

Supplementary Material

for

Latondaine-1-Amino-Hydantoin Hybrid, Triazole-Fused Latondaine Schiff Bases and Their Metal Complexes: Synthesis, X-ray and Electron Diffraction, Molecular Docking Studies and Antiproliferative Activity

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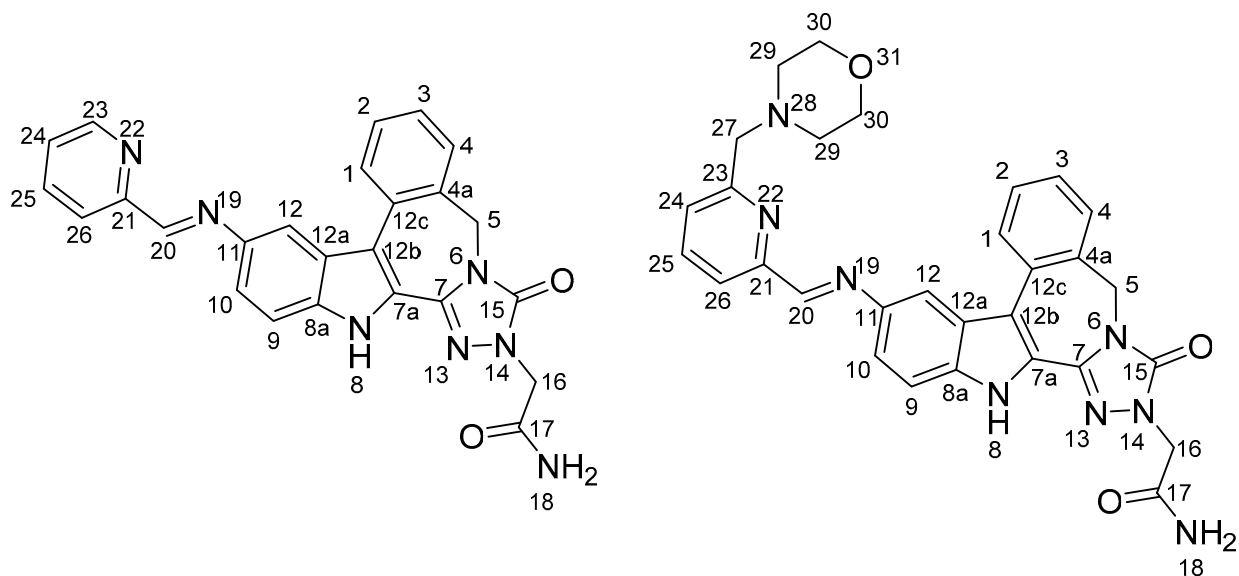
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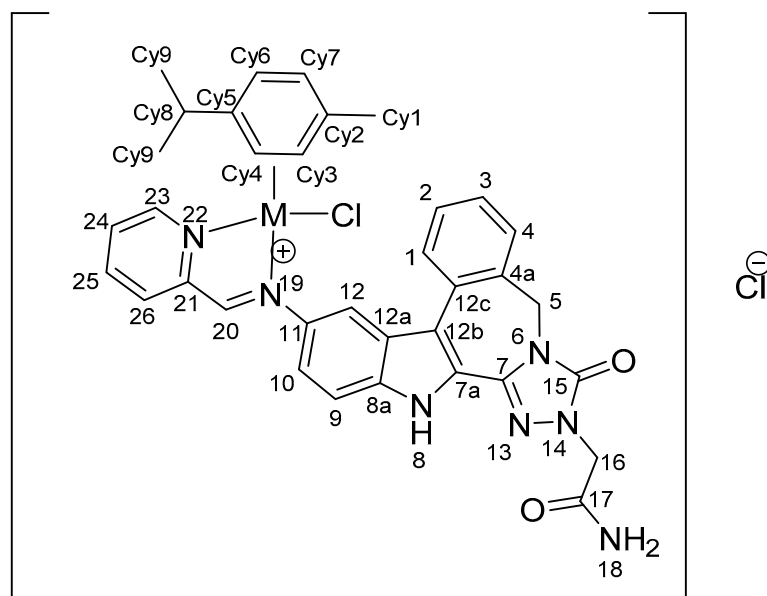
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Numbering schemes



Scheme S1. Numbering scheme of **HL**¹ and **HL**².



Scheme S2. Numbering scheme for complexes **1** (M = Ru) and **2** (M = Os).

Table S1. Yields and analytical data for **HL¹** and **HL²**, and complexes **1–3**.

		B	HL¹	HL²	1	2	3
Yield (%)		60	69	71	91	76	89
Brutto formula		C ₁₉ H ₁₄ N ₆ O ₄ - 0.8H ₂ O- 0.4CH ₃ OH	C ₂₅ H ₁₉ N ₇ O ₂ - 1.1H ₂ O	C ₃₀ H ₂₈ N ₈ O ₃ - 2H ₂ O	C ₃₅ H ₃₃ N ₇ O ₂ RuCl ₂ - 2.5H ₂ O	C ₃₅ H ₃₃ N ₇ O ₂ OsCl ₂ - 2.4H ₂ O	C ₃₀ H ₂₈ N ₈ O ₃ CuCl ₂ - H ₂ O·0.3C ₃ H ₈ O
M_r		417.58	469.28	584.62	800.69	888.05	719.09
C (%)	calcd	55.79	63.98	61.63	52.5	47.33	51.61
	found	55.92	63.78	61.85	52.52	47.65	51.51
H (%)	calcd	4.15	4.55	5.51	4.78	4.29	4.54
	found	3.98	4.3	5.21	4.52	4.05	4.35
N (%)	calcd	20.12	20.89	19.16	12.24	11.04	15.58
	found	19.76	20.71	19.26	11.85	11.01	15.36
O (%)	calcd			9.56		7.92	
	found			9.96		8.11	
ESI-MS	[M + H]⁺	391.17	450.25	549.33			
	[M + Na]⁺		472.22				
	[M - Cl]⁺				720.21	810.26	646.2
X-ray structure		YES	YES				YES

Crystallographic data obtained from both X-ray diffraction and electron diffraction techniques

Table S2. Crystallographic data and refinement details for B·CH₃OH and HL¹·2DMF and 3.

Compound	B·CH ₃ OH	HL ¹ ·2DMF	3
empirical formula	C ₂₀ H ₁₈ N ₆ O ₅	C ₃₁ H ₃₃ N ₉ O ₄	C ₃₀ H ₂₈ Cl ₂ CuN ₈ O ₃
fw	422.40	595.66	683.05
space group	monoclinic, <i>C2/c</i>	triclinic, <i>P</i> $\bar{1}$	monoclinic, <i>P2₁/n</i>
<i>a</i> , Å	34.647(14)	8.7154(7)	11.16(3)
<i>b</i> , Å	11.809(5)	11.8767(10)	9.602(13)
<i>c</i> , Å	19.252(7)	15.5370(16)	26.92(7)
α , °	90	111.185(5)	90
β , °	98.83(2)	95.529(4)	94.6(5)
γ , °	90	90.420(4)	90
<i>V</i> [Å ³]	7784(5)	1491.0(2)	2877(11)
<i>Z</i>	16	2	4
λ [Å]	0.71073	0.71073	0.02508
ρ_{calcd} , g cm ⁻³	1.442	1.327	1.577
cryst size, mm ³	0.44 × 0.13 × 0.04	0.20 × 0.20 × 0.20	0.005 × 0.0013 × 0.0005
<i>T</i> [K]	100(2)	100(2)	148(1)
μ , mm ⁻¹	0.107	0.092	
<i>R</i> ₁ ^a	0.0616	0.0577	0.1985
<i>wR</i> ₂ ^b	0.1569	0.1655	0.5049
GOF ^c	0.956	1.011	1.514

^a $R_1 = \Sigma ||F_o| - |F_c|| / \Sigma |F_o|$. ^b $wR_2 = \{\Sigma [w(F_o^2 - F_c^2)^2] / \Sigma [w(F_o^2)^2]\}^{1/2}$. ^c $\text{GOF} = \{\Sigma [w(F_o^2 - F_c^2)^2] / (n - p)\}^{1/2}$, where *n* is the number of reflections and *p* is the total number of parameters refined.

