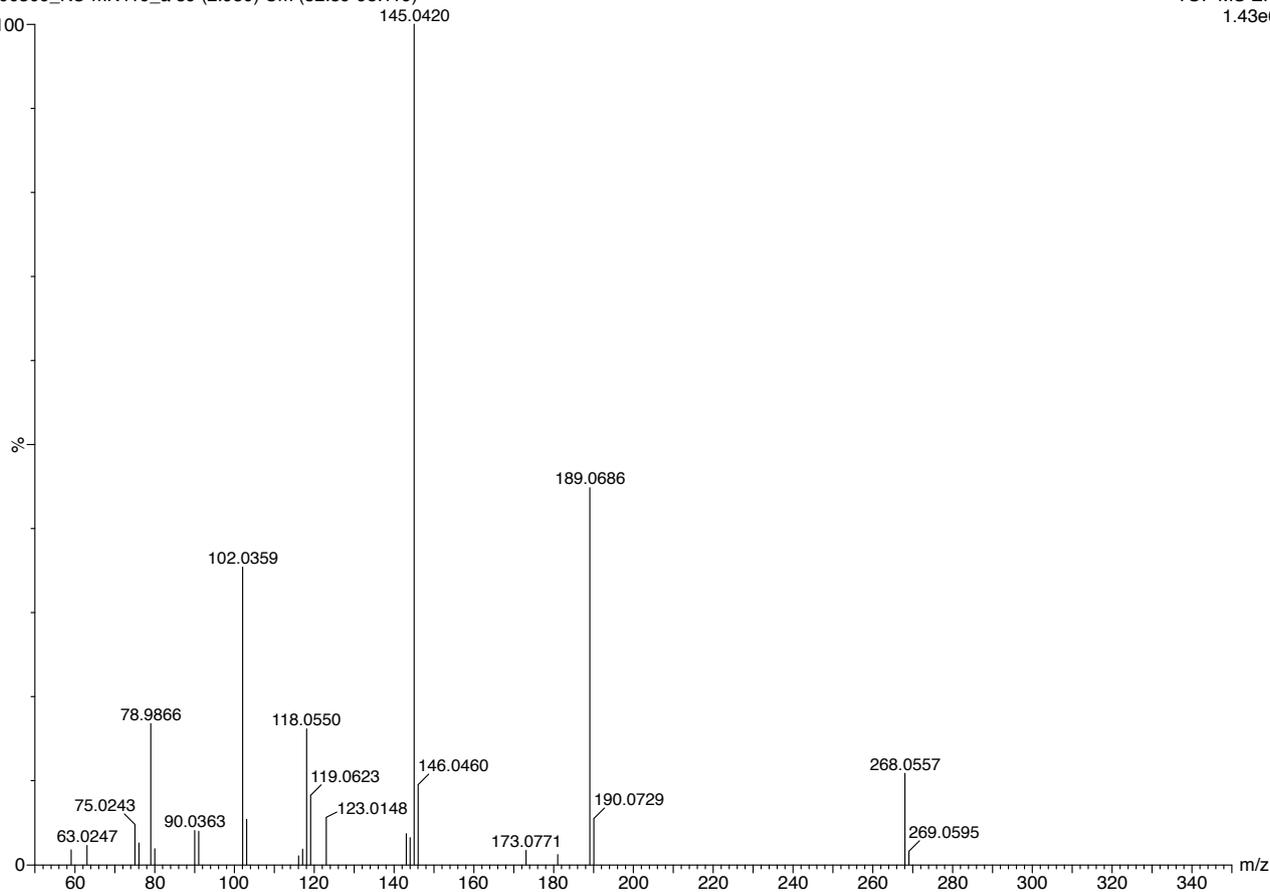
EI MS Spectrum: 5 (positive mode).

09-Mar-2010 12:06:28 EI_Feststoff

09-Mar-2010
GCT Premier CAB163

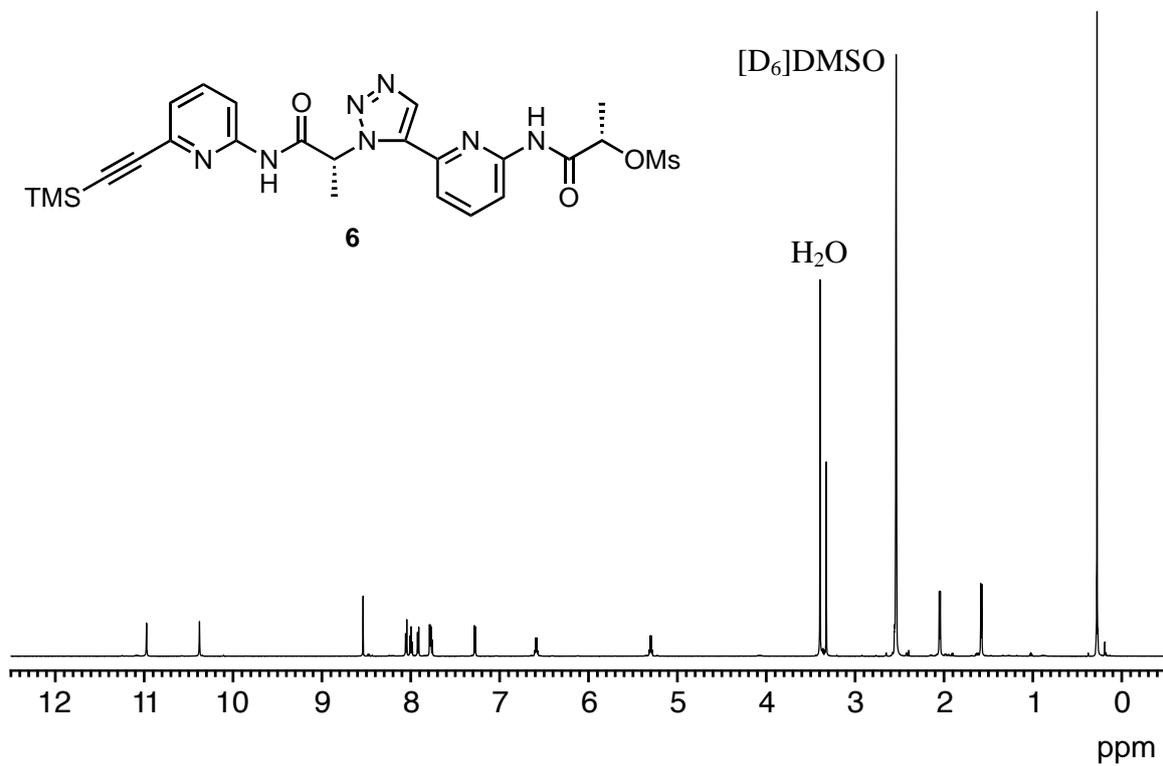
12:06:28
mk419, M 268.29, Ramp2
TOF MS EI+
1.43e6

100309_KU-mK419_a 89 (2.950) Cm (82:89-98:119)

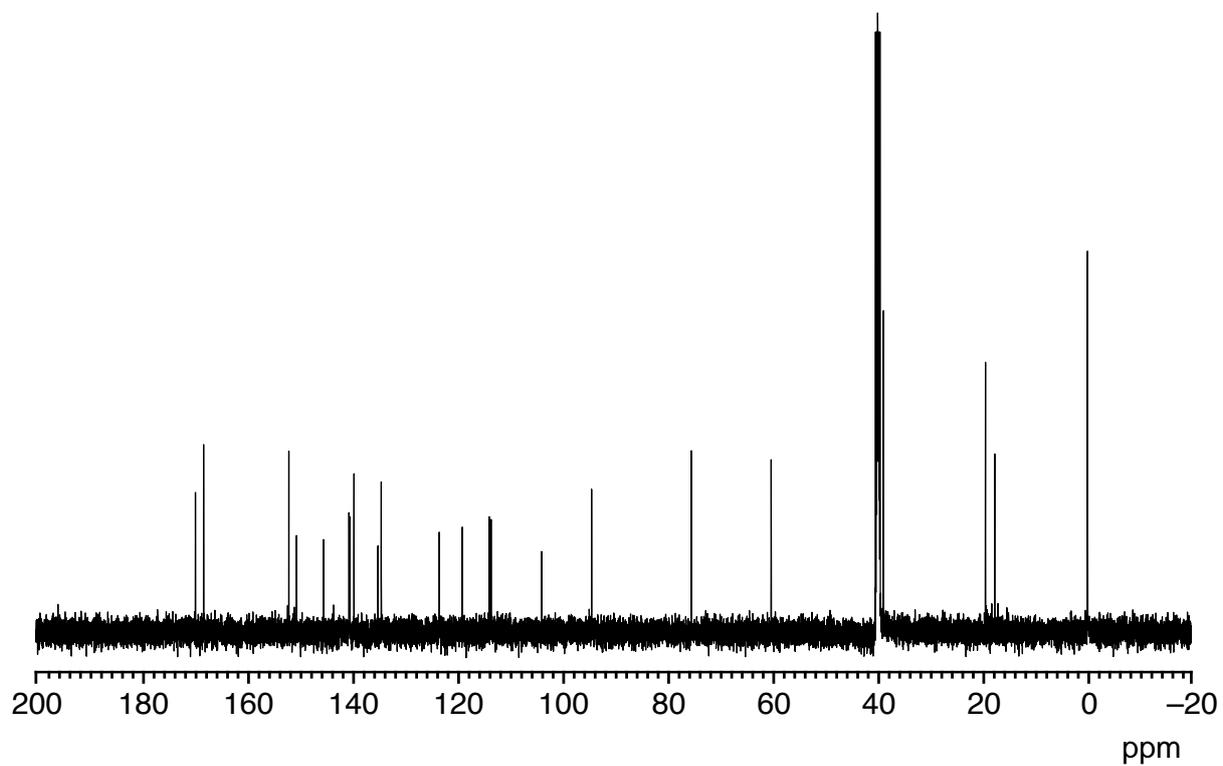


		<i>m/z calcd.</i>	<i>m/z exp.</i>
5^+ -CH(CH ₃)OSO ₂ CH ₃	C ₁₁ H ₁₂ N ₂ O ₄ S ⁺ -CH(CH ₃)OSO ₂ CH ₃	145.0402	145.0420
5^+ -SO ₂ CH ₃	C ₁₁ H ₁₂ N ₂ O ₄ S ⁺ -SO ₂ CH ₃	189.0664	189.0686
5^+	C ₁₁ H ₁₂ N ₂ O ₄ S ⁺	268.0518	268.0557

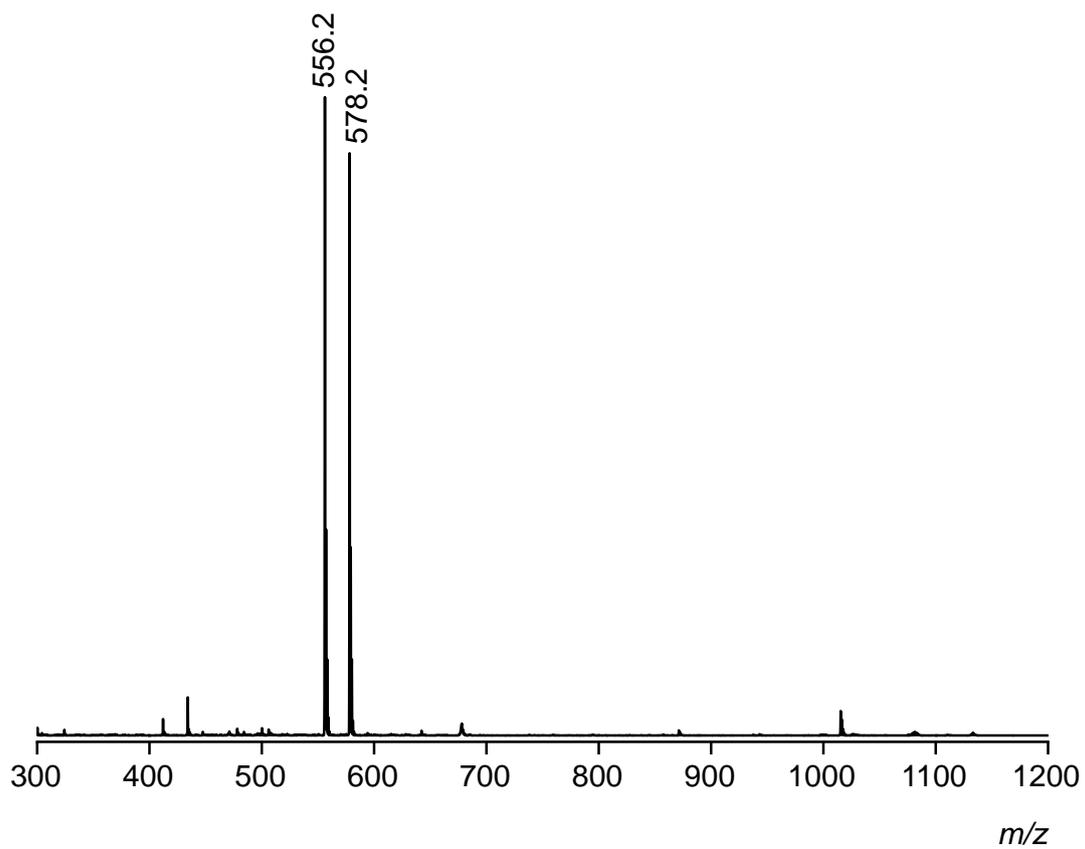
¹H NMR Spectrum: 6 in [D₆]DMSO (600 MHz, 25°C).



¹³C NMR Spectrum: 6 in [D₆]DMSO (151 MHz, 25°C).

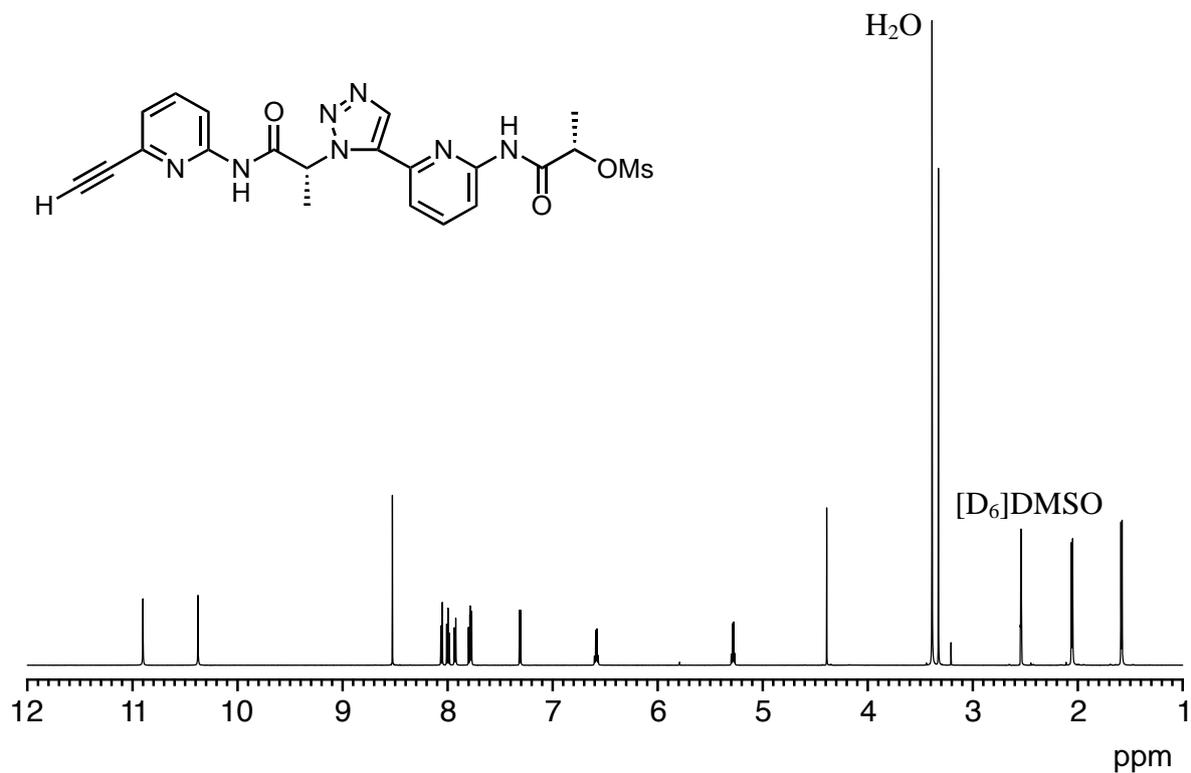


ESI-TOF MS Spectrum: **6** (positive mode).

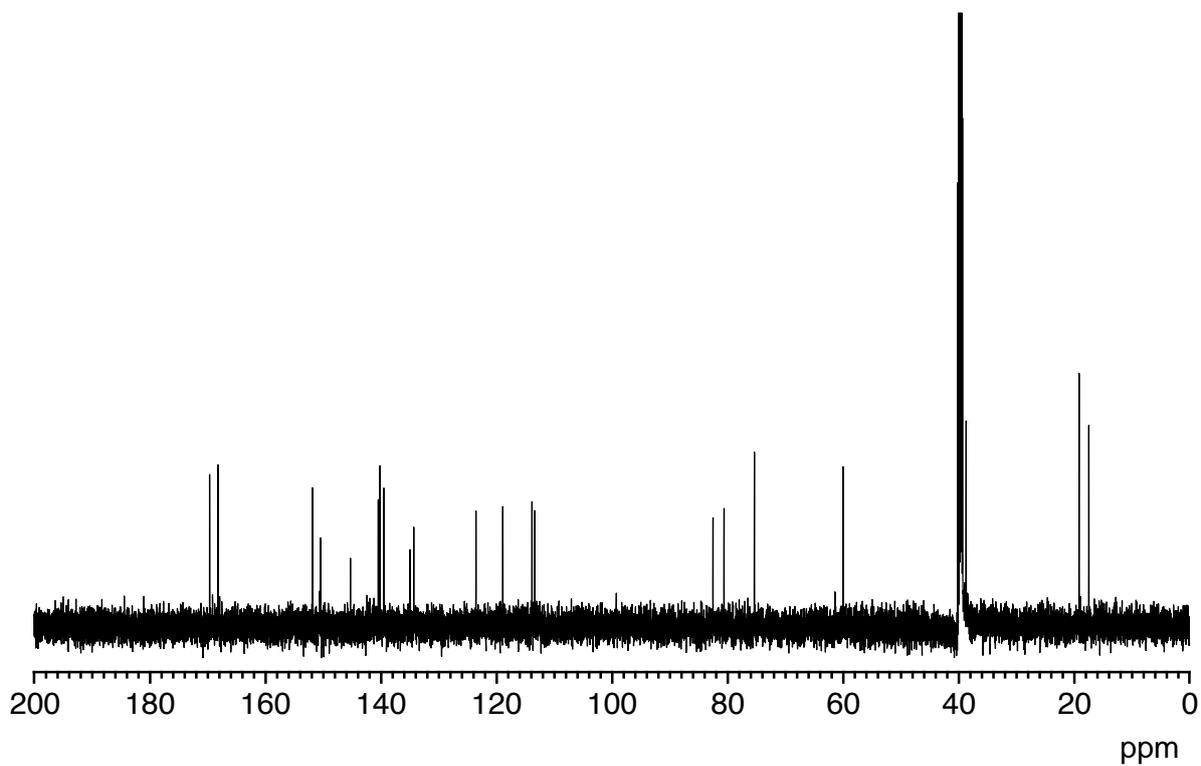


		<i>m/z calcd.</i>	<i>m/z exp.</i>
6 + H ⁺	C ₂₄ H ₂₉ N ₇ O ₅ SSi + H ⁺	556.18	556.2
6 + Na ⁺	C ₂₄ H ₂₉ N ₇ O ₅ SSi + Na ⁺	578.16	578.2

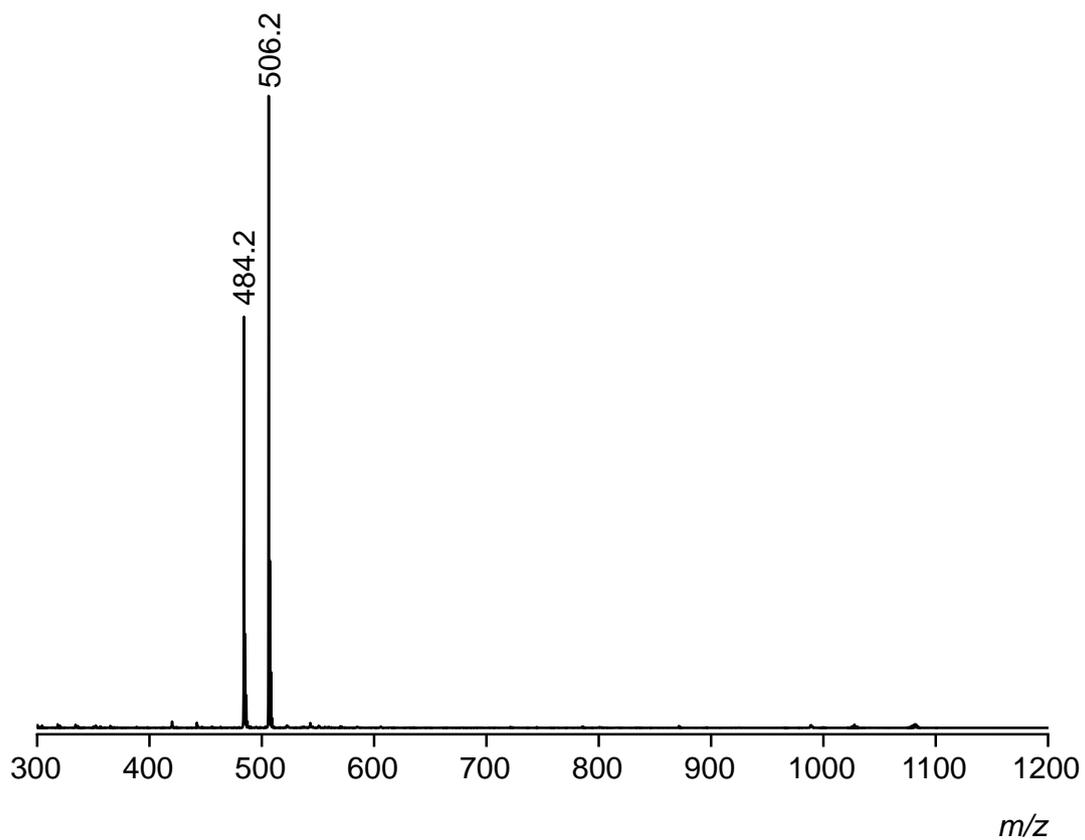
¹H NMR Spectrum: TMS-deprotected **6** in [D₆]DMSO (600 MHz, 25°C).



¹³C NMR Spectrum: TMS-deprotected **6** in [D₆]DMSO (151 MHz, 25°C).

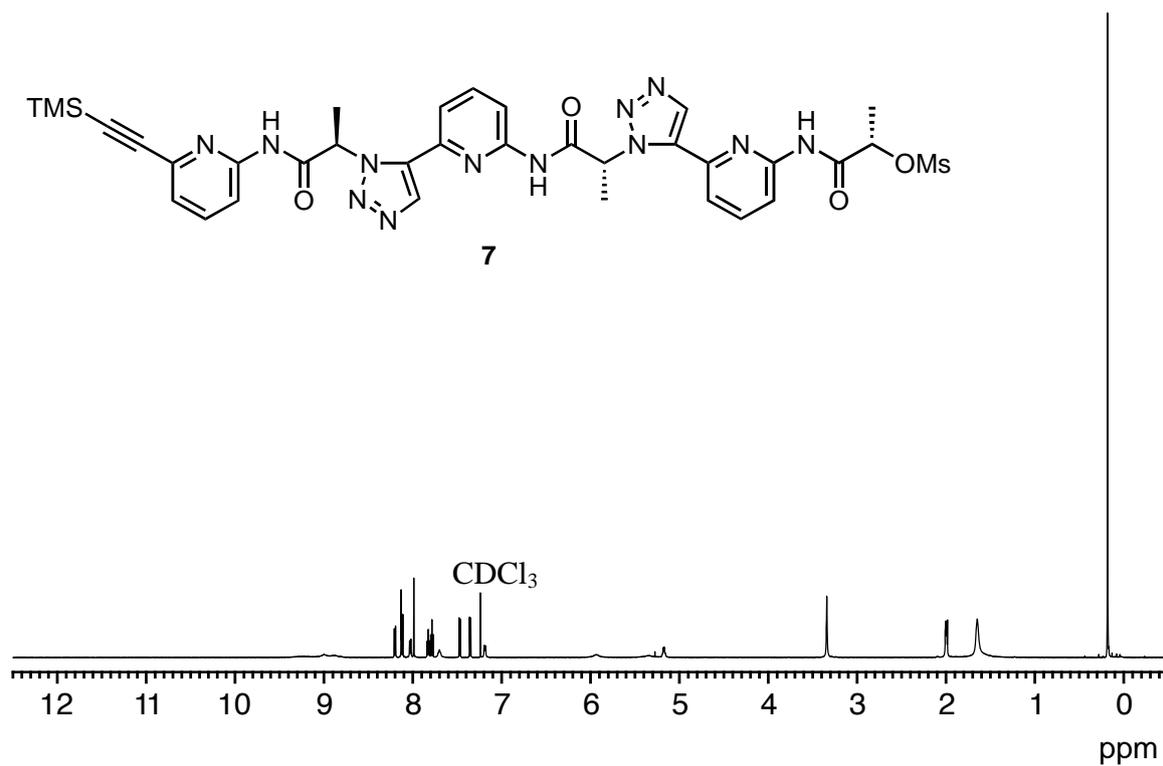


ESI-TOF MS Spectrum: TMS-deprotected **6** (positive mode).

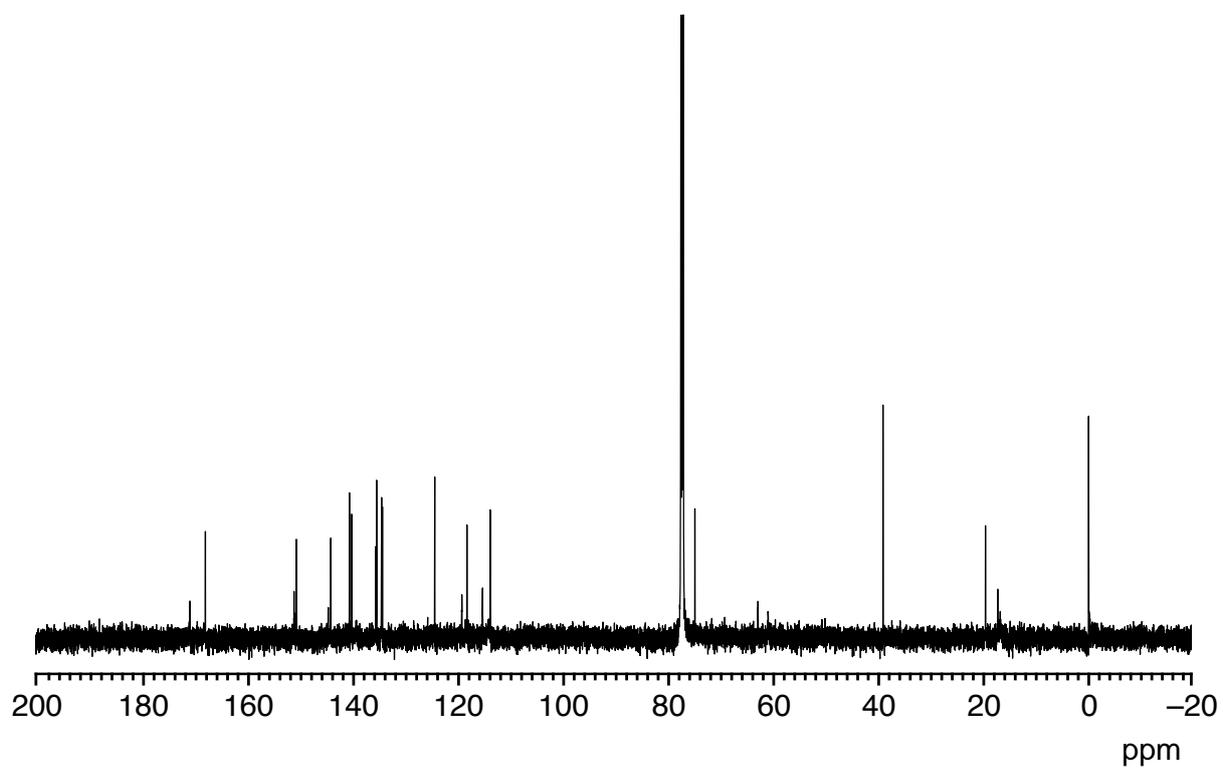


		<i>m/z calcd.</i>	<i>m/z exp.</i>
6-TMS + H ⁺	C ₂₁ H ₂₁ N ₇ O ₅ S + H ⁺	484.14	484.2
6-TMS + Na ⁺	C ₂₁ H ₂₁ N ₇ O ₅ S + Na ⁺	506.12	506.2

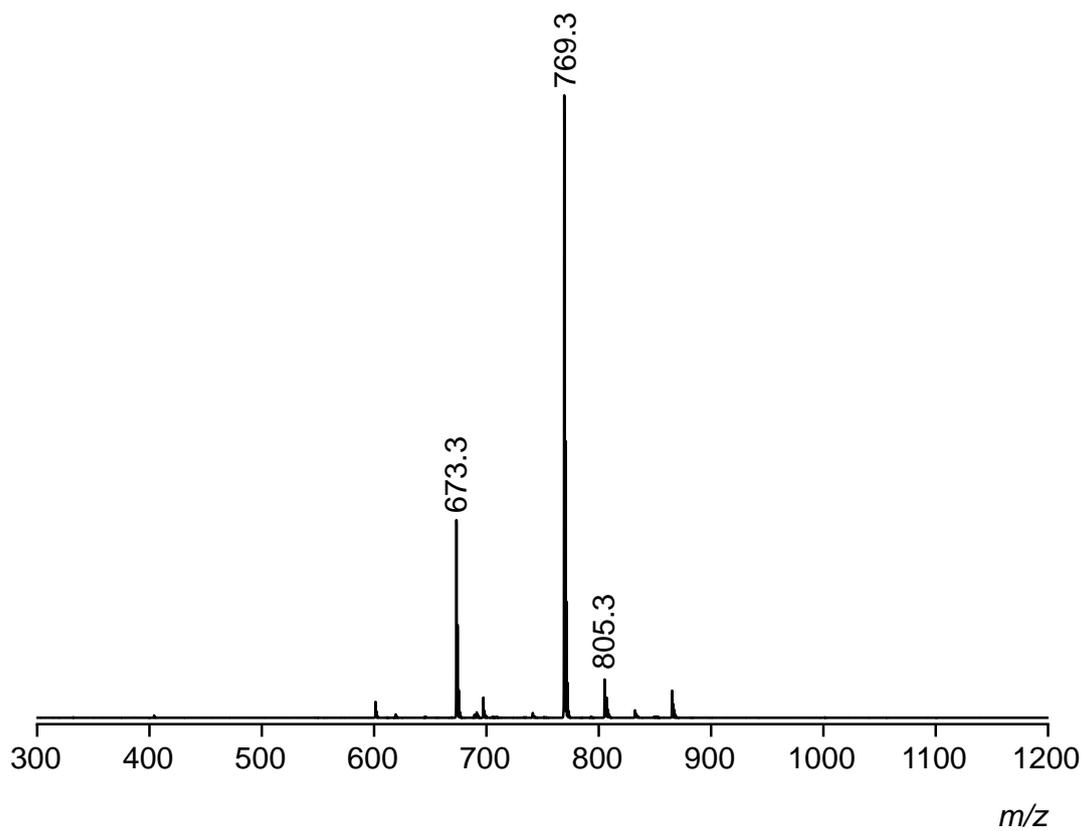
^1H NMR Spectrum: 7 in CDCl_3 (600 MHz, 25°C).



^{13}C NMR Spectrum: 7 in CDCl_3 (151 MHz, 25°C).