Synthesis of C*

To species **C** (225 mg, 0.61 mmol) in MeOH (20 mL) triethylamine (1 mL) was added. The solution was stirred at 80 °C for 3 h. The precipitate was filtered off after cooling to room temperature and washed with Et₂O (1 mL). The product was dried to give a light-beige solid. Yield: 156 mg. ¹H NMR (500 MHz, DMSO-*d*₆) δ 11.88 (s, 1H), 7.88 (t, *J* = 7.4 Hz, 1H), 7.56 (d, *J* = 7.4 Hz, 2H), 7.50 (td, *J* = 7.7, 1.2 Hz, 1H), 7.35 – 7.26 (m, 2H), 7.24 (t, *J* = 6.8 Hz, 1H), 7.14 (d, *J* = 1.7 Hz, 1H), 6.72 (dt, *J* = 11.8, 5.9 Hz, 1H), 4.81 (s, 2H), 4.69 (s, 2H), 4.36 (s, 2H).

UV-vis stability data

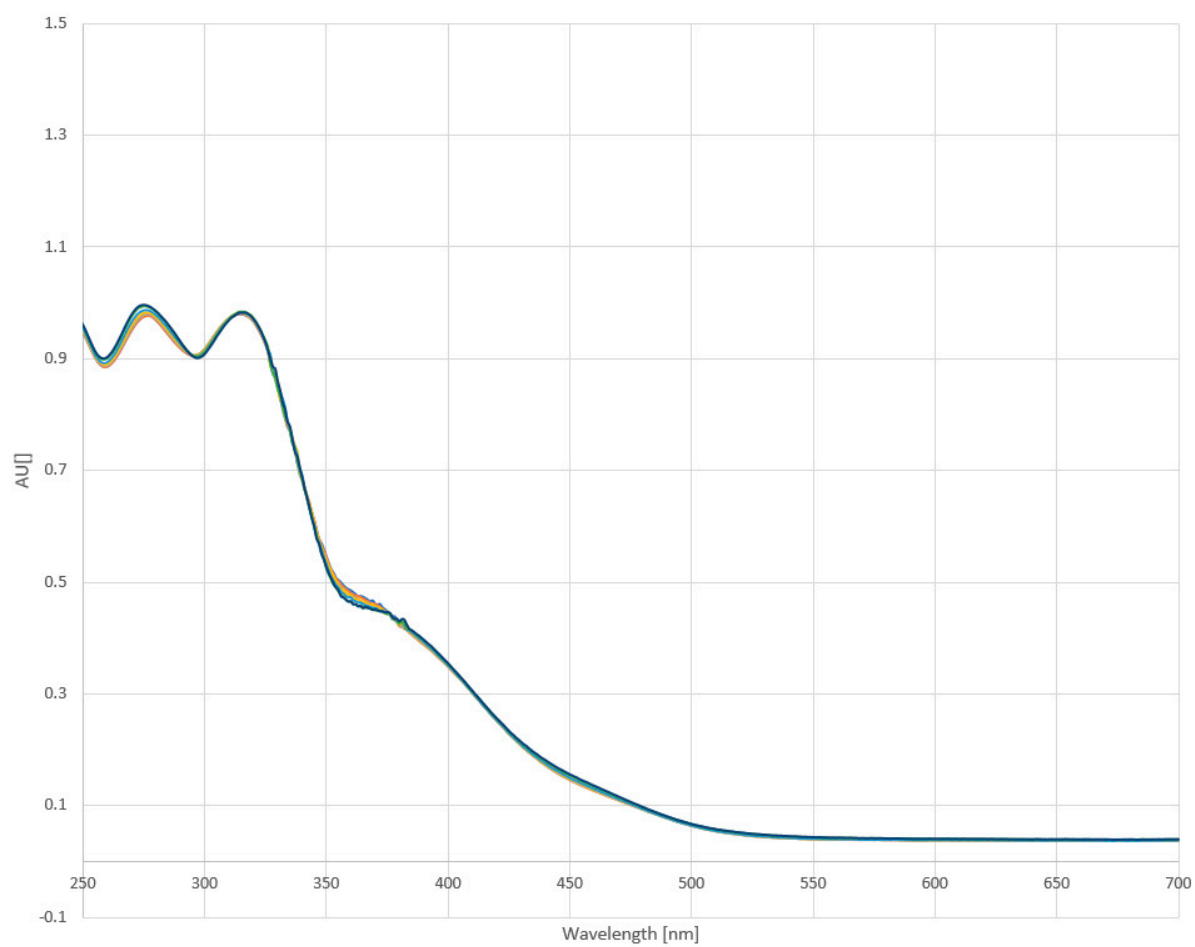


Figure S1. UV-vis absorption spectra of **1** (35 μ M) in DMSO/H₂O 1/99 monitored over 72 h.

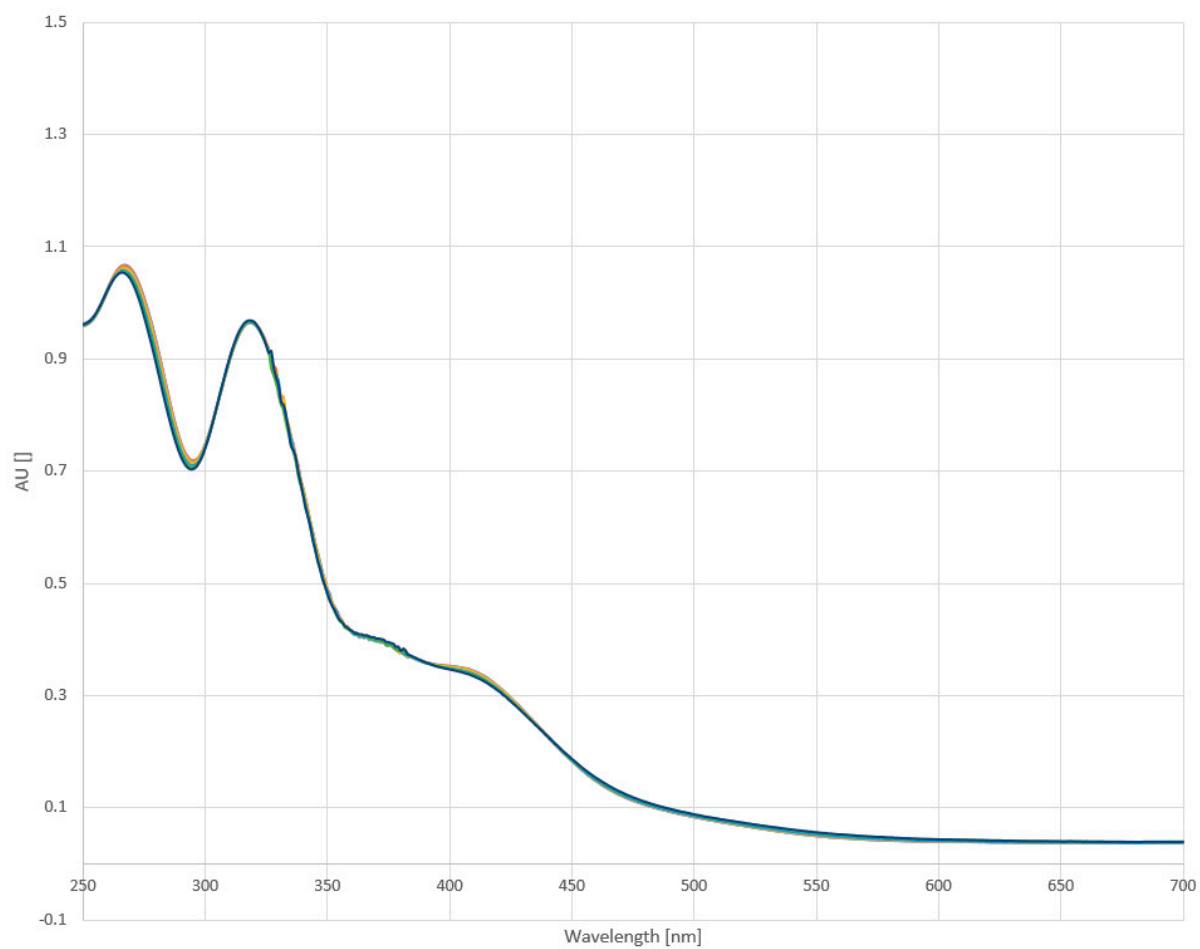


Figure S2. UV–vis absorption spectra of **2** (37 μM) in DMSO/H₂O 1/99 monitored over 72 h.