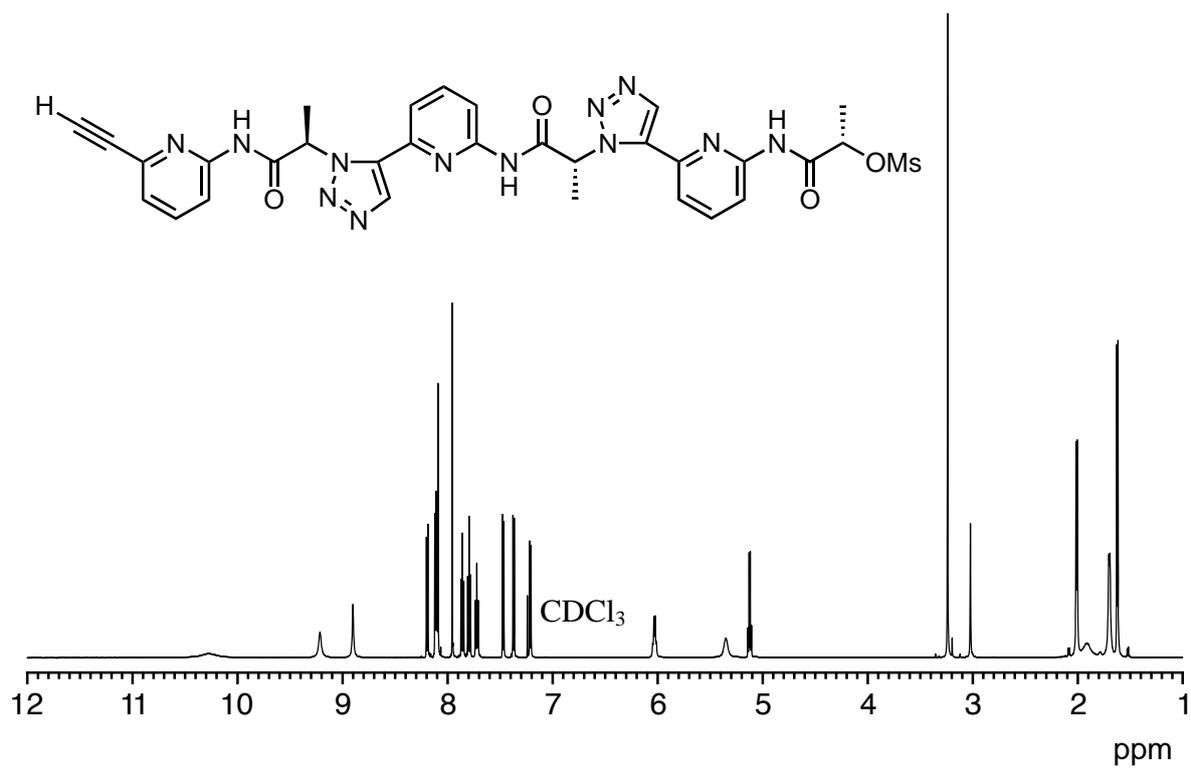


ESI-TOF MS Spectrum: **7** (negative mode).

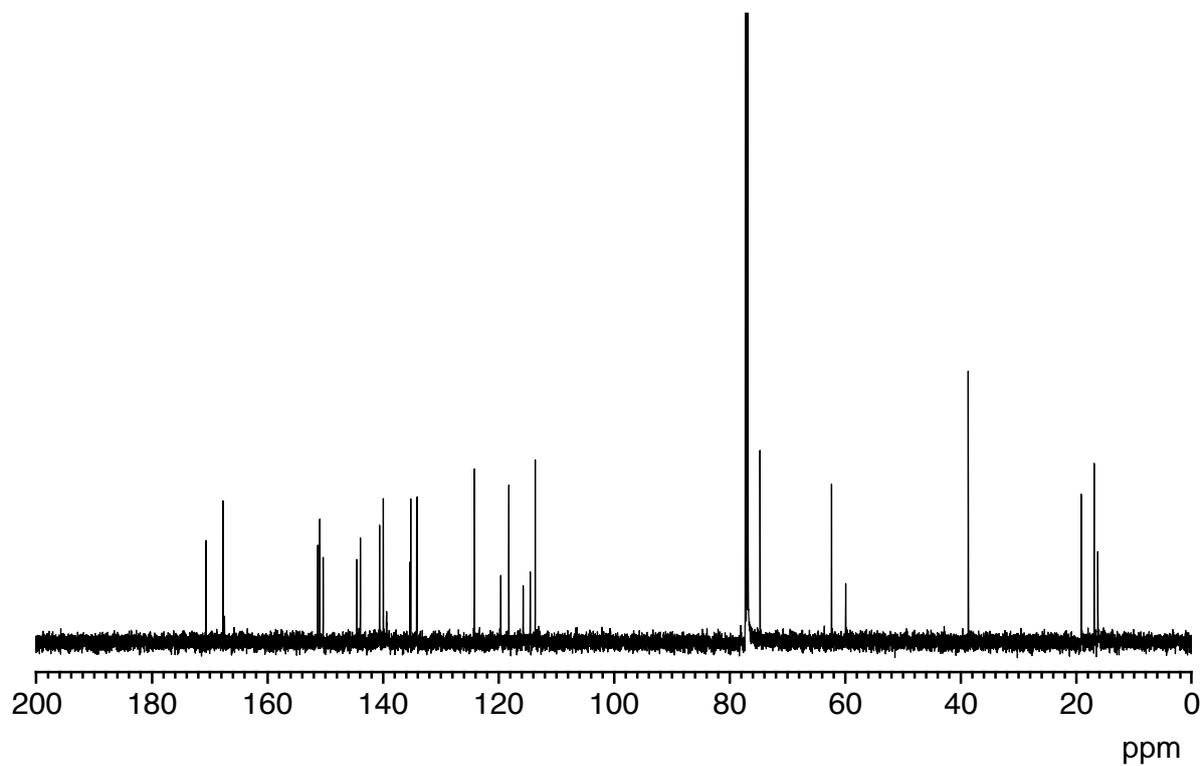


		<i>m/z calcd.</i>	<i>m/z exp.</i>
7 - CH ₃ SO ₃ H - H ⁺	C ₃₄ H ₃₈ N ₁₂ O ₆ SSi - CH ₃ SO ₃ H - H ⁺	673.26	673.3
7 - H ⁺	C ₃₄ H ₃₈ N ₁₂ O ₆ SSi - H ⁺	769.24	769.3
7 + Cl ⁻	C ₃₄ H ₃₈ N ₁₂ O ₆ SSi + Cl ⁻	805.22	805.3

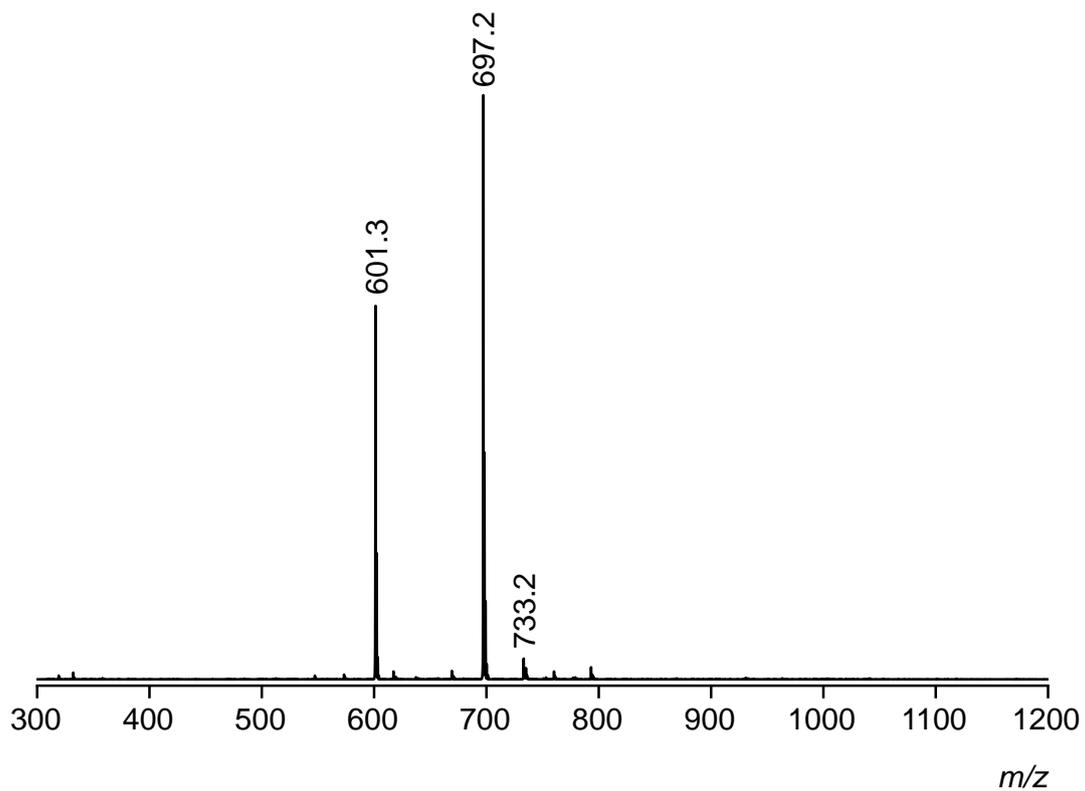
¹H NMR Spectrum: TMS-deprotected **7** in CDCl₃ (600 MHz, 25°C).



¹³C NMR Spectrum: TMS-deprotected **7** in CDCl₃ (151 MHz, 25°C).

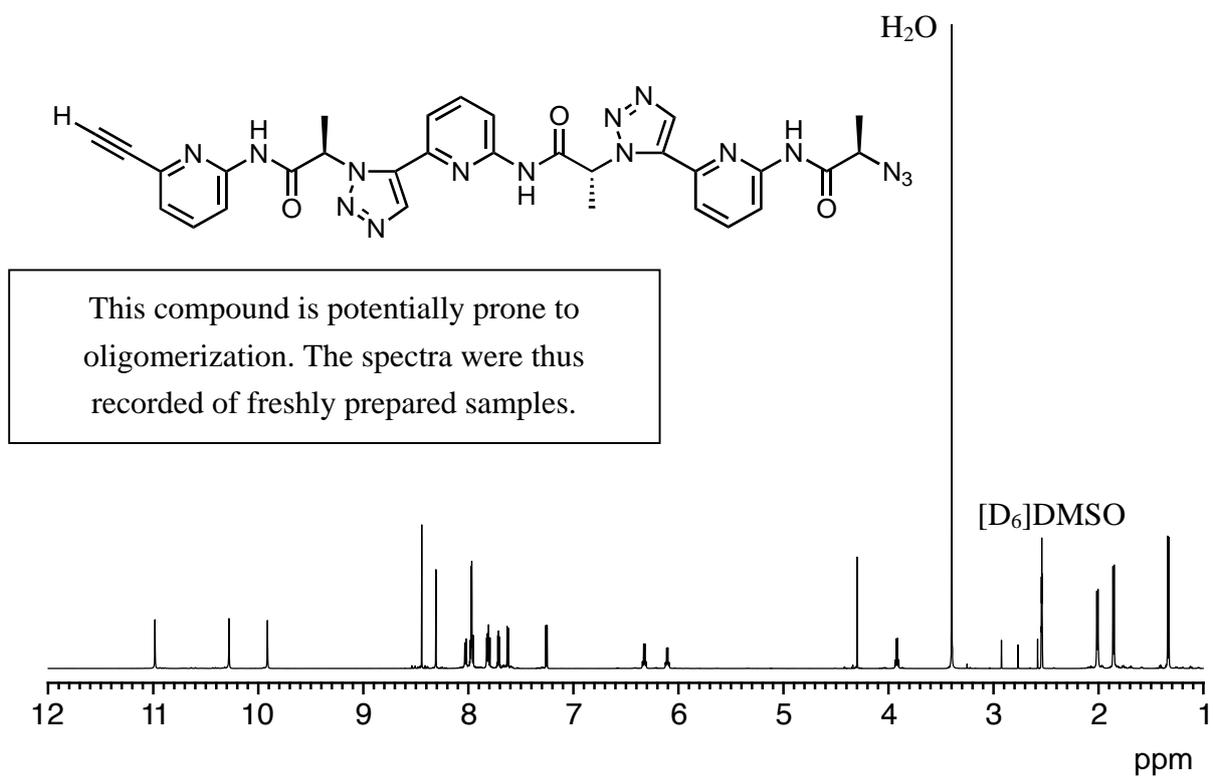


ESI-TOF MS Spectrum: TMS-deprotected **7** (negative mode).

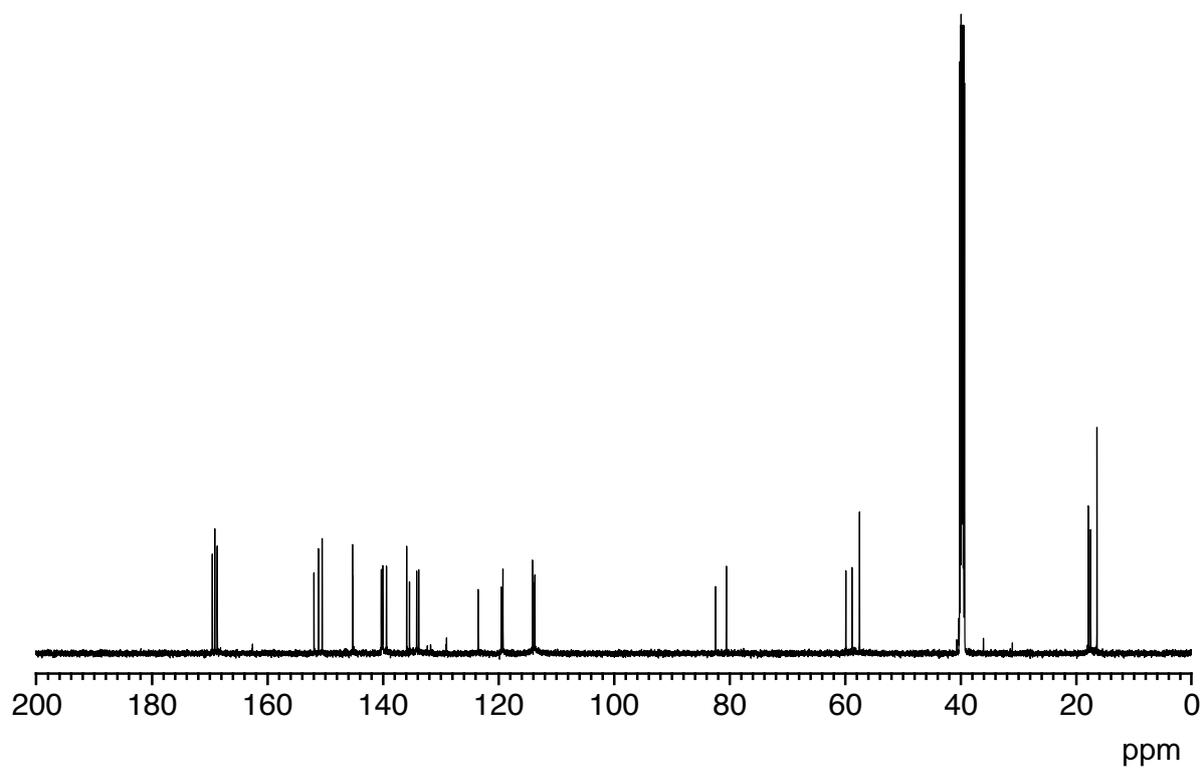


		<i>m/z, calcd.</i>	<i>m/z, exp.</i>
7-TMS – CH ₃ SO ₃ H – H ⁺	C ₃₁ H ₃₀ N ₁₂ O ₆ S – CH ₃ SO ₃ H – H ⁺	601.22	601.3
7-TMS – H ⁺	C ₃₁ H ₃₀ N ₁₂ O ₆ S – H ⁺	697.21	697.2
7-TMS + Cl ⁻	C ₃₁ H ₃₀ N ₁₂ O ₆ S + Cl ⁻	733.18	733.2

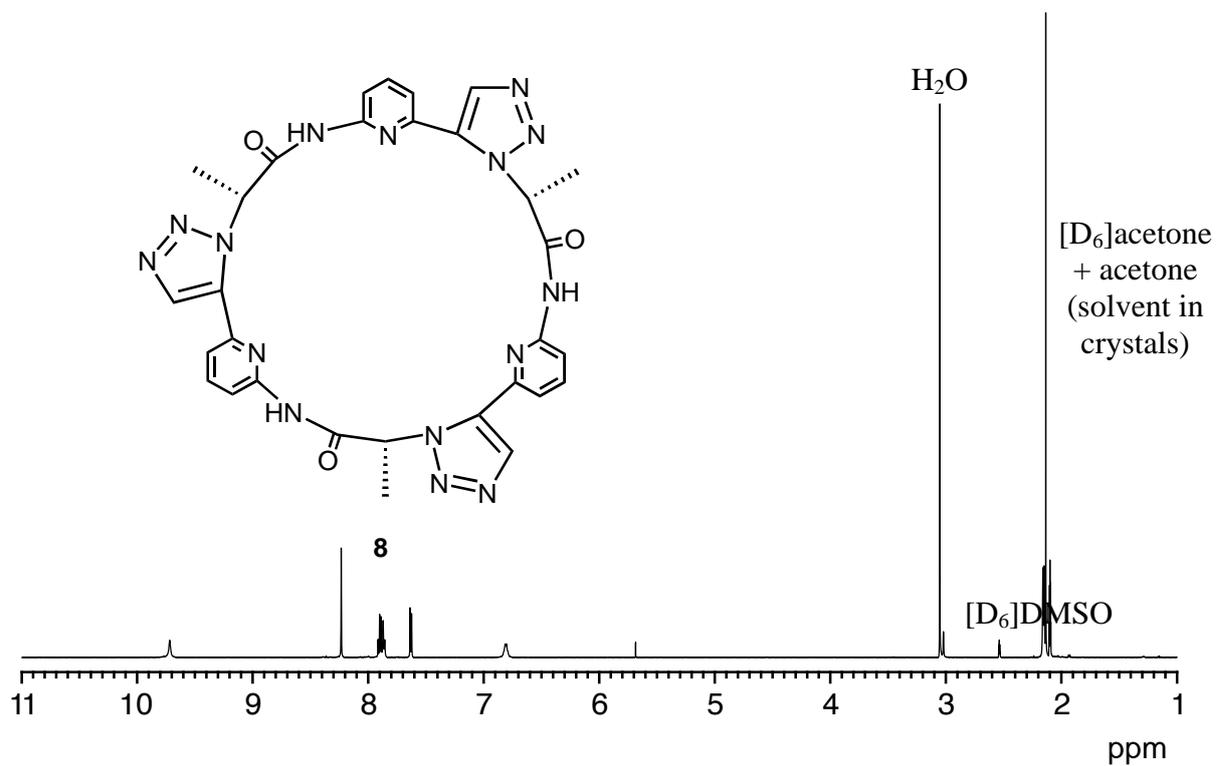
¹H NMR Spectrum: Linear precursor of **8** in [D₆]DMSO (600 MHz, 25°C).



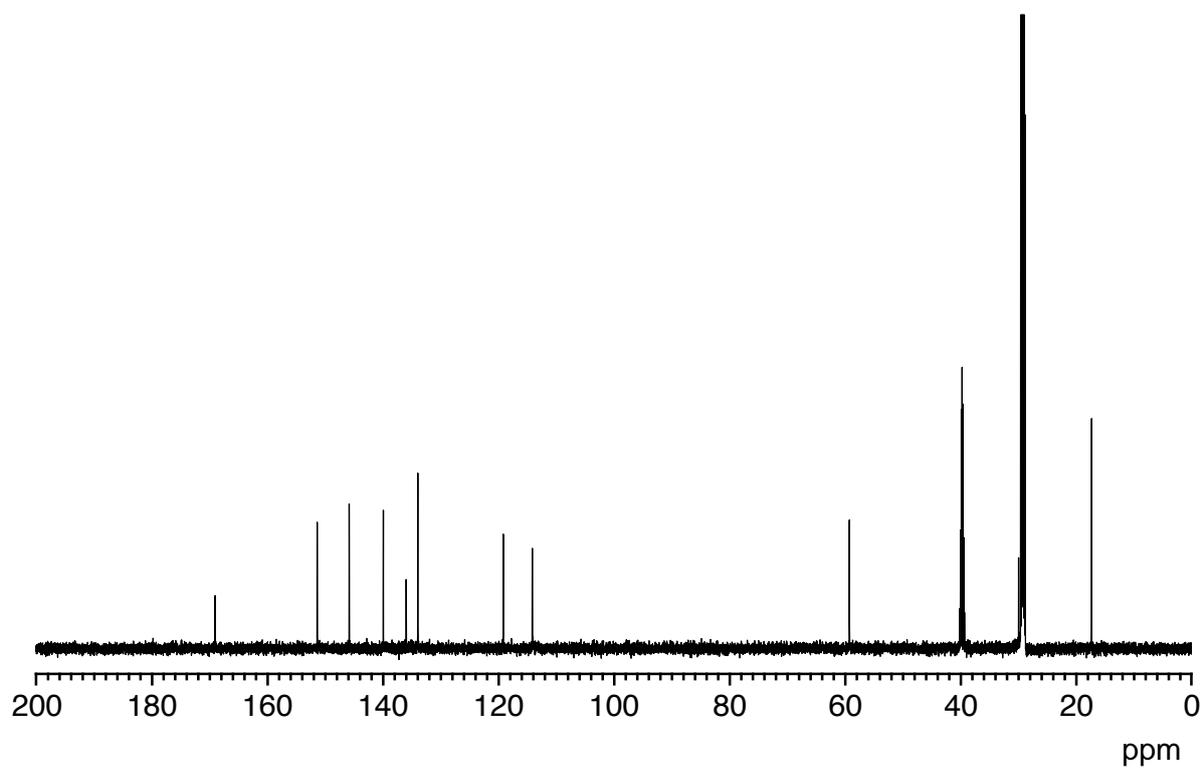
¹³C NMR Spectrum: Linear precursor of **8** in [D₆]DMSO (151 MHz, 25°C).



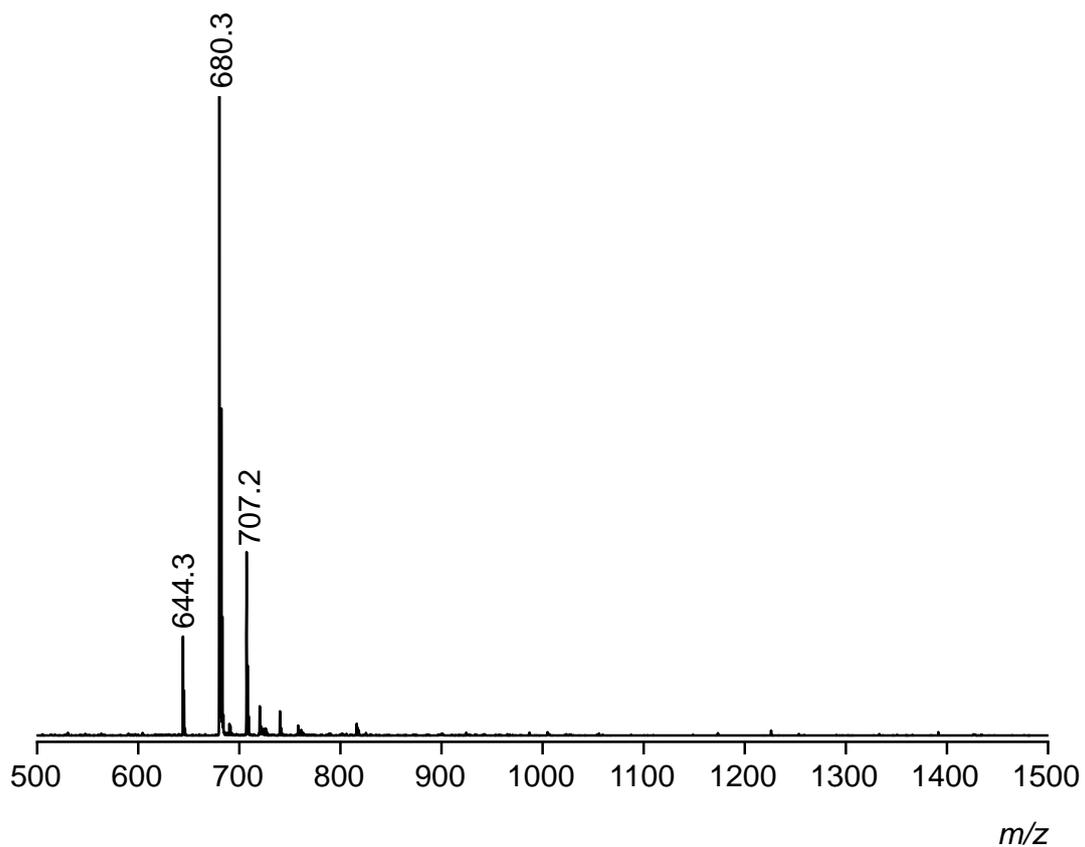
¹H NMR Spectrum: **8** in [D₆]DMSO/[D₆]acetone 1:9 (v/v) (600 MHz, 25°C).



¹³C NMR Spectrum: **8** in [D₆]DMSO/[D₆]acetone 1:9 (v/v) (151 MHz, 25°C).

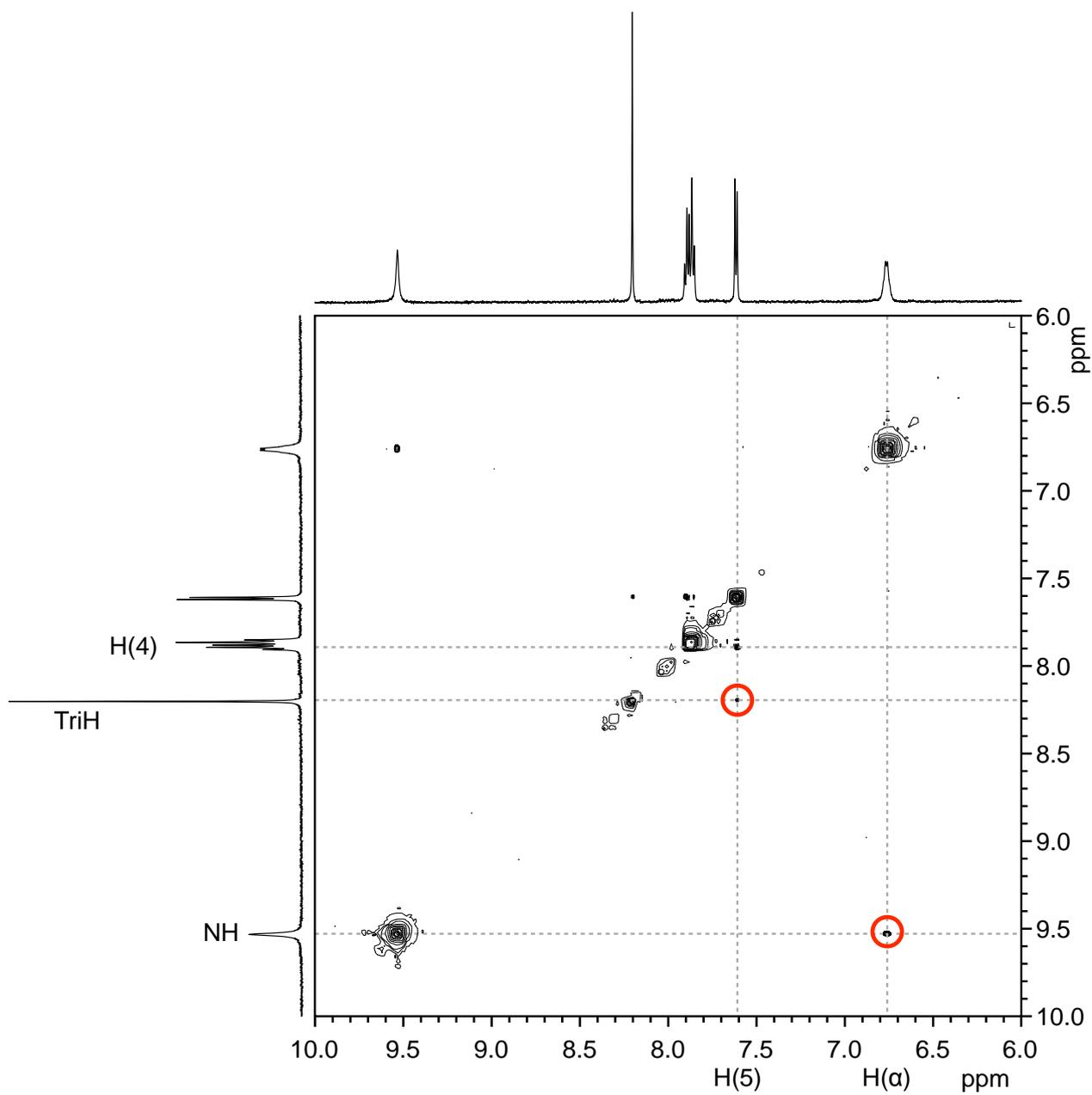


ESI-TOF MS Spectrum: **8** (negative mode).

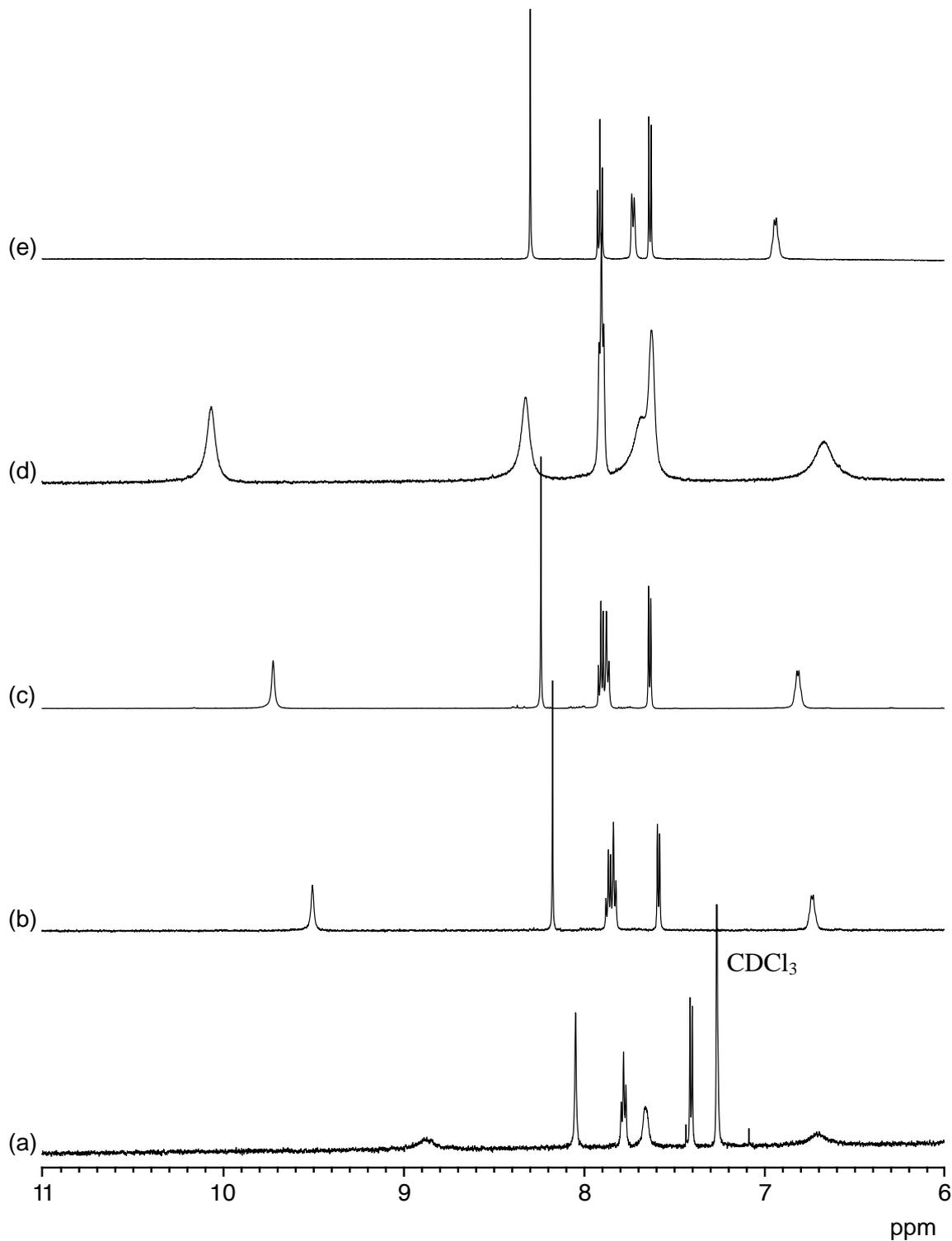


		<i>m/z calcd.</i>	<i>m/z exp.</i>
8 - H ⁺	(C ₃₀ H ₂₇ N ₁₅ O ₃) - H ⁺	644.23	644.3
8 + Cl ⁻	(C ₃₀ H ₂₇ N ₁₅ O ₃) + Cl ⁻	680.21	680.3
8 + NO ₃ ⁻	(C ₃₀ H ₂₇ N ₁₅ O ₃) + NO ₃ ⁻	707.23	707.2