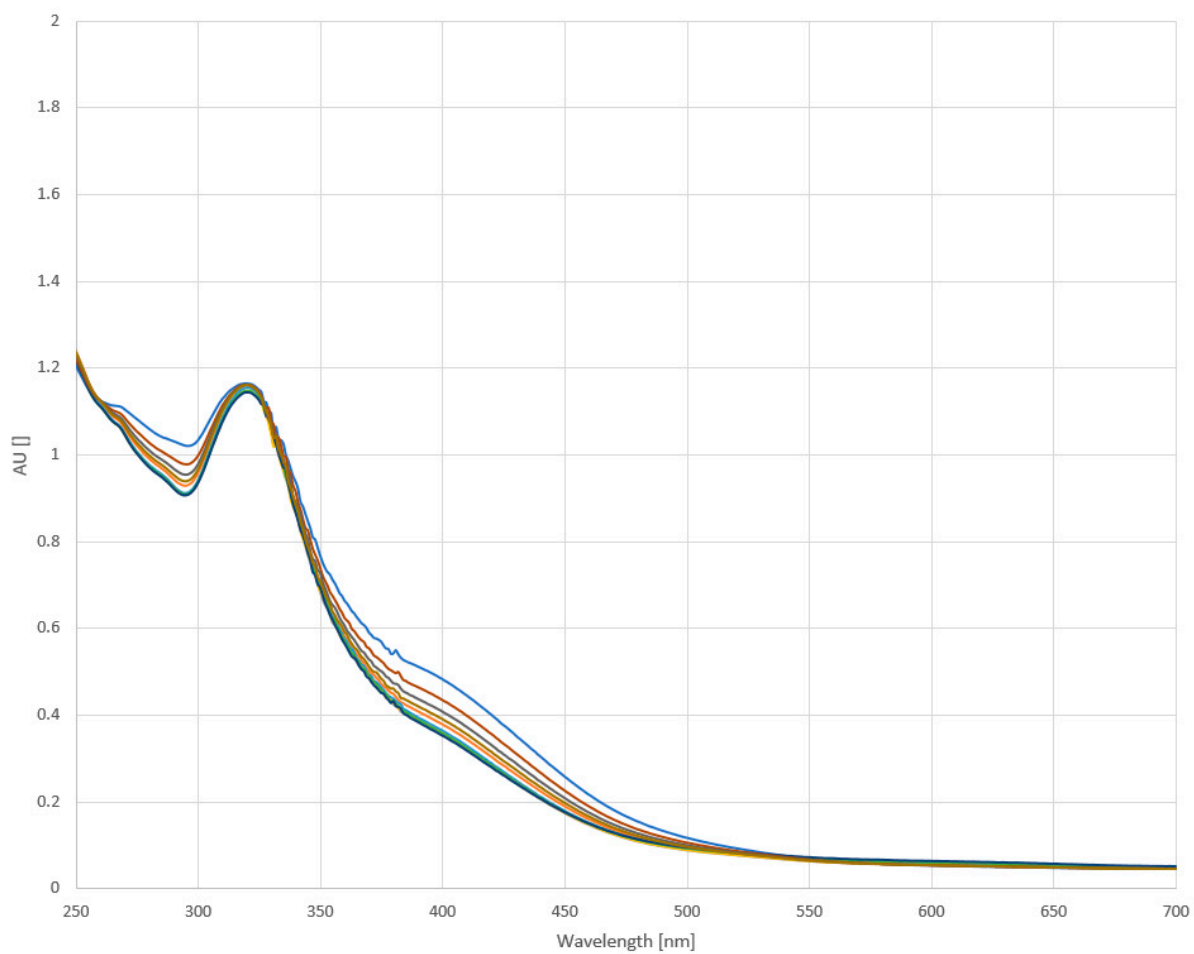


Figure S3. UV-vis absorption spectra of **3** (49 μM) in DMSO/H₂O 1/99 monitored over 72 h.

NMR spectra

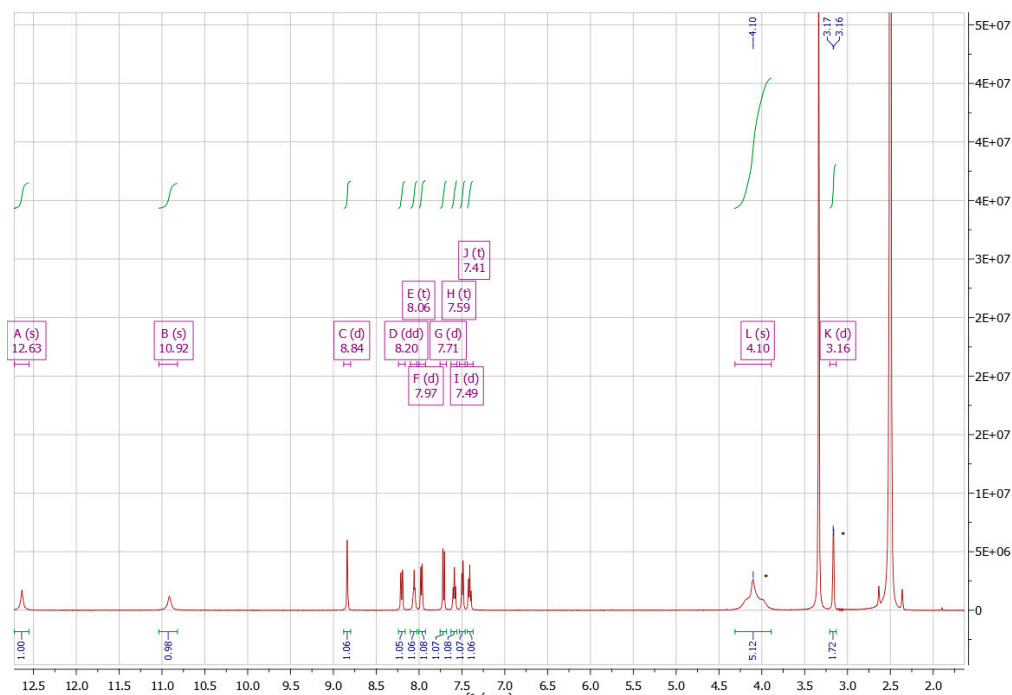


Figure S4. ^1H NMR spectrum of **B**. Residual MeOH is marked with an asterisk (*).

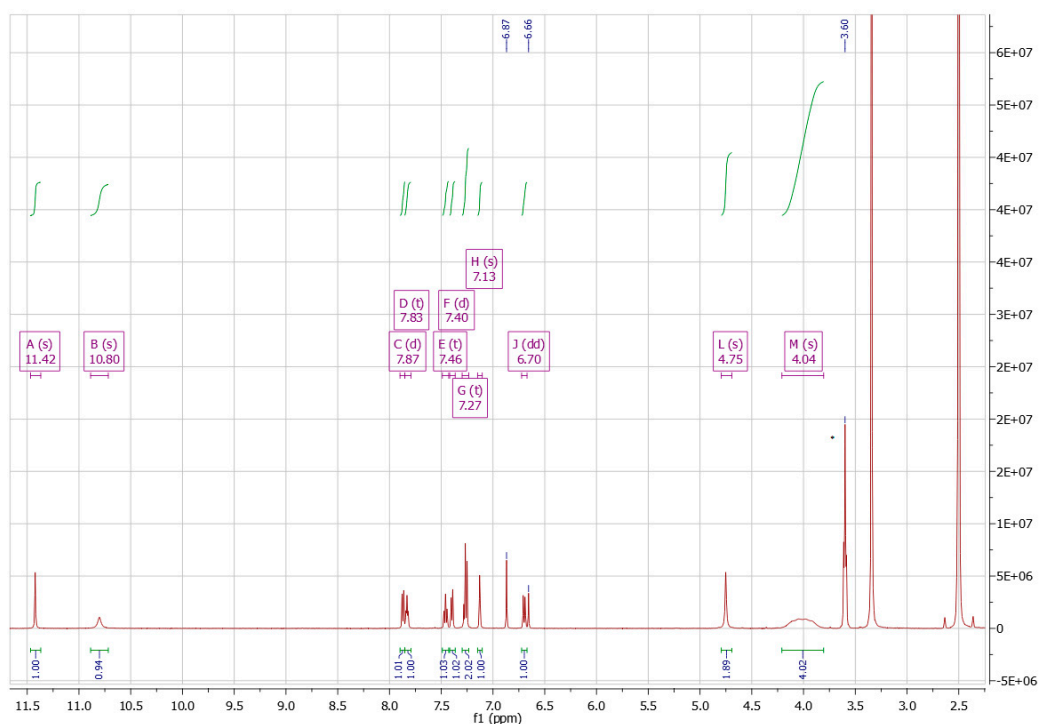


Figure S5. ^1H NMR spectrum of **C**. Residual THF is marked with an asterisk (*).

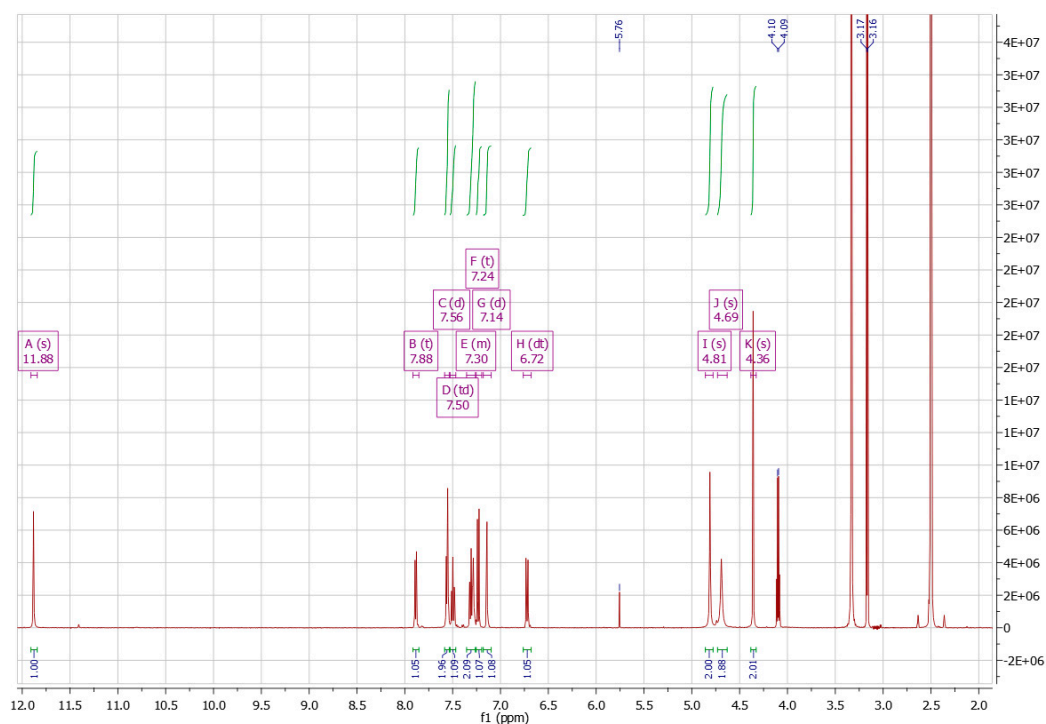


Figure S6. ^1H NMR spectrum of **C***. Residual solvents are identified by their respective chemical shifts on the top.

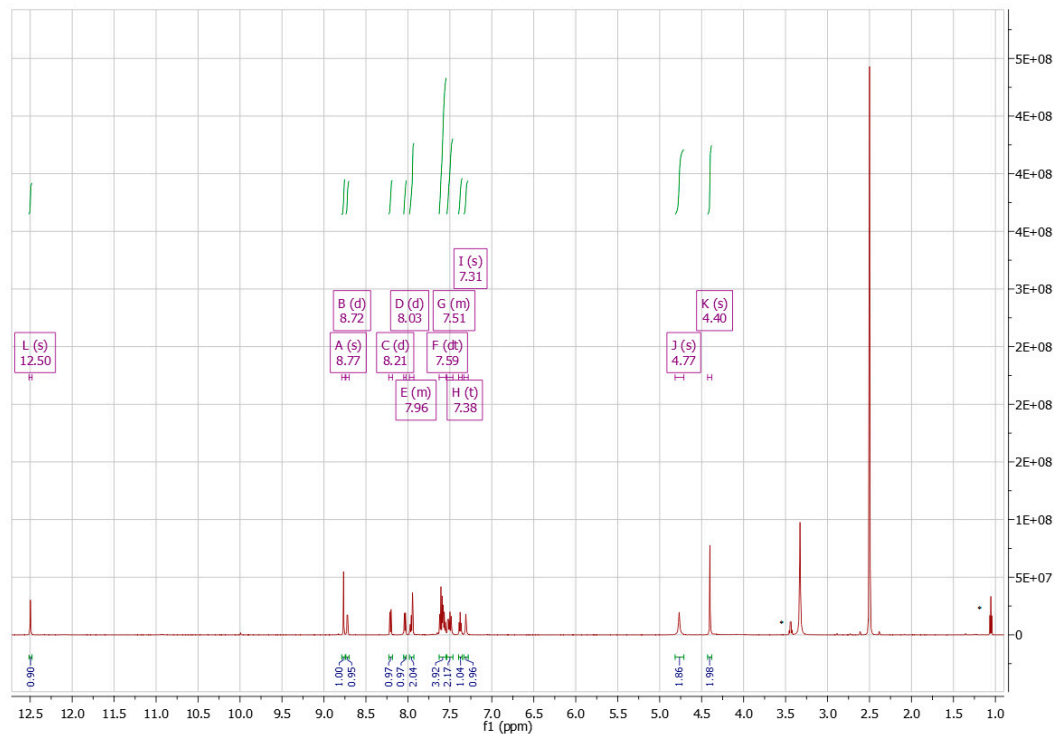


Figure S7. ^1H NMR spectrum of **HL¹**. Residual EtOH is marked with an asterisk (*).



Figure S8. ^1H NMR spectrum of **HL**². Residual EtOH is marked with an asterisk (*).

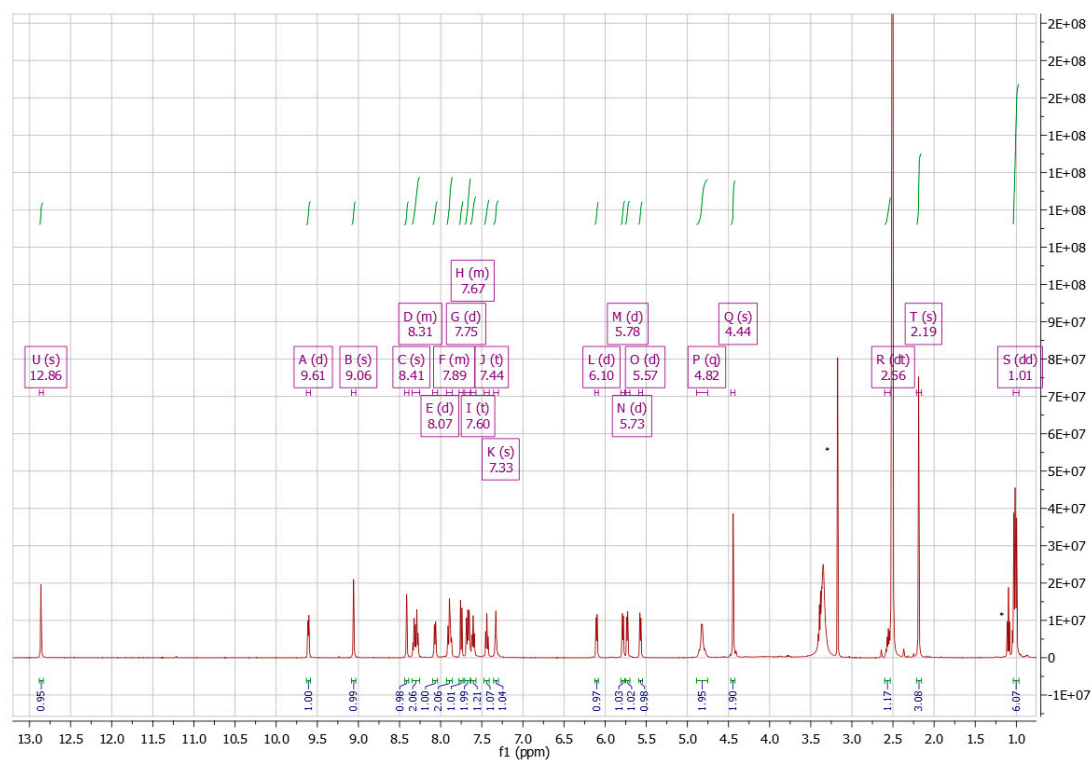


Figure S9. ^1H NMR spectrum of **1**. Residual $i\text{PrOH}$ is marked with an asterisk (*).