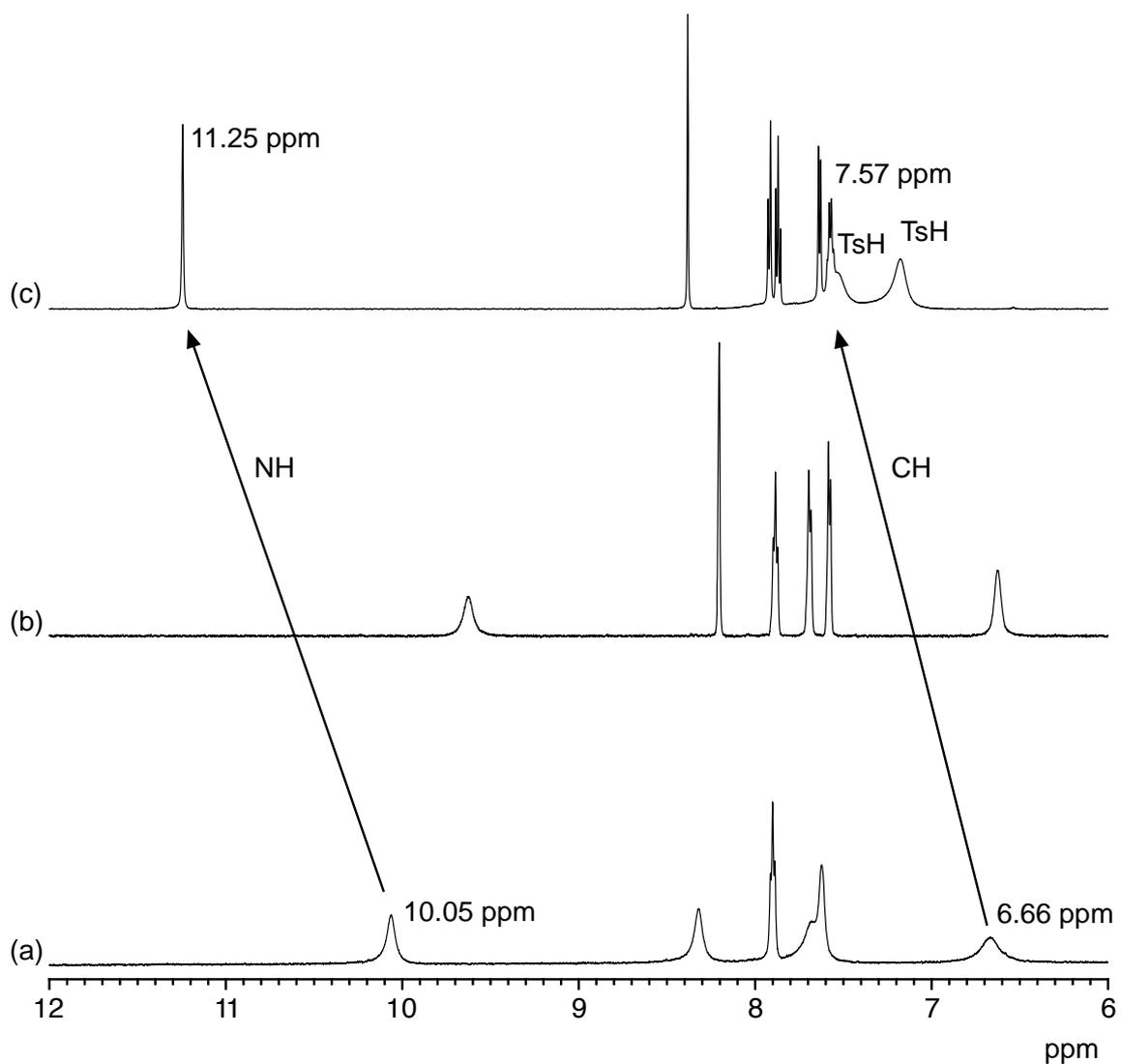
NOESY NMR Spectrum: 8 (2 mM) in [D₆]acetone (mixing time 150 ms) (600 MHz, 25 °C).



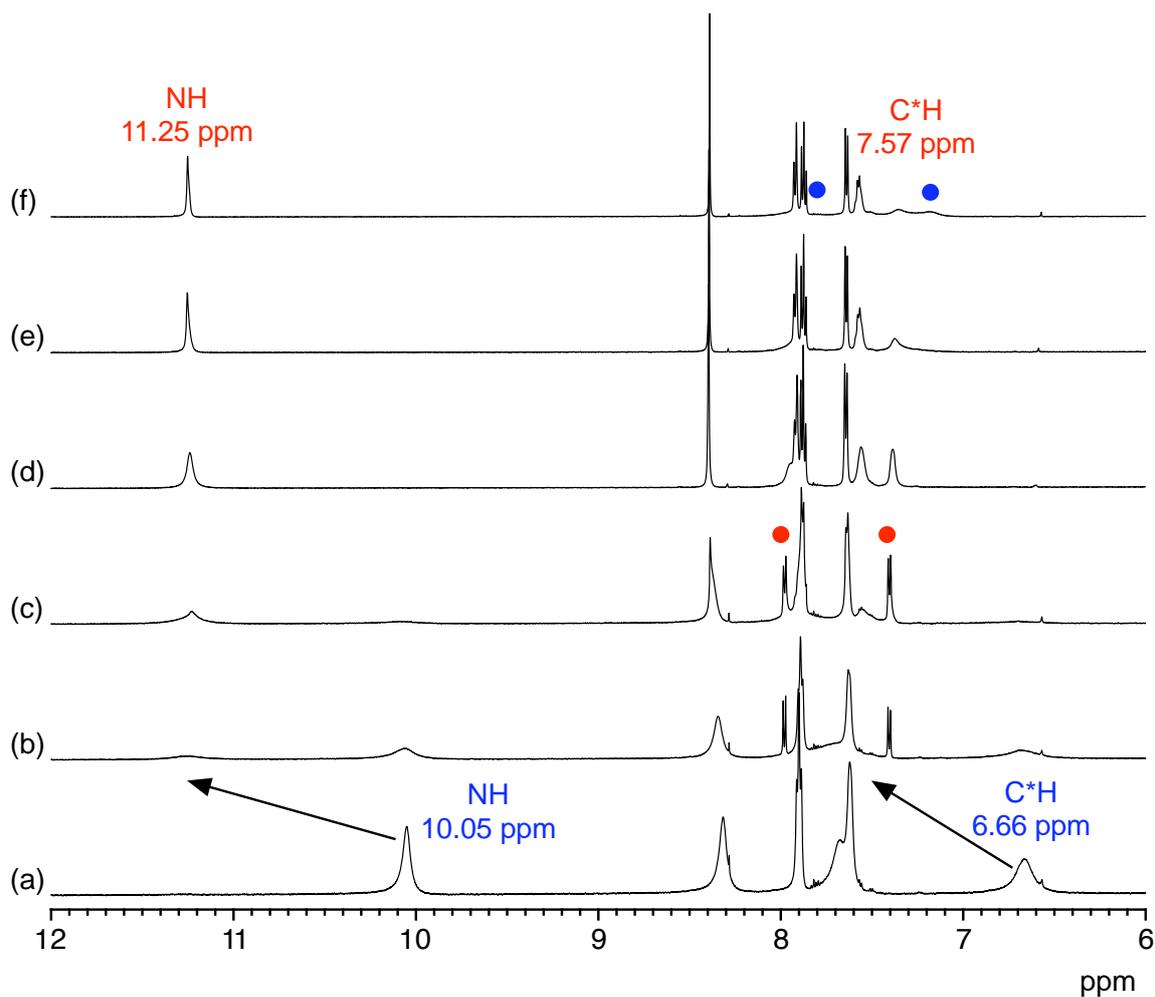
¹H NMR Spectrum: 8 in CDCl₃ (a), in [D₆]acetone (b), in 10% [D₆]DMSO/[D₆]acetone (c), in [D₆]DMSO (d), and in D₂O/CD₃OD 1:1 (v/v) (e) (600 MHz, 25°C).



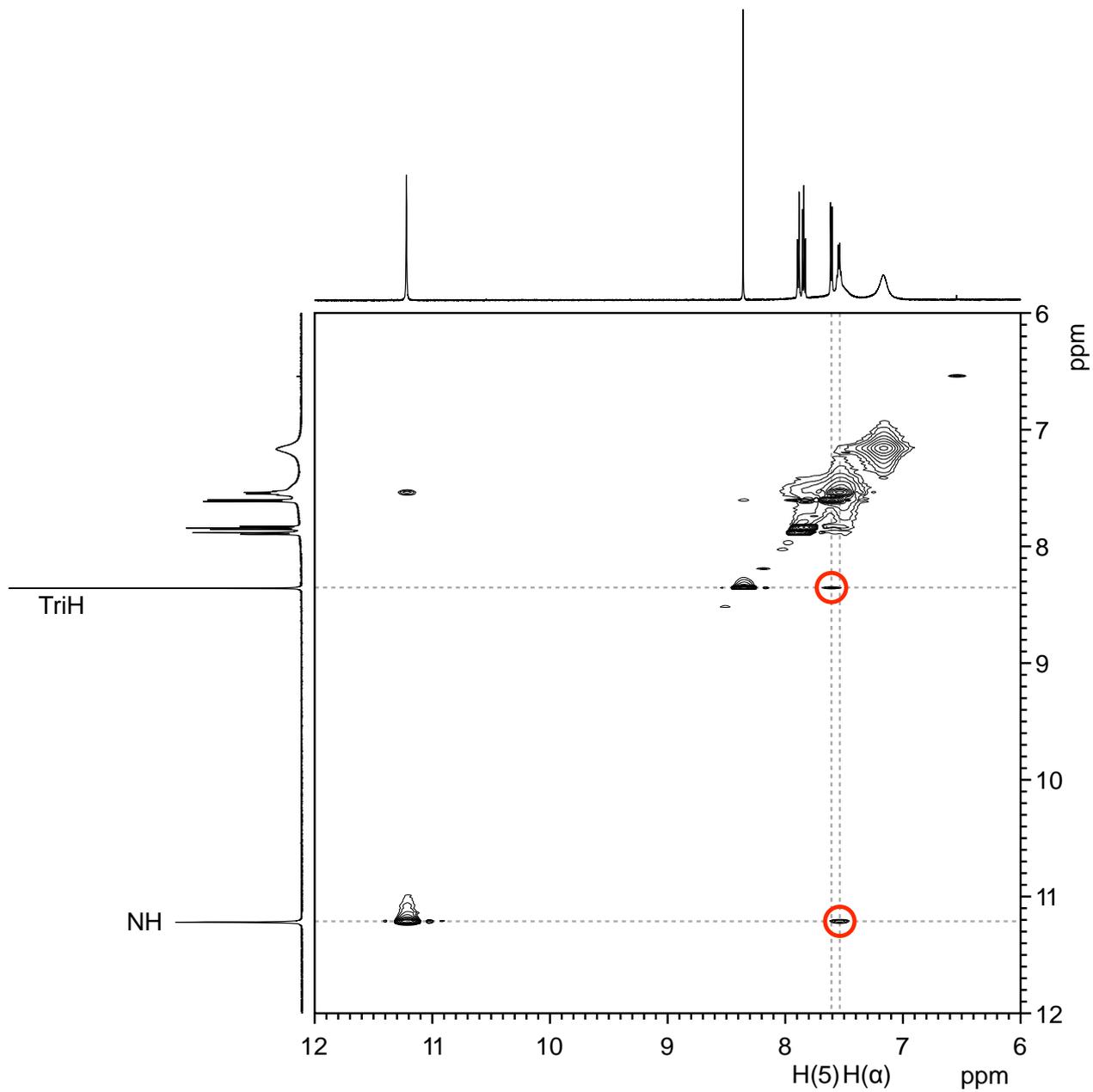
¹H NMR Spectrum: **8** (2 mM) in [D₆]DMSO at 25 °C (a), at 100 °C (b), and at 25 °C in the presence of 3.5 equivalents of *n*-butyltrimethylammonium tosylate (c) (600 MHz).



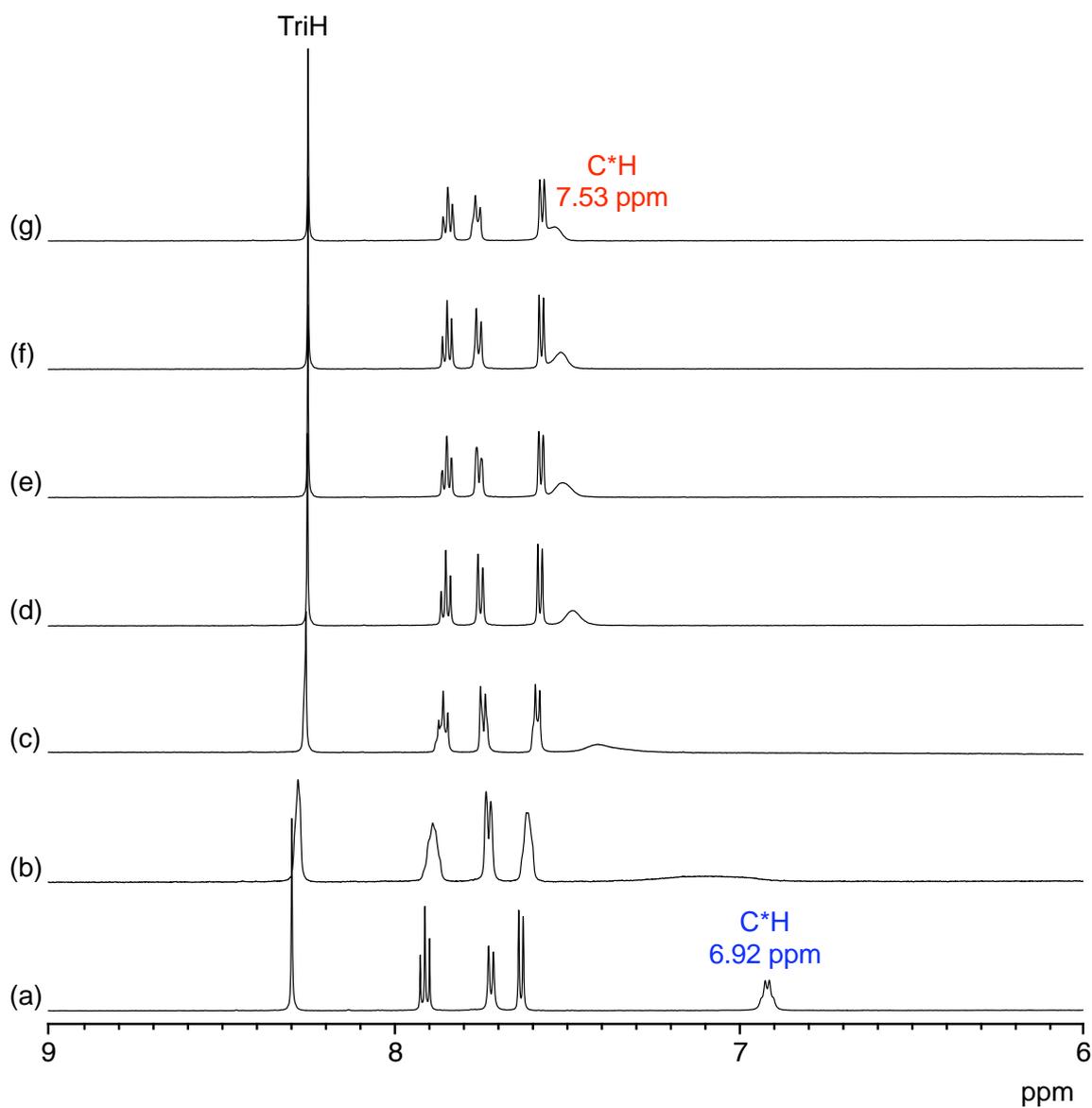
¹H NMR Spectrum: **8** (2 mM) in [D₆]DMSO in the presence of 0 equiv (a), 0.4 equiv (b), 0.8 equiv (c), 1.0 equiv (d), 1.4 equiv (e), and 1.8 equiv (f) of *n*-butyltrimethylammonium tosylate at 25 °C. The blue assignments correspond to the signals of the free receptor and the red ones to those of the complex. The dots mark the aromatic tosylate signals with red indicating complexed and blue indicating free anion (600 MHz, 25°C).