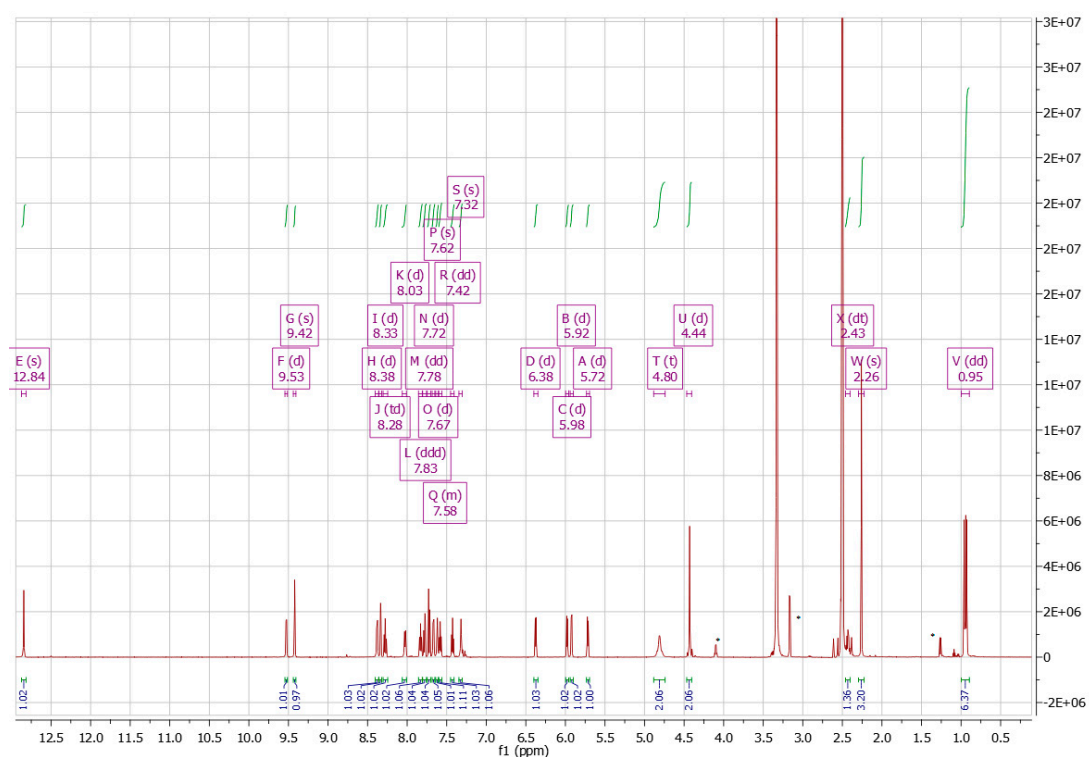


Figure S10. ^1H NMR spectrum of **2**. Residual $i\text{PrOH}$ is marked with an asterisk (*).

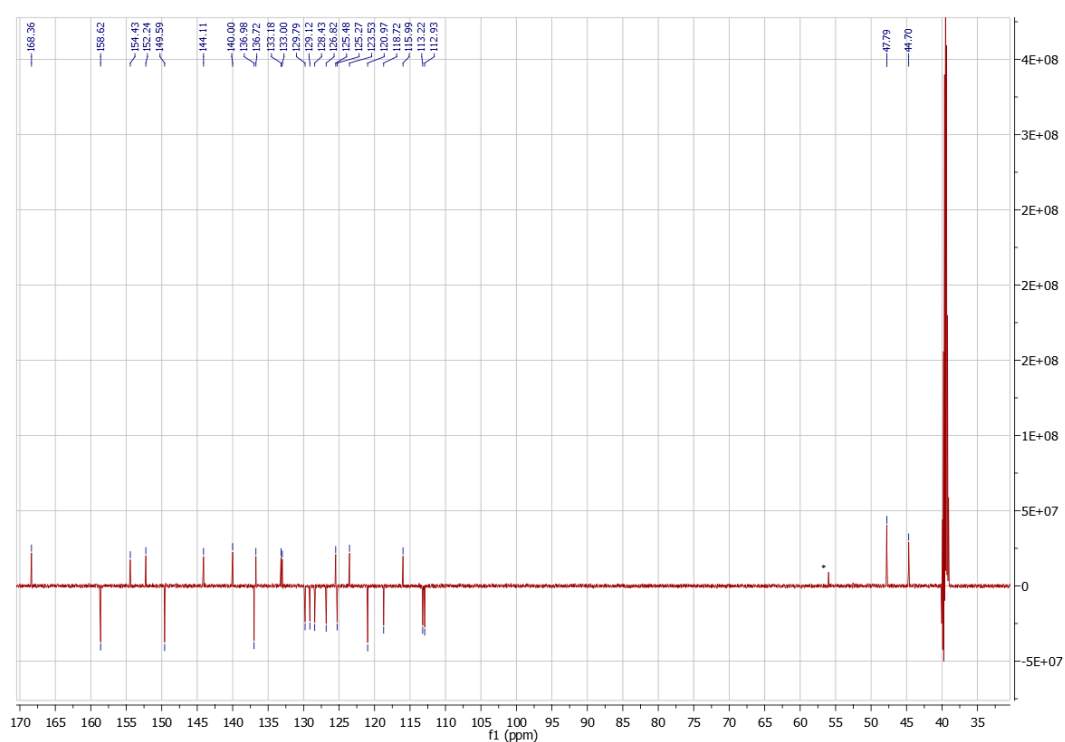


Figure S11. ^{13}C NMR spectrum of **HL¹**. Residual EtOH is marked with an asterisk (*).

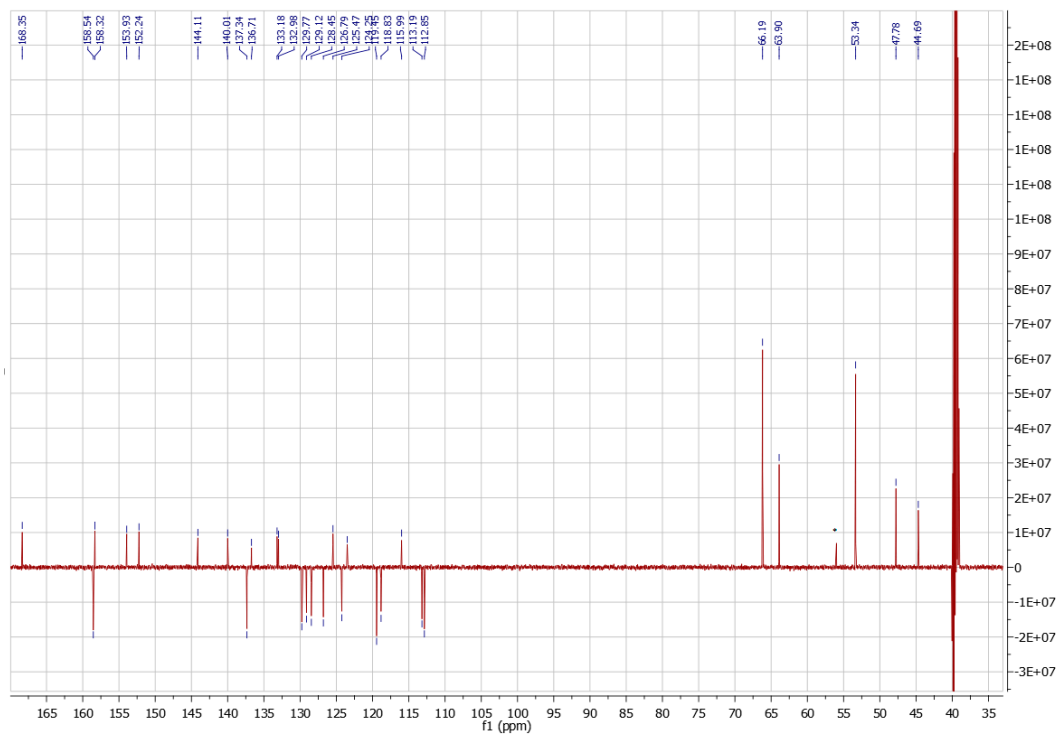


Figure S12. ¹³C NMR spectrum of **HL**². Residual EtOH is marked with an asterisk (*).

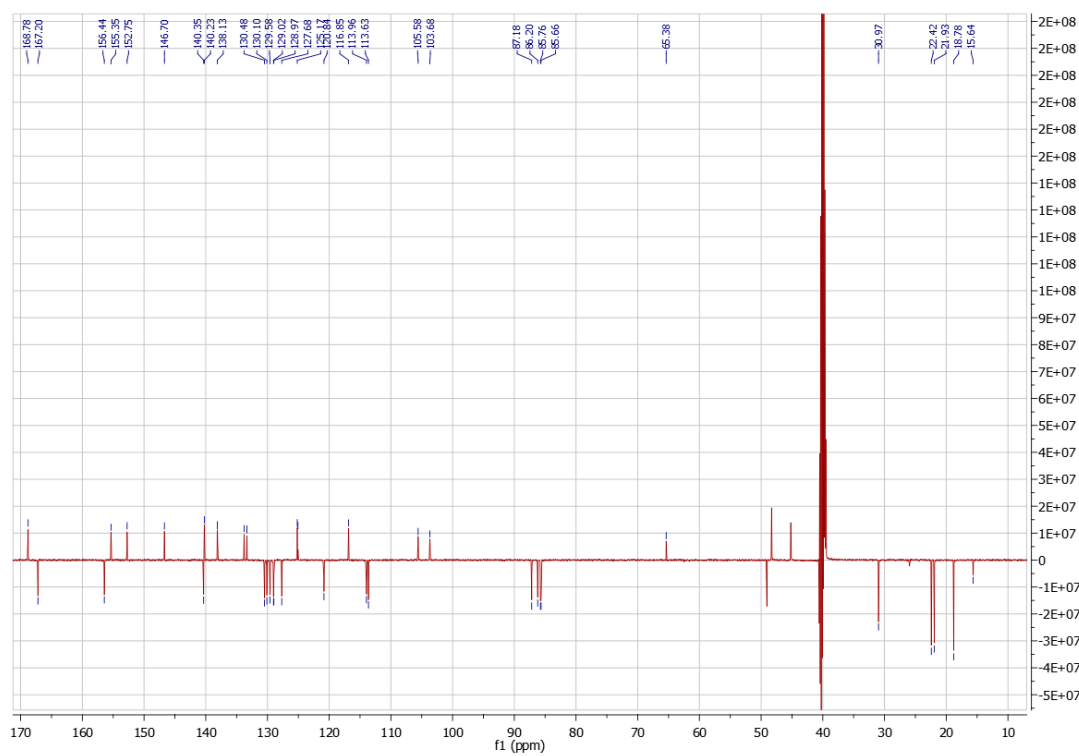


Figure S13. ¹³C NMR spectrum of **1**.

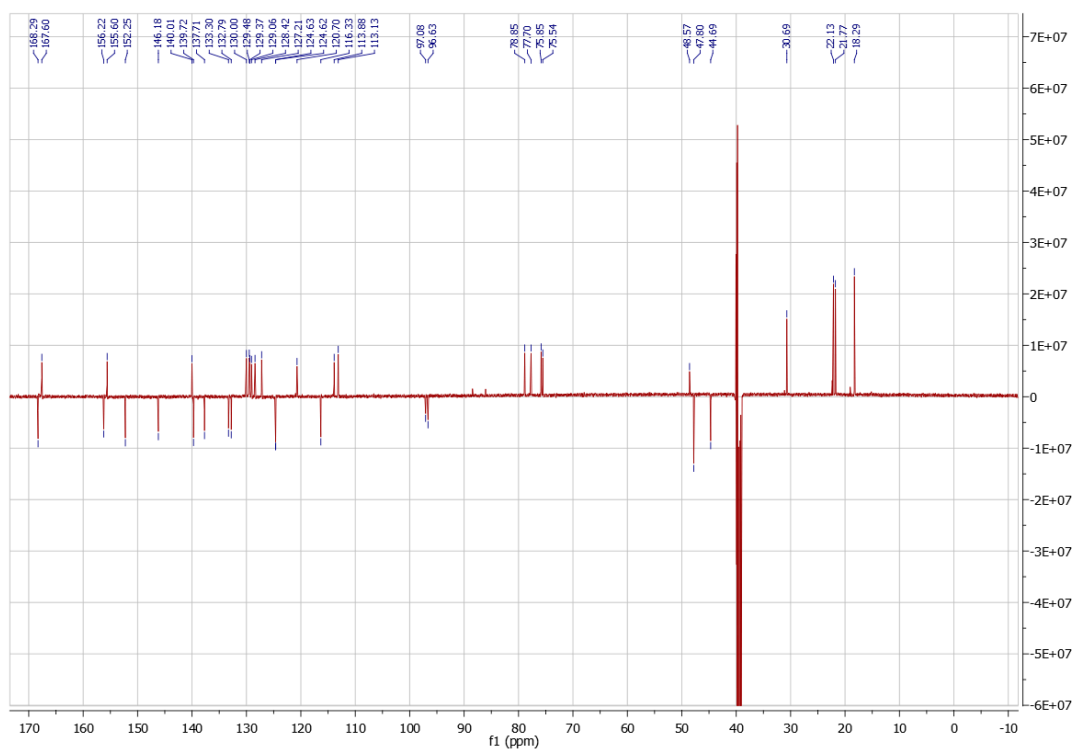


Figure S14. ^{13}C NMR spectrum of **2**.

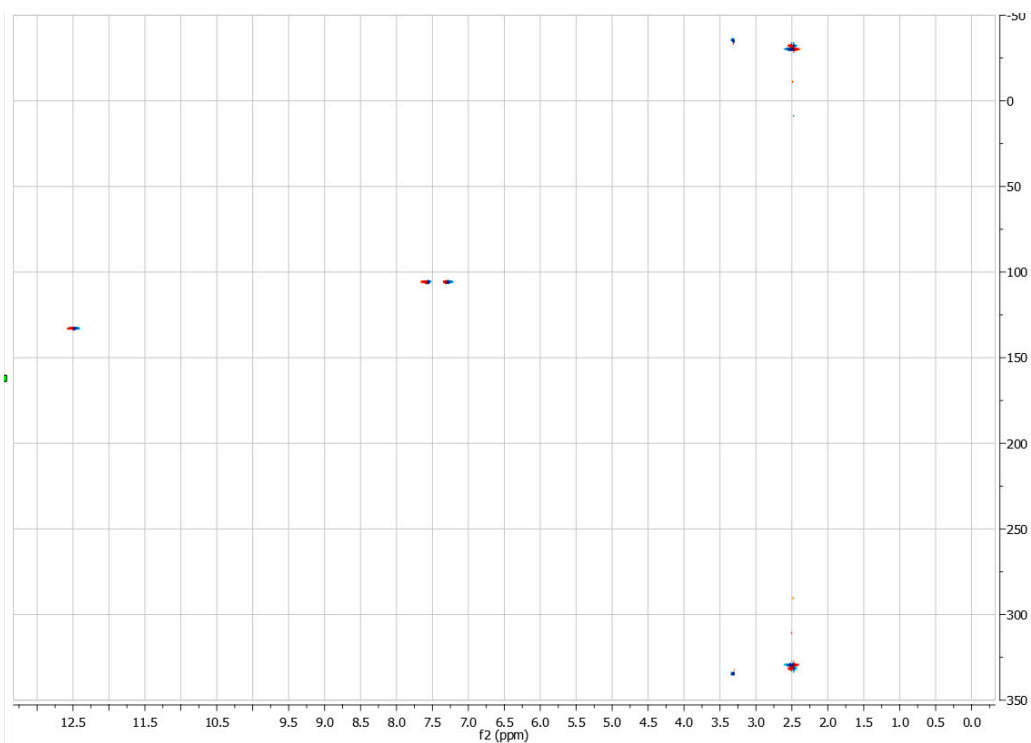


Figure S15. ^{15}N - ^1H HSQC NMR spectrum of **HL¹**.