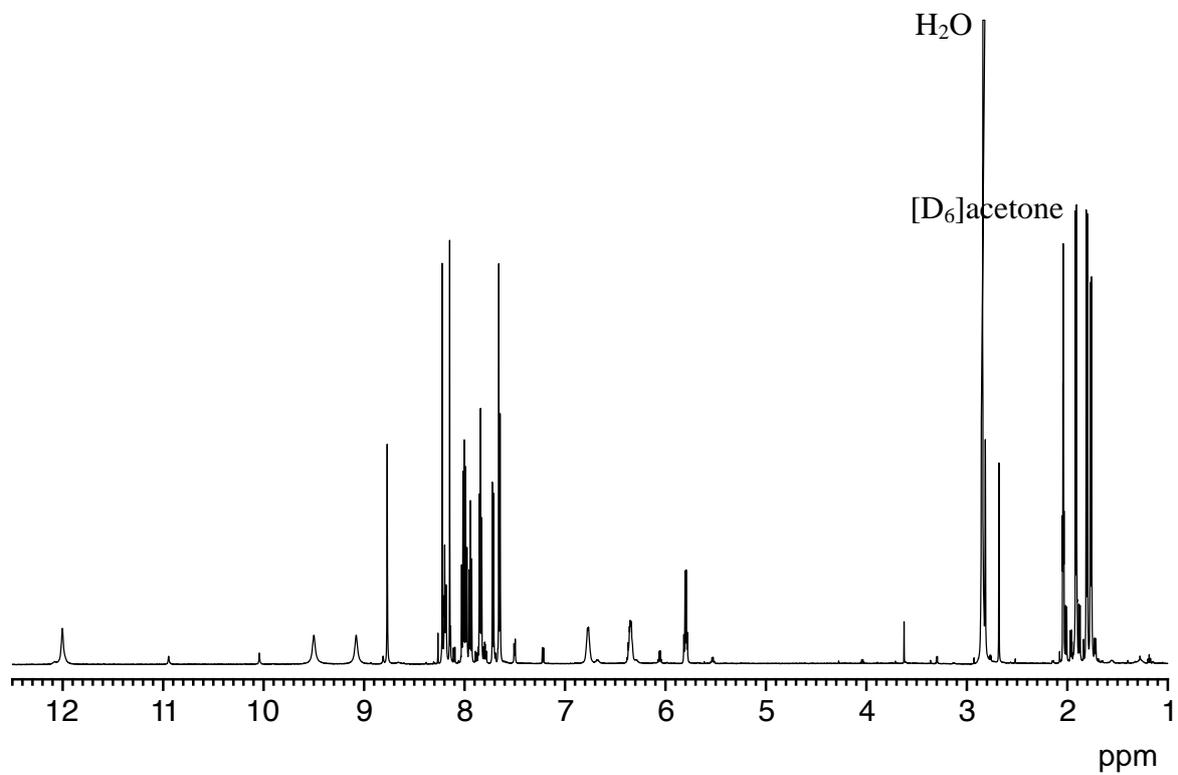
NOESY NMR Spectrum: **8** (2 mM) in [D₆]DMSO in the presence of 3.5 equivalents of *n*-butyltrimethylammonium tosylate (mixing time 150 ms) (600 MHz, 25 °C).



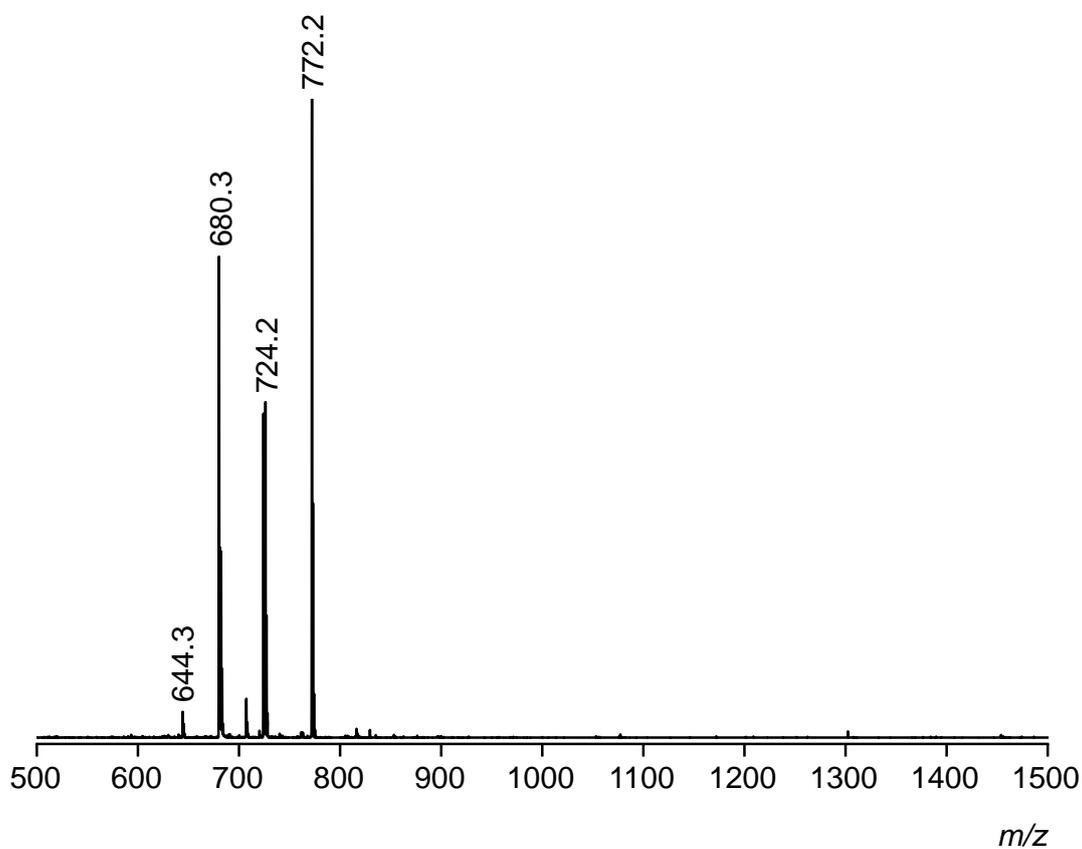
¹H NMR Spectrum: **8** (1 mM) in CD₃OD/D₂O 2:1 (v/v) in the presence of 0 equiv (a), 0.25 equiv (b), 0.5 equiv (c), 0.75 equiv (d), 1.0 equiv (e), 1.25 equiv (f), and 1.50 equiv of Na₂SO₄. The blue assignments correspond to the signals of the free receptor and the red ones to those of the complex (600 MHz, 25°C).



^1H NMR Spectrum: Side product formed besides **8** during the cyclization in $[\text{D}_6]$ acetone (600 MHz, 25°C).

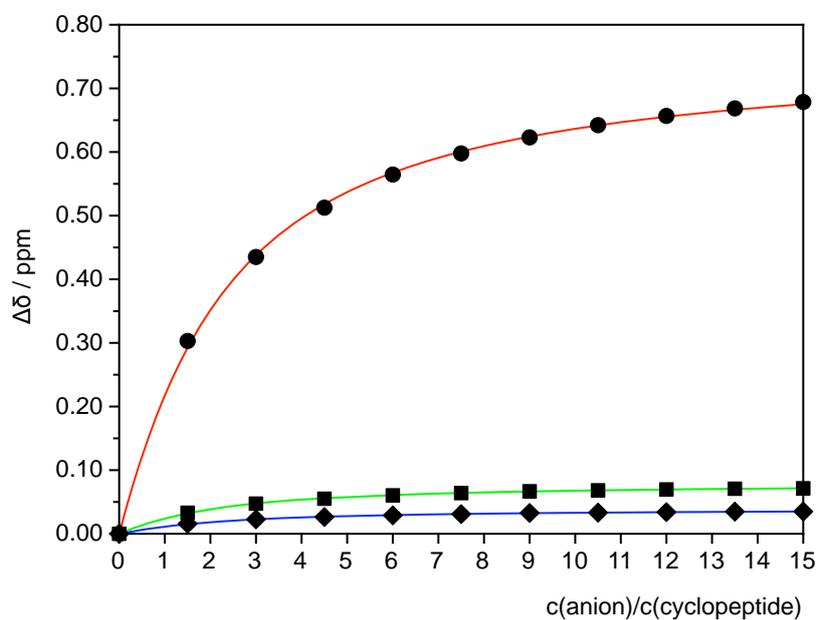


ESI-TOF MS Spectrum: **8** (10 μ M) in the presence of 0.33 equivalents of each NaCl, NaBr, NaI in 1:1 (v/v) H₂O/CH₃OH (negative mode).



		<i>m/z calcd.</i>	<i>m/z exp.</i>
8 - H ⁺	(C ₃₀ H ₂₇ N ₁₅ O ₃) - H ⁺	644.23	644.3
8 + Cl ⁻	(C ₃₀ H ₂₇ N ₁₅ O ₃) + Cl ⁻	680.21	680.3
8 + Br ⁻	(C ₃₀ H ₂₇ N ₁₅ O ₃) + Br ⁻	724.16	724.2
8 + I ⁻	(C ₃₀ H ₂₇ N ₁₅ O ₃) + I ⁻	772.15	772.2

NMR Titration: **1b** with tetramethylammonium bromide in CD₃OD ($T = 298$ K)

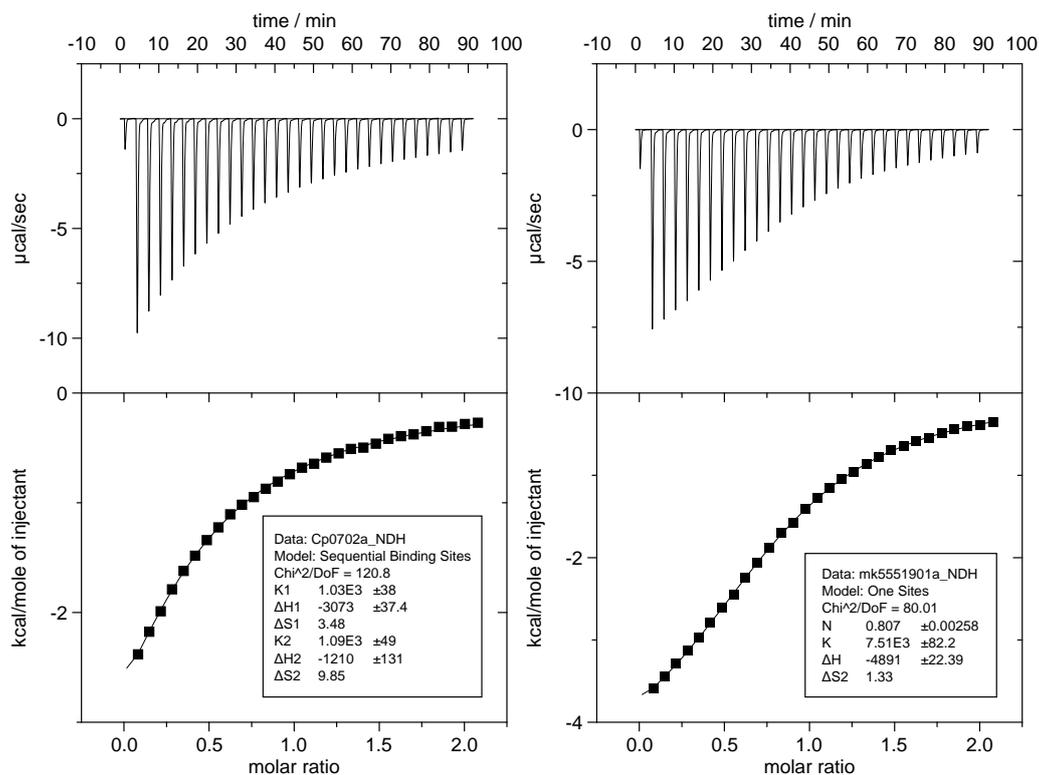


H(α) **Parameters:** **Standard deviations:**
 K = 559 ΔK = 16
 $\Delta\delta_{\text{max}}$ = 0.7607 $\Delta\Delta\delta_{\text{max}}$ = 0.0045

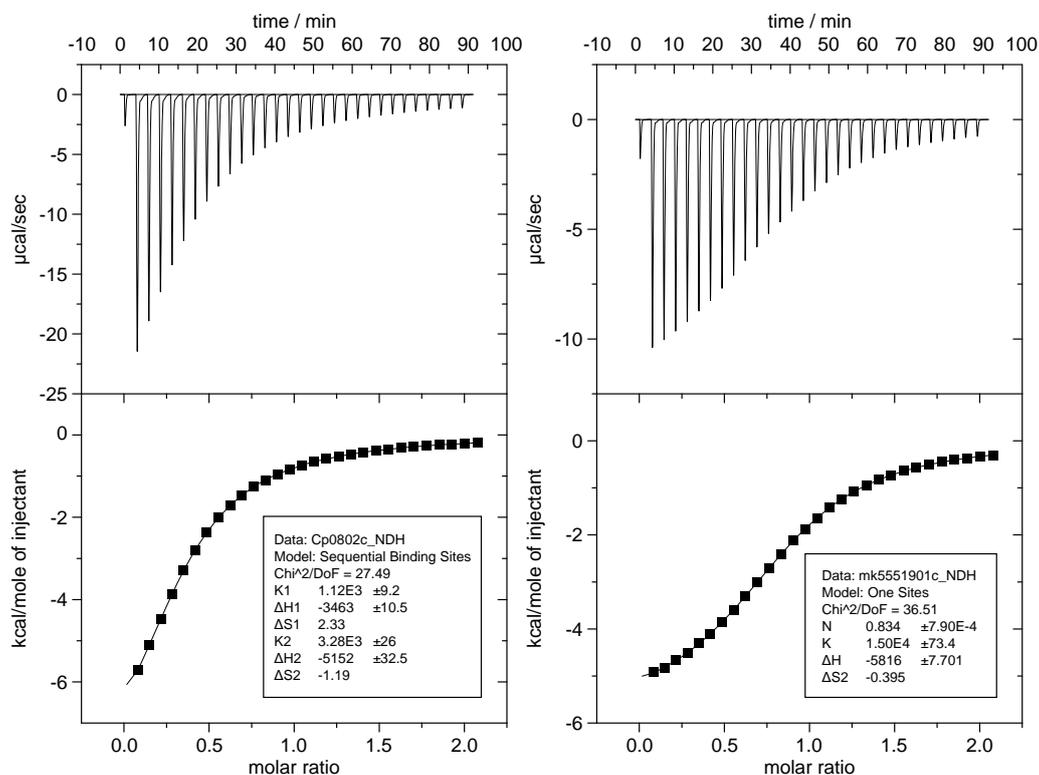
ApaH(5) **Parameters:** **Standard deviations:**
 K = 558 ΔK = 24
 $\Delta\delta_{\text{max}}$ = 0.0394 $\Delta\Delta\delta_{\text{max}}$ = 0.0004

ApaH(3) **Parameters:** **Standard deviations:**
 K = 616 ΔK = 17
 $\Delta\delta_{\text{max}}$ = 0.0797 $\Delta\Delta\delta_{\text{max}}$ = 0.0004

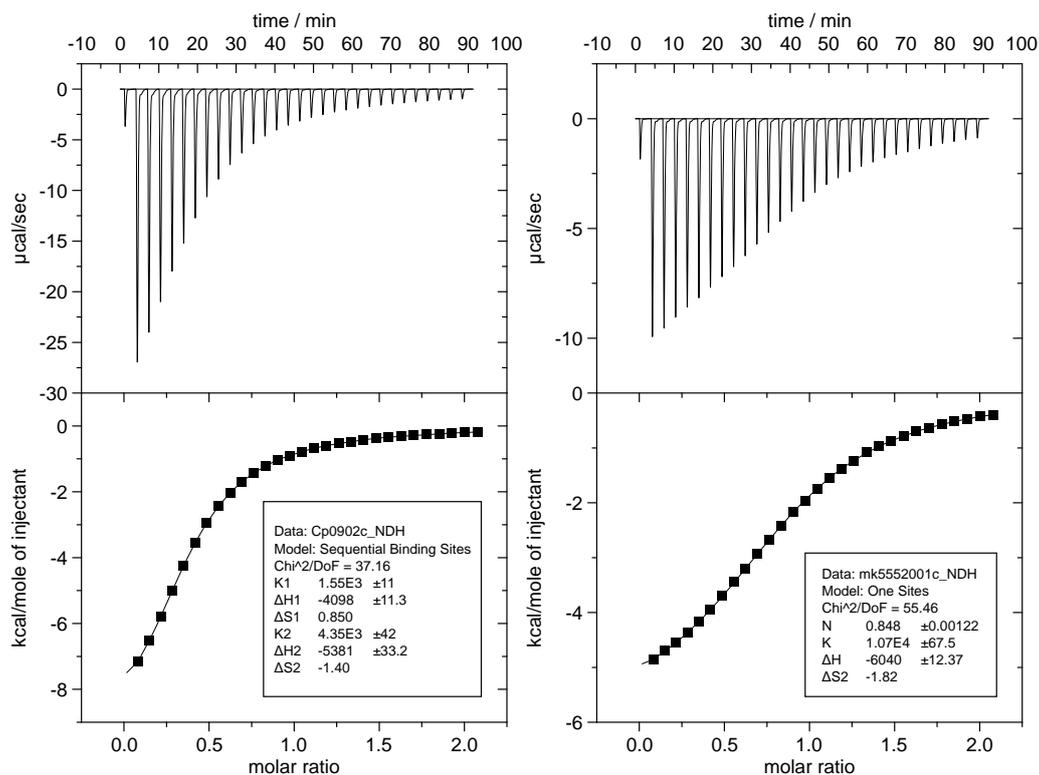
ITC Traces and Binding Isotherms: Titrations of receptors **1a** (left) **8** (right) with (CH₃)₄NCl in CH₃OH at 298 K.