Antiproliferative Activity

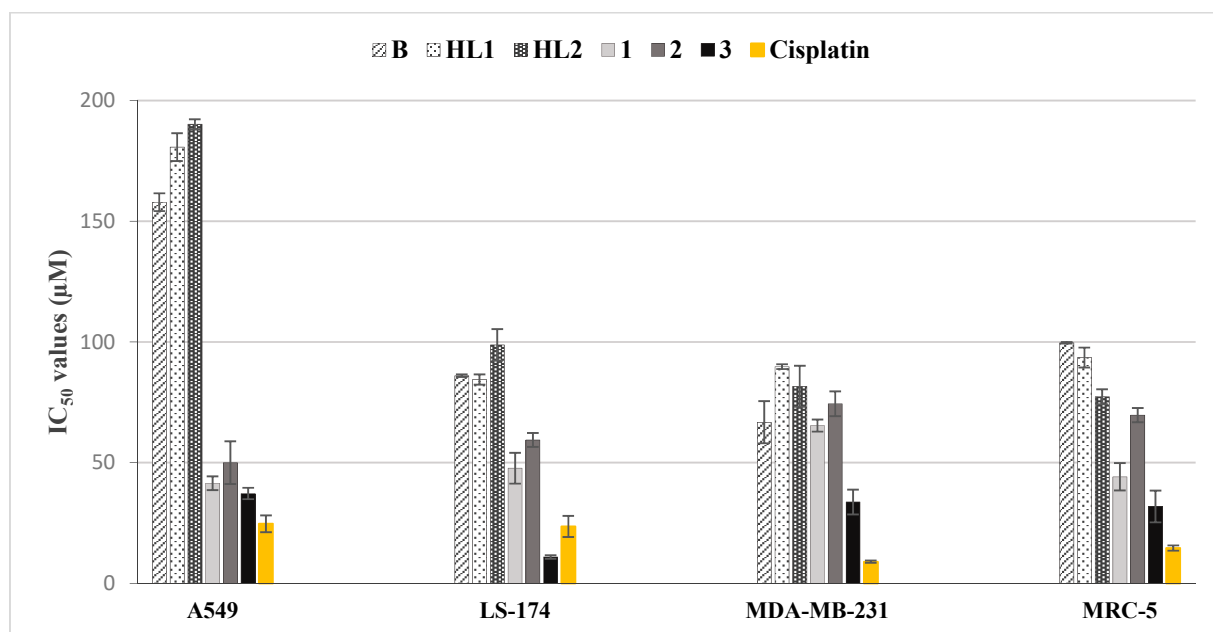


Figure S16. Bar graphs present IC₅₀ values (μM) of tested complexes and ligands vs cisplatin in three human tumor cell lines (A549, LS-174, MDA-MB-231) and non-tumor MRC-5 cells. Each IC₅₀ value presents average of at least three independent experiments ± SD, obtained for 72 h action using MTT assay.

ESI-mass spectra

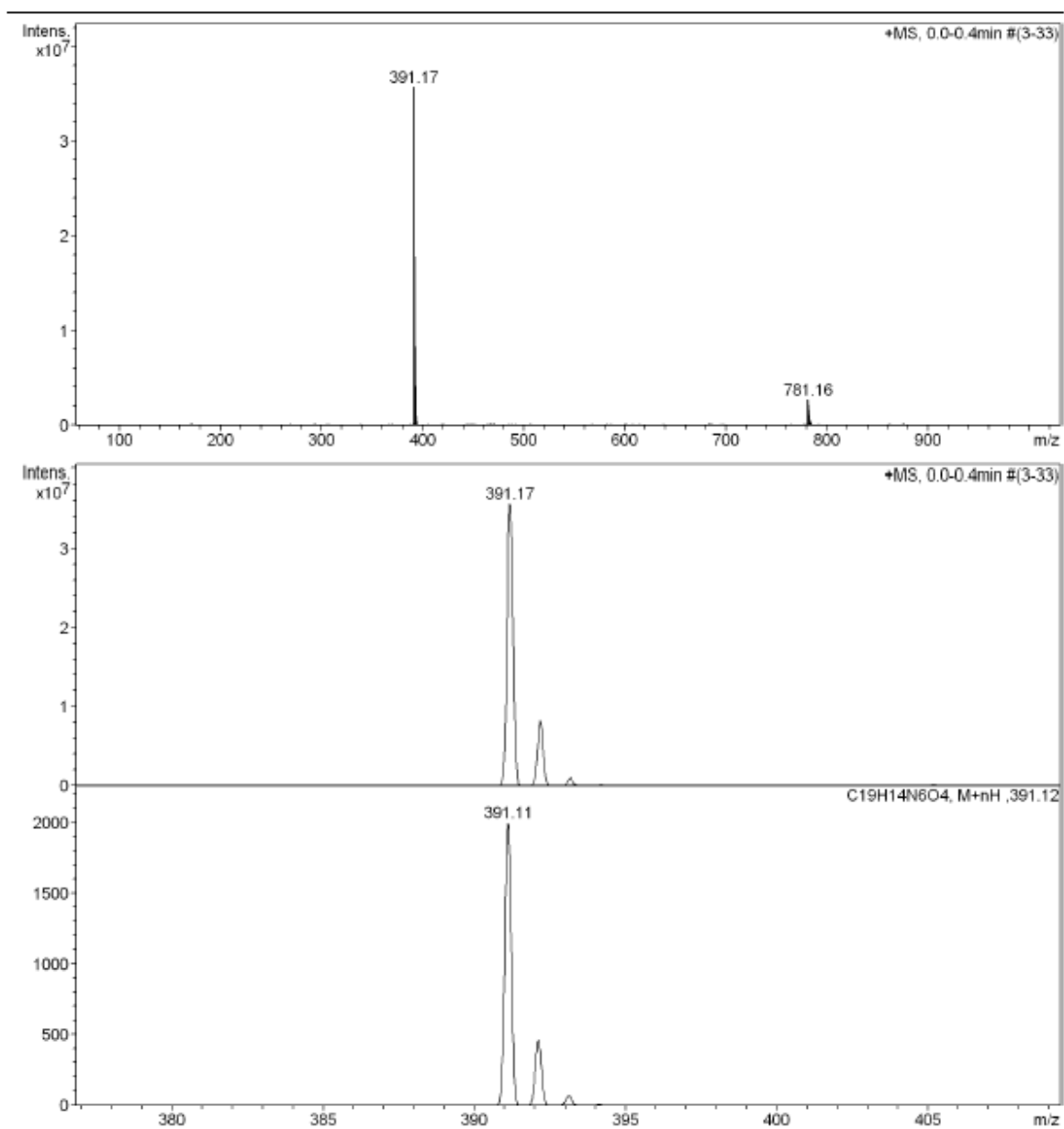


Figure S17. Positive ion ESI-mass spectrum of **B**.

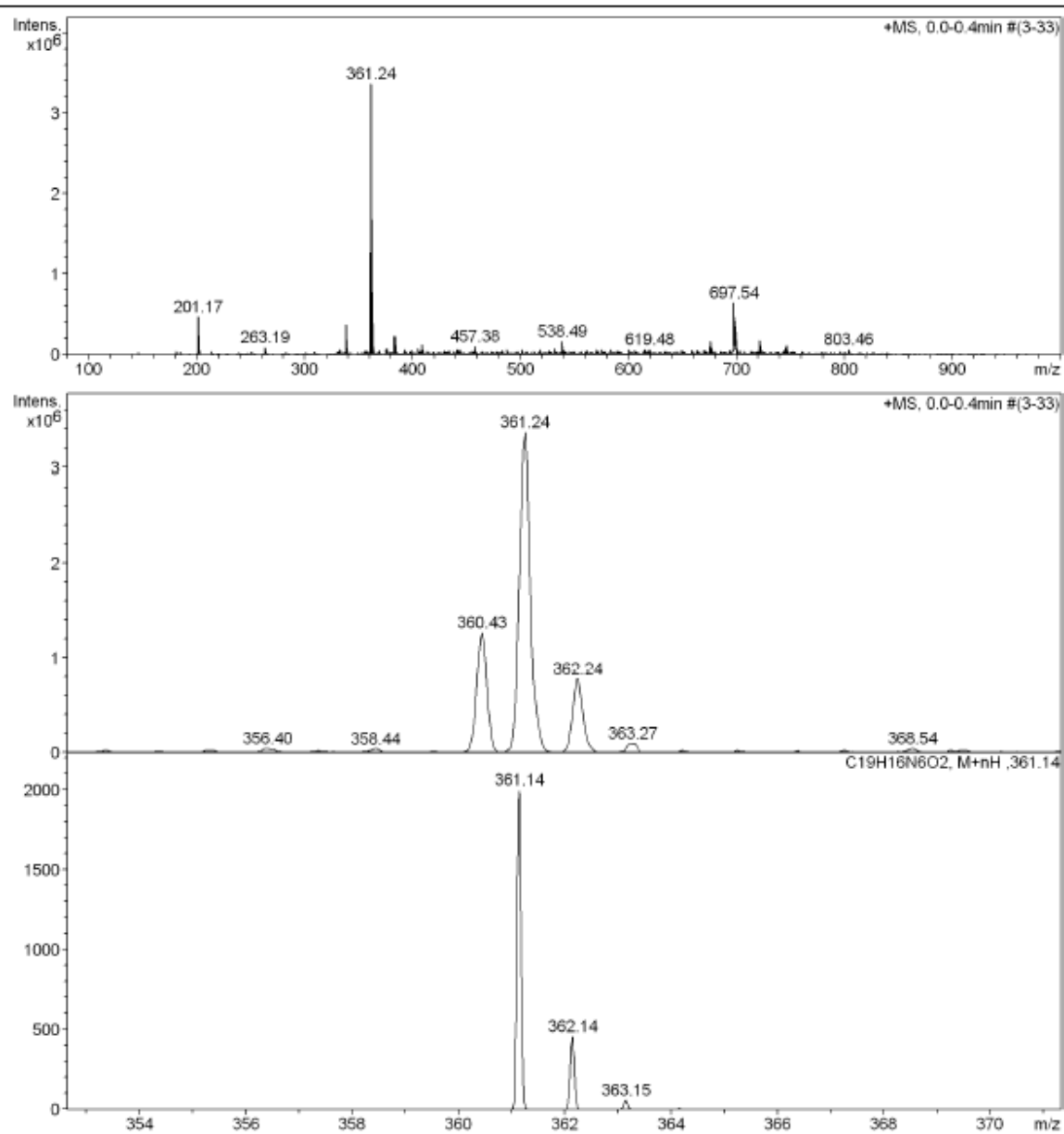


Figure S18. Positive ion mode of ESI-mass spectrum of C.