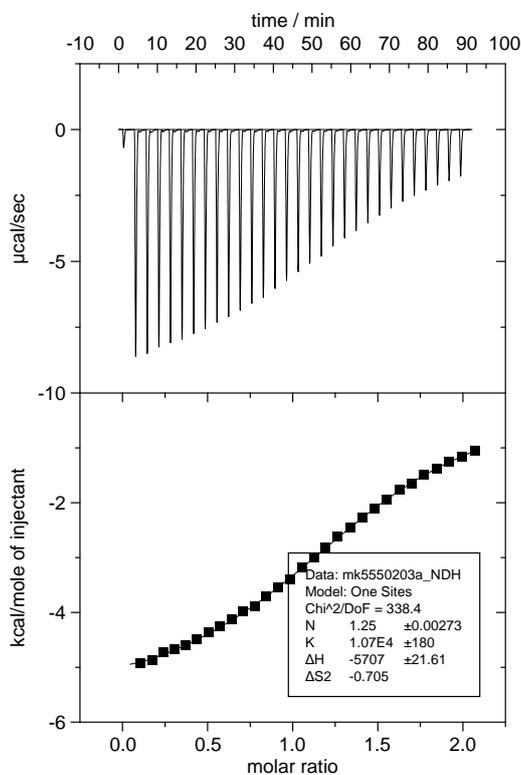
ITC Traces and Binding Isotherms: Titrations of receptors **1a** (left) **8** (right) with (CH₃)₄NBr in CH₃OH at 298 K.



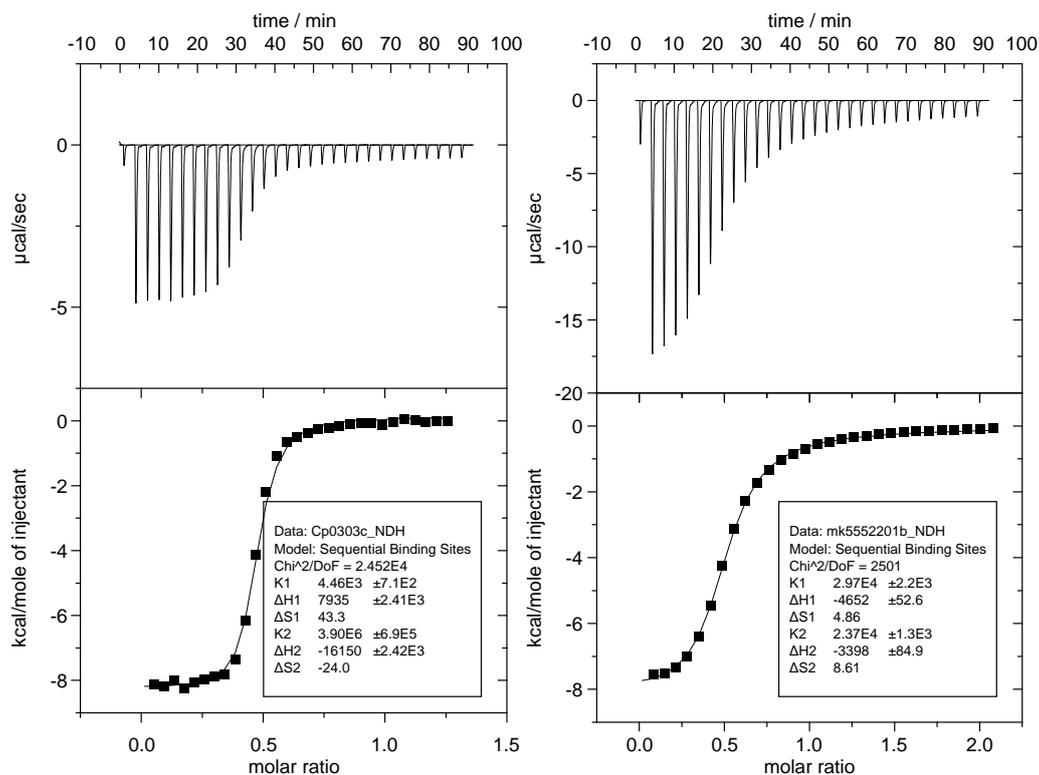
ITC Traces and Binding Isotherms: Titrations of receptors **1a** (left) **8** (right) with (CH₃)₄NI in CH₃OH at 298 K.



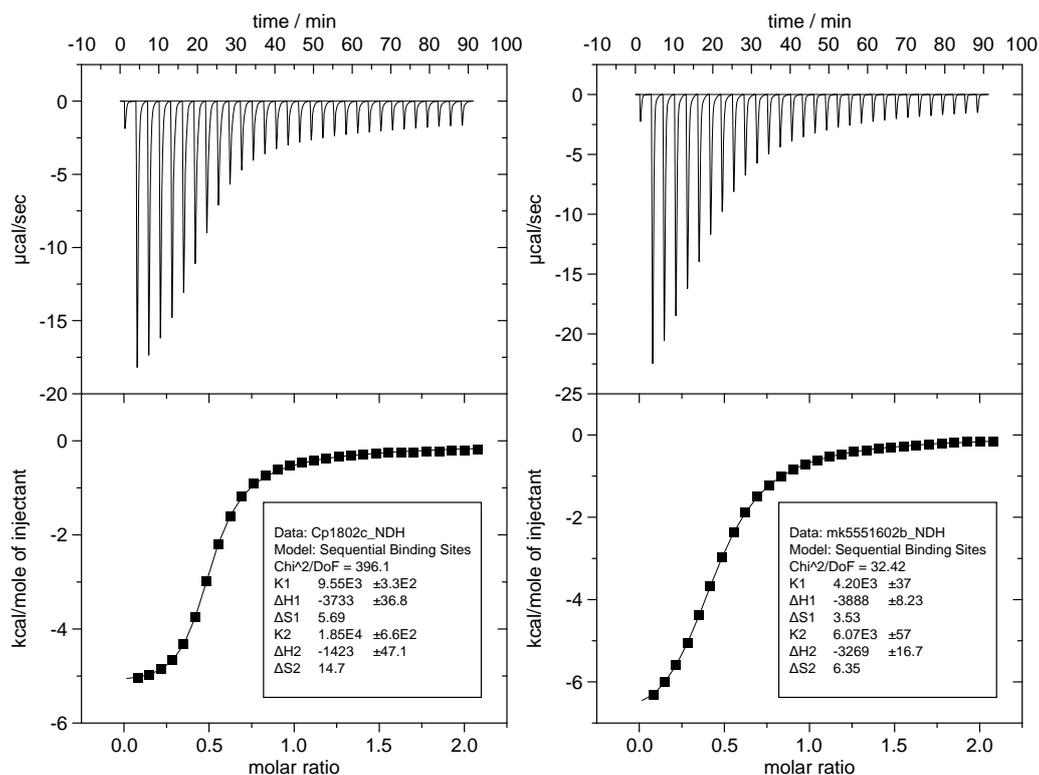
ITC Traces and Binding Isotherms: Titration of (CH₃)₄NI with receptor **8** in CH₃OH at 298 K.



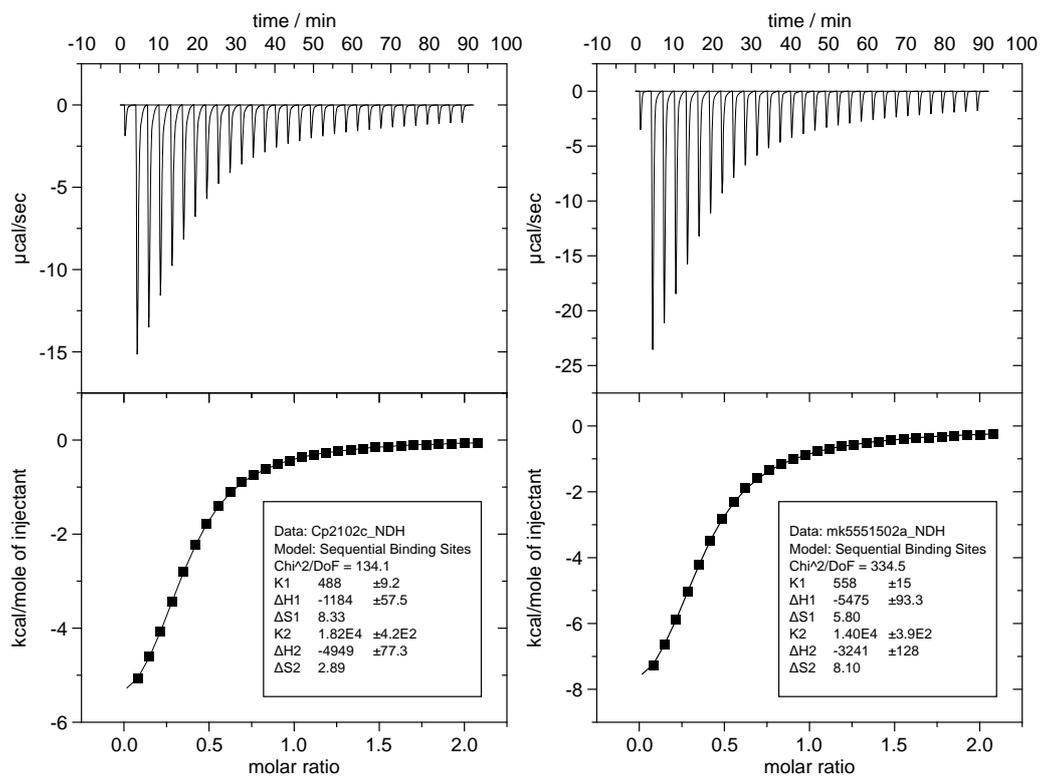
ITC Traces and Binding Isotherms: Titrations of receptors **1a** (left) **8** (right) with $[(\text{CH}_3)_4\text{N}]_2\text{SO}_4$ in CH_3OH at 298 K.



ITC Traces and Binding Isotherms: Titrations of receptors **1a** (left) **8** (right) with $[(\text{CH}_3)_4\text{N}]_2\text{SO}_4$ in $\text{CH}_3\text{OH}/\text{H}_2\text{O}$ 4:1 (v/v) at 298 K.

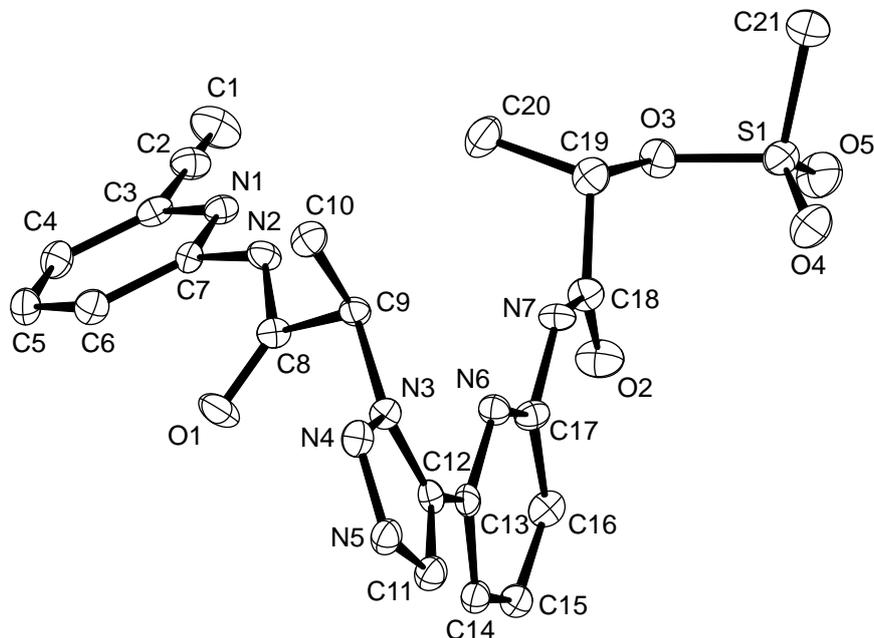


ITC Traces and Binding Isotherms: Titrations of receptors **1a** (left) **8** (right) with $[(\text{CH}_3)_4\text{N}]_2\text{SO}_4$ in $\text{CH}_3\text{OH}/\text{H}_2\text{O}$ 2:1 (v/v) at 298 K.



X-ray Crystal Structure Analysis: TMS-Deprotected Derivative of 6

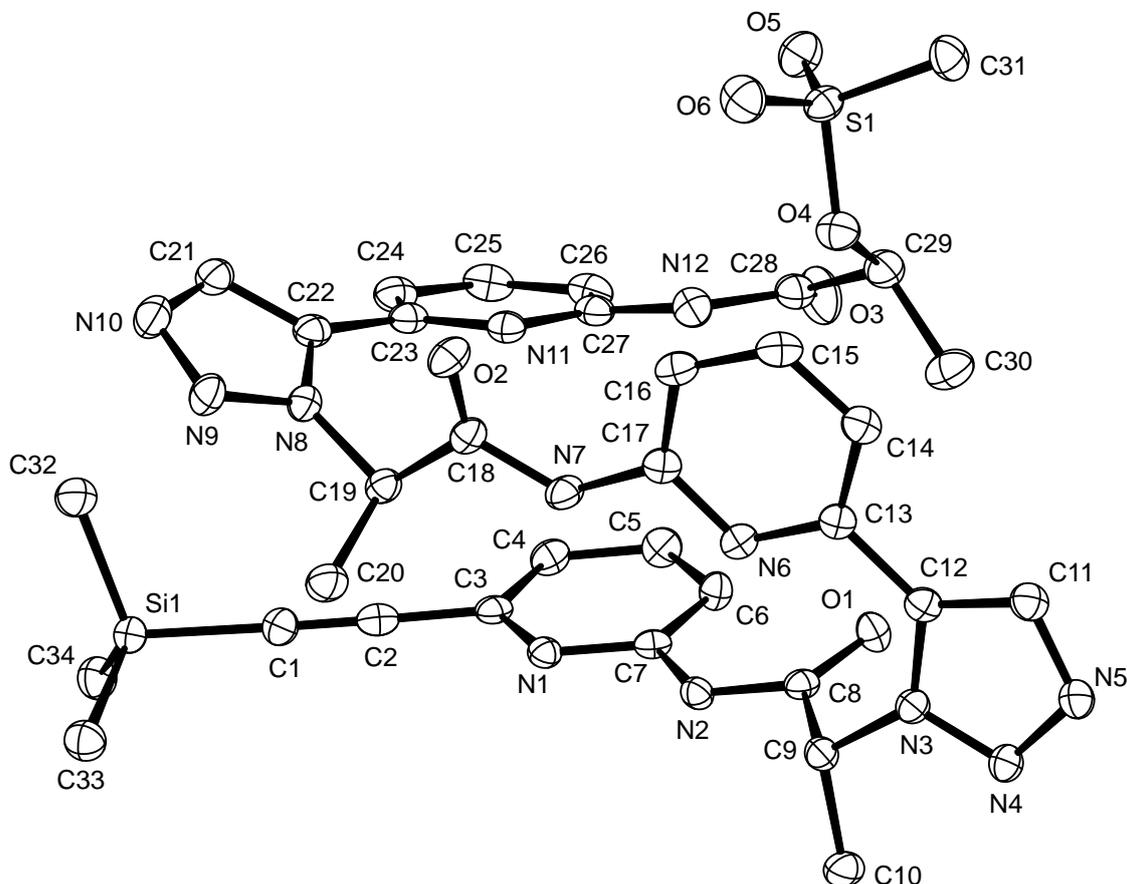
Structure of TMS-deprotected 6 in the crystal. Hydrogen atoms omitted for clarity. Ellipsoids are shown at the 50 % probability level.



Crystal Data for TMS-deprotected 6: $C_{21}H_{21}N_7O_5S$, $M_r = 483.51 \text{ g}\cdot\text{mol}^{-1}$, colorless needle from methanol/dichloromethane, crystal size $0.47 \times 0.09 \times 0.04 \text{ mm}^3$, monoclinic, space group $C2$ [No. 5], $a = 18.534(4) \text{ \AA}$, $b = 8.8696(17) \text{ \AA}$, $c = 13.761(3) \text{ \AA}$, $\beta = 97.078(8)^\circ$, $V = 2244.8(7) \text{ \AA}^3$, $T = 100 \text{ K}$, $Z = 4$, $D_{\text{calc}} = 1.431 \text{ g}\cdot\text{cm}^{-3}$, $\lambda = 1.54178 \text{ \AA}$, $\mu(\text{Cu-K}\alpha) = 1.711 \text{ mm}^{-1}$, Gaussian absorption correction ($T_{\text{min}} = 0.6510$, $T_{\text{max}} = 0.9353$), $3.24 < \theta < 66.47^\circ$, 24475 measured reflections, 3821 independent reflections ($R_{\text{int}} = 0.106$), 3169 reflections with $I > 2\sigma(I)$. Structure solved by direct methods (SHELXS) and refined by full-matrix least-squares (SHELXL) against F^2 to $R_1 = 0.041$ [$I > 2\sigma(I)$], $wR_2 = 0.092$ (all data), 311 parameters, H atoms riding, $S = 1.169$, residual electron density $+0.26/-0.33 \text{ e \AA}^{-3}$.

X-ray Crystal Structure Analysis: Compound 7

Structure of 7 in the crystal. Hydrogen atoms omitted for clarity. Ellipsoids are shown at the 50 % probability level.



Crystal Data for 7: $C_{34}H_{38}N_{12}O_6SSi$, $M_r = 770.91 \text{ g}\cdot\text{mol}^{-1}$, colorless prism from methanol/dichloromethane, crystal size $0.061 \times 0.042 \times 0.030 \text{ mm}^3$, tetragonal, space group $P4_3$ [No. 78], $a = 17.235(2) \text{ \AA}$, $c = 12.720(3) \text{ \AA}$, $V = 3778.4(11) \text{ \AA}^3$, $T = 100 \text{ K}$, $Z = 4$, $D_{\text{calc}} = 1.355 \text{ g}\cdot\text{cm}^{-3}$, $\lambda = 0.71073 \text{ \AA}$, $\mu(Mo-K\alpha) = 0.179 \text{ mm}^{-1}$, Gaussian absorption correction ($T_{\text{min}} = 0.9922$, $T_{\text{max}} = 0.9961$), $1.99 < \theta < 26.75^\circ$, 81307 measured reflections, 8005 independent reflections ($R_{\text{int}} = 0.060$), 7394 reflections with $I > 2\sigma(I)$. Structure solved by direct methods (SHELXS) and refined by full-matrix least-squares (SHELXL) against F^2 to $R_1 = 0.033$ [$I > 2\sigma(I)$], $wR_2 = 0.082$ (all data), 494 parameters, H atoms riding, absolute structure parameter = $-0.29(5)$, $S = 1.038$, residual electron density $+0.39/-0.24 \text{ e \AA}^{-3}$. Both the Flack parameter and the Hooft parameter are significantly negative (ca. -0.3) in spite of measuring the data with a high redundancy (Hooft, R.; Straver, L.; Spek, A. *J. Appl. Cryst.* **2008**, *41*, 96-103). We currently have no explanation for this, apart from

the fact that the normal probability plot has a slope of 1.49 (correlation coefficient 0.98) indicating that the standard uncertainties of the reflections may be underestimated due to averaging:

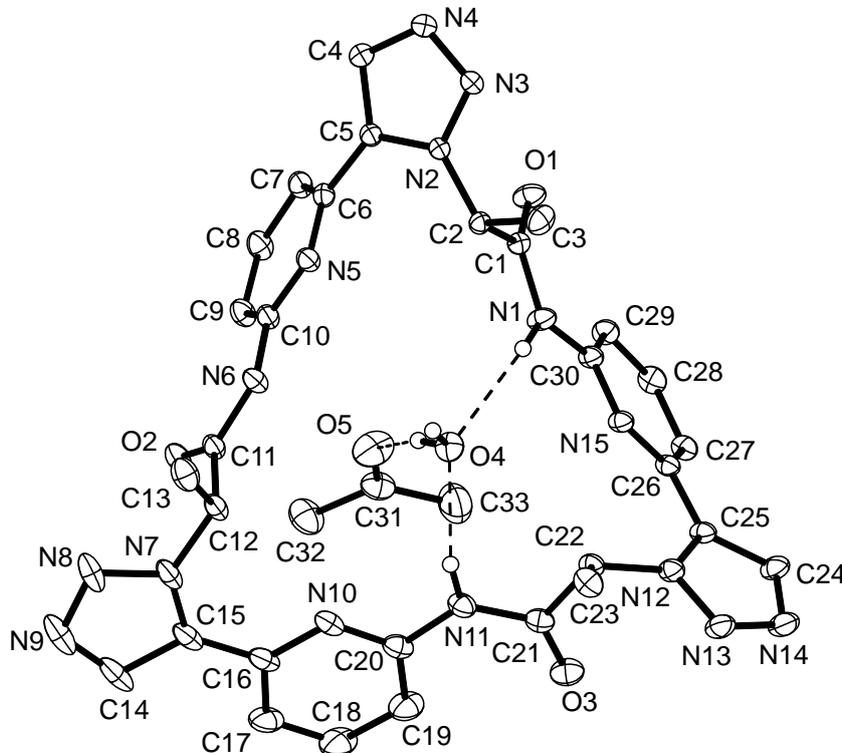
Intensity Statistics for Dataset

Resolution	#Data	#Theory	%Complete	Redundancy	Mean I	Mean I/s	R(int)	Rsigma
Inf - 2.20	211	212	99.5	22.57	85.2	87.17	0.0381	0.0101
2.20 - 1.73	215	215	100.0	23.87	32.7	79.96	0.0423	0.0109
1.73 - 1.50	228	228	100.0	23.61	21.6	69.06	0.0475	0.0122
1.50 - 1.35	223	223	100.0	23.71	14.2	59.04	0.0596	0.0140
1.35 - 1.25	228	228	100.0	23.01	14.6	55.52	0.0602	0.0147
1.25 - 1.17	236	236	100.0	22.07	14.1	51.52	0.0652	0.0159
1.17 - 1.11	233	233	100.0	21.17	10.9	42.51	0.0754	0.0185
1.11 - 1.06	221	221	100.0	20.12	10.4	38.97	0.0819	0.0206
1.06 - 1.01	278	278	100.0	19.54	9.2	36.58	0.0855	0.0224
1.01 - 0.97	258	258	100.0	18.86	6.1	27.68	0.1108	0.0299
0.97 - 0.94	240	240	100.0	18.09	5.9	25.02	0.1146	0.0321
0.94 - 0.91	250	250	100.0	17.70	4.6	21.17	0.1361	0.0396
0.91 - 0.88	297	297	100.0	17.05	4.2	19.14	0.1539	0.0446
0.88 - 0.86	216	216	100.0	16.46	4.3	18.54	0.1586	0.0464
0.86 - 0.84	258	258	100.0	16.18	3.7	16.22	0.1707	0.0525
0.84 - 0.82	259	259	100.0	15.81	3.2	14.04	0.1945	0.0614
0.82 - 0.80	294	294	100.0	15.24	2.9	12.49	0.2173	0.0698
0.80 - 0.79	58	66	87.9	7.82	3.5	10.82	0.1949	0.0953
0.89 - 0.79	1274	1282	99.4	15.63	3.5	15.33	0.1804	0.0573
Inf - 0.79	4203	4212	99.8	19.32	13.4	37.65	0.0633	0.0183

Merged [A], lowest resolution = 12.65 Angstroms, 4149 outliers downweighted

X-ray Crystal Structure Analysis: Compound 8·C₃H₆O·H₂O

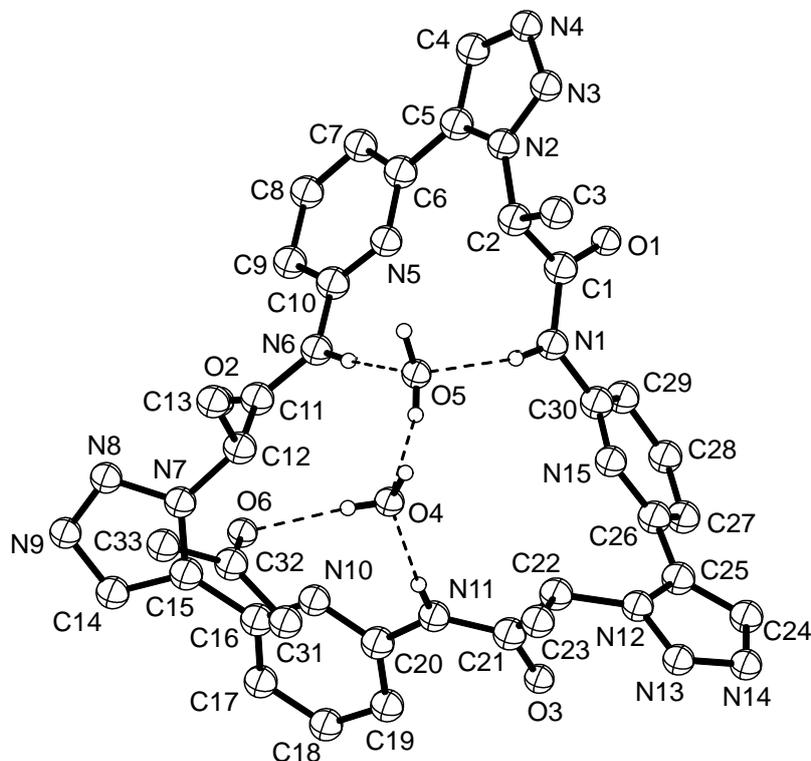
Structure of 8·C₃H₆O·H₂O in the crystal. Except for those involved in hydrogen bonding to water and acetone, hydrogen atoms omitted for clarity. Ellipsoids are shown at the 50 % probability level.



Crystal Data for 8·C₃H₆O·H₂O: C₃₀H₂₇N₁₅O₃·C₃H₆O·H₂O, $M_r = 721.76 \text{ g}\cdot\text{mol}^{-1}$, colorless prism from acetone/hexane/trace water, crystal size 0.11 x 0.05 x 0.04 mm³, monoclinic, space group P2₁ [No. 4], $a = 9.9601(9) \text{ \AA}$, $b = 9.3717(9) \text{ \AA}$, $c = 19.2123(18) \text{ \AA}$, $\beta = 101.469(2)^\circ$, $V = 1757.5(3) \text{ \AA}^3$, $T = 100 \text{ K}$, $Z = 2$, $D_{\text{calc}} = 1.364 \text{ g}\cdot\text{cm}^{-3}$, $\lambda = 0.71073 \text{ \AA}$, $\mu(\text{Mo-K}\alpha) = 0.098 \text{ mm}^{-1}$, Gaussian absorption correction ($T_{\text{min}} = 0.9911$, $T_{\text{max}} = 0.9967$), $2.43 < \theta < 32.89^\circ$, 103103 measured reflections, 6846 independent reflections ($R_{\text{int}} = 0.078$), 6524 reflections with $I > 2\sigma(I)$. Structure solved by direct methods (SHELXS) and refined by full-matrix least-squares (SHELXL) against F^2 to $R_1 = 0.037$ [$I > 2\sigma(I)$], $wR_2 = 0.106$ (all data), 489 parameters. The positions of the H atoms on the water molecule were located on a difference Fourier synthesis and the atoms were refined with isotropic atomic displacement parameters, otherwise H atoms riding, absolute structure was not determined (total number of reflections without merging Friedel pairs (MERG 3) is 12993), $S = 0.820$, residual electron density $+0.40/-0.20 \text{ e \AA}^{-3}$.

X-ray Crystal Structure Analysis: Compound 8·C₃H₆O·2H₂O

Structure of 8·C₃H₆O·2H₂O in the crystal. Except for those involved in hydrogen bonding to water and acetone and neighboring triazole N atoms in neighboring molecules (not shown), hydrogen atoms omitted for clarity. Ellipsoids are shown at the 50 % probability level.



Crystal Data for 8·C₃H₆O·2H₂O: C₃₀H₂₇N₁₅O₃·C₃H₆O·2H₂O, $M_r = 739.78 \text{ g}\cdot\text{mol}^{-1}$, colorless needle from water/acetone (1:1), crystal size 1.024 (see below) x 0.052 x 0.052 mm³, orthorhombic, space group P2₁2₁2₁ [No. 18], $a = 12.4653(5) \text{ \AA}$, $b = 17.1970(7) \text{ \AA}$, $c = 17.6158(8) \text{ \AA}$, $V = 3776.2(3) \text{ \AA}^3$, $T = 100 \text{ K}$, $Z = 4$, $D_{\text{calc}} = 1.301 \text{ g}\cdot\text{cm}^{-3}$, $\lambda = 1.54178 \text{ \AA}$, $\mu(\text{Cu-K}\alpha) = 0.785 \text{ mm}^{-1}$, Gaussian absorption correction ($T_{\text{min}} = 0.6789$, $T_{\text{max}} = 0.9666$), $3.59 < \theta < 54.35^\circ$, 63933 measured reflections, 4532 independent reflections ($R_{\text{int}} = 0.086$), 4203 reflections with $I > 2\sigma(I)$. Structure solved by direct methods (SHELXS) and refined by full-matrix least-squares (SHELXL) against F^2 to $R_1 = 0.056 [I > 2\sigma(I)]$, $wR_2 = 0.146$ (all data), 505 parameters. The positions of the four H atoms on the two water molecules were located on a difference Fourier synthesis and the atoms were refined with isotropic atomic displacement parameters, otherwise H atoms riding, absolute structure parameter = $-0.1(4)$, $S = 1.143$, residual electron density $+0.37/-0.22 \text{ e \AA}^{-3}$. The crystals grew as very fine needles, which were difficult to cut. In order to obtain adequate diffraction

intensity, it was necessary to choose a long crystal, since the longer crystals were thicker. In order to minimize the effects of a needle that was longer than the diameter of the X-ray beam, data were measured at the constant chi angle 54.74°, and the frames were scaled using SADABS. Data were collected with a high redundancy in order to reduce bias in the scaling process. Intensity statistics are shown below.

Intensity Statistics for Dataset

Resolution	#Data	#Theory	%Complete	Redundancy	Mean I	Mean I/s	R(int)	Rsigma
Inf - 3.96	70	70	100.0	6.60	104.9	20.94	0.0410	0.0454
3.96 - 2.62	163	163	100.0	11.26	49.9	27.46	0.0538	0.0392
2.62 - 2.07	223	223	100.0	23.17	24.1	40.19	0.0569	0.0223
2.07 - 1.80	232	232	100.0	23.08	17.5	38.71	0.0567	0.0229
1.80 - 1.63	232	233	99.6	21.00	13.1	33.20	0.0639	0.0256
1.63 - 1.52	220	220	100.0	19.92	8.7	30.42	0.0836	0.0282
1.52 - 1.43	227	227	100.0	19.12	7.2	26.13	0.1018	0.0322
1.43 - 1.35	259	259	100.0	17.91	6.2	22.24	0.1131	0.0355
1.35 - 1.30	192	192	100.0	16.68	5.8	22.73	0.1282	0.0372
1.30 - 1.24	261	261	100.0	16.39	5.8	19.95	0.1394	0.0391
1.24 - 1.20	234	234	100.0	14.81	5.6	19.10	0.1537	0.0441
1.20 - 1.16	242	242	100.0	12.49	5.0	16.53	0.1463	0.0489
1.16 - 1.13	195	196	99.5	11.53	6.1	17.37	0.1391	0.0486
1.13 - 1.10	237	237	100.0	10.86	5.6	15.92	0.1491	0.0518
1.10 - 1.07	253	256	98.8	10.85	4.1	13.94	0.1747	0.0581
1.07 - 1.05	196	197	99.5	9.89	3.1	11.50	0.1977	0.0762
1.05 - 1.02	310	310	100.0	9.82	3.1	11.43	0.1996	0.0742
1.02 - 1.00	234	242	96.7	9.24	2.6	10.55	0.2304	0.0816
1.00 - 0.98	218	222	98.2	8.72	2.5	9.71	0.2408	0.0896
0.98 - 0.97	138	138	100.0	8.43	1.7	7.66	0.2944	0.1169
0.97 - 0.95	219	306	71.6	3.55	1.8	5.75	0.2505	0.1664
1.05 - 0.95	1119	1218	91.9	7.77	2.4	9.33	0.2274	0.0956
Inf - 0.95	4555	4660	97.7	13.75	9.9	20.07	0.0849	0.0403