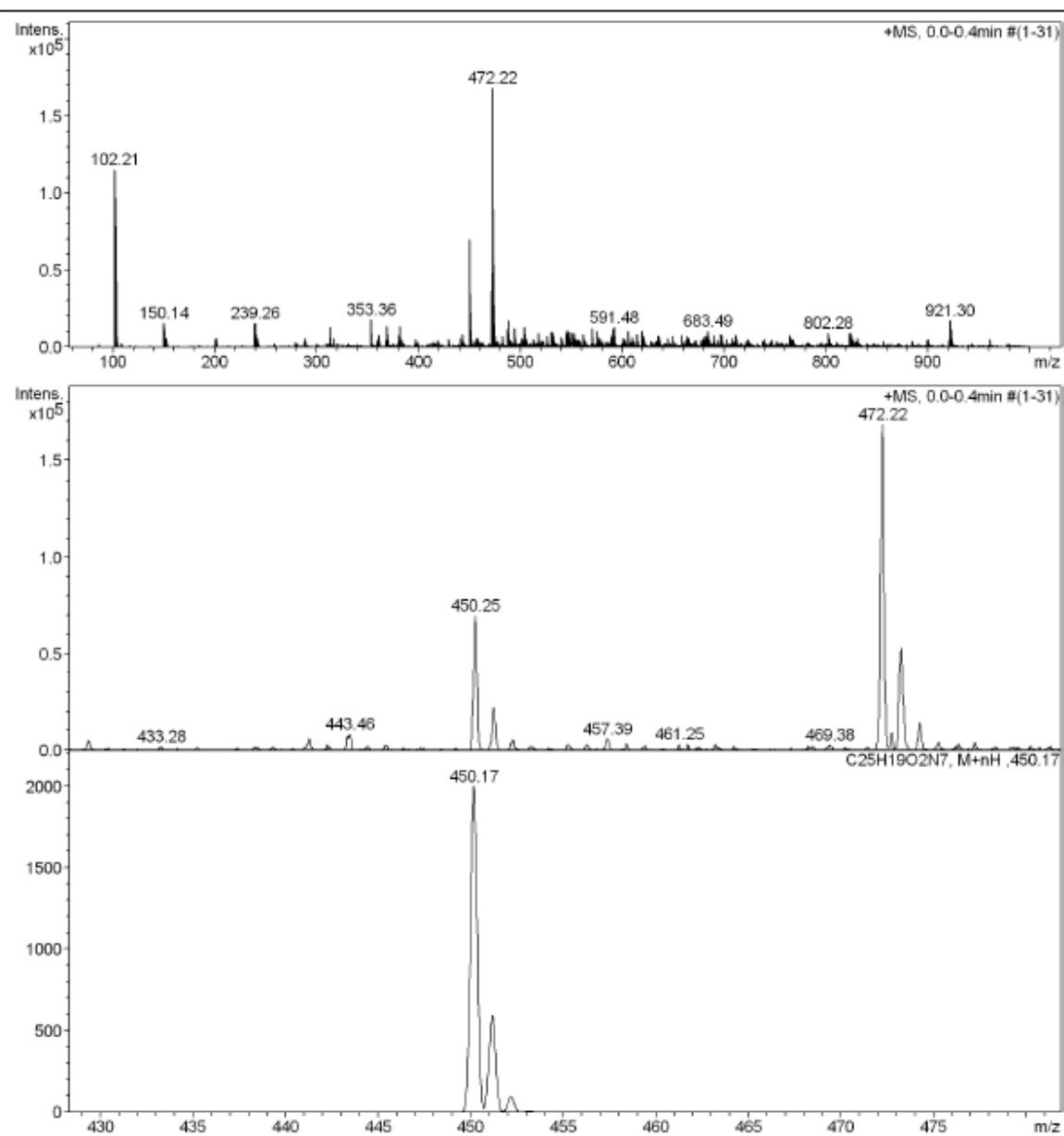


Figure S19. Positive ion mode ESI-mass spectrum of **HL**¹.

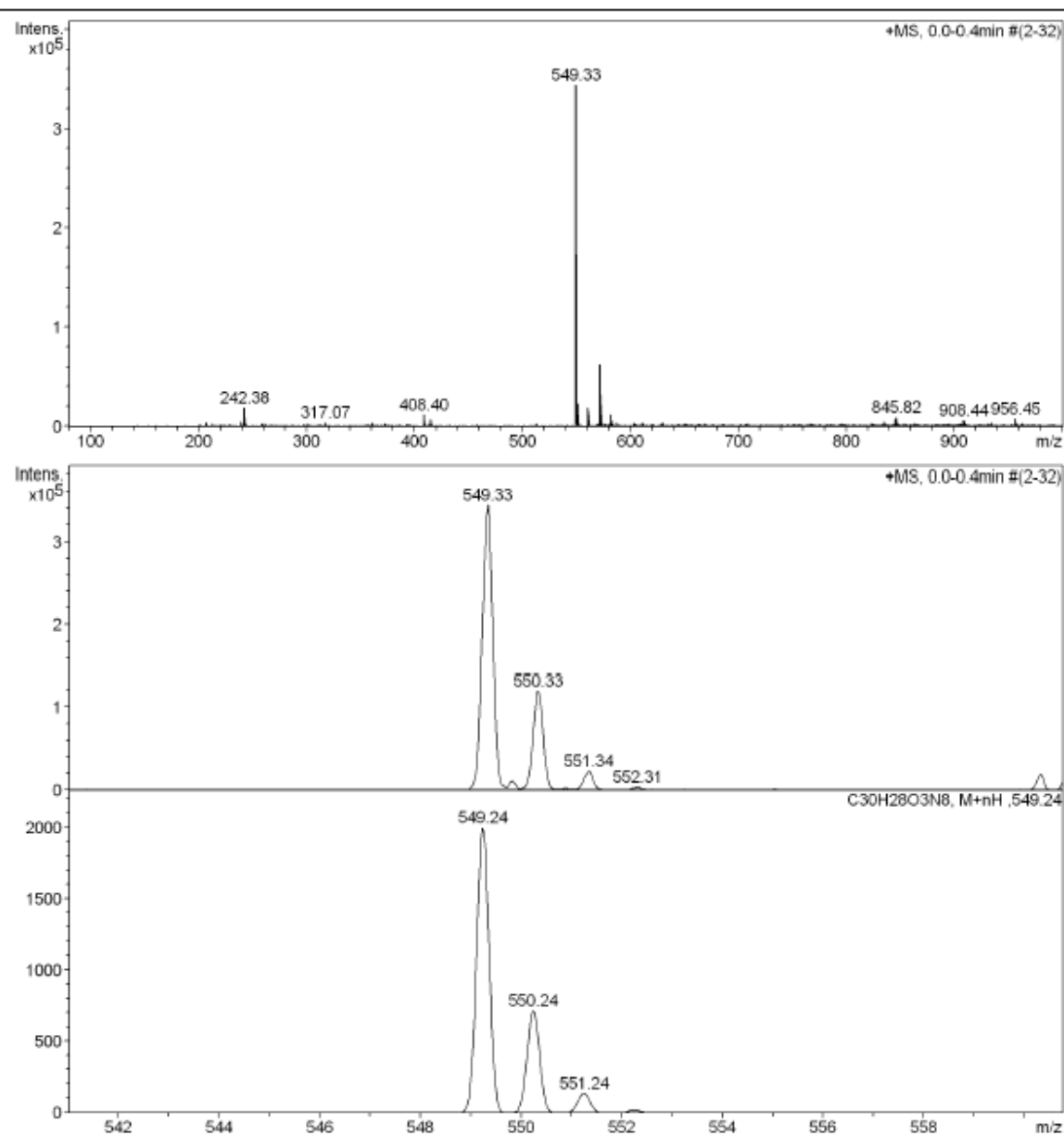


Figure S20. Positive ion mode ESI-mass spectrum of **HL²**.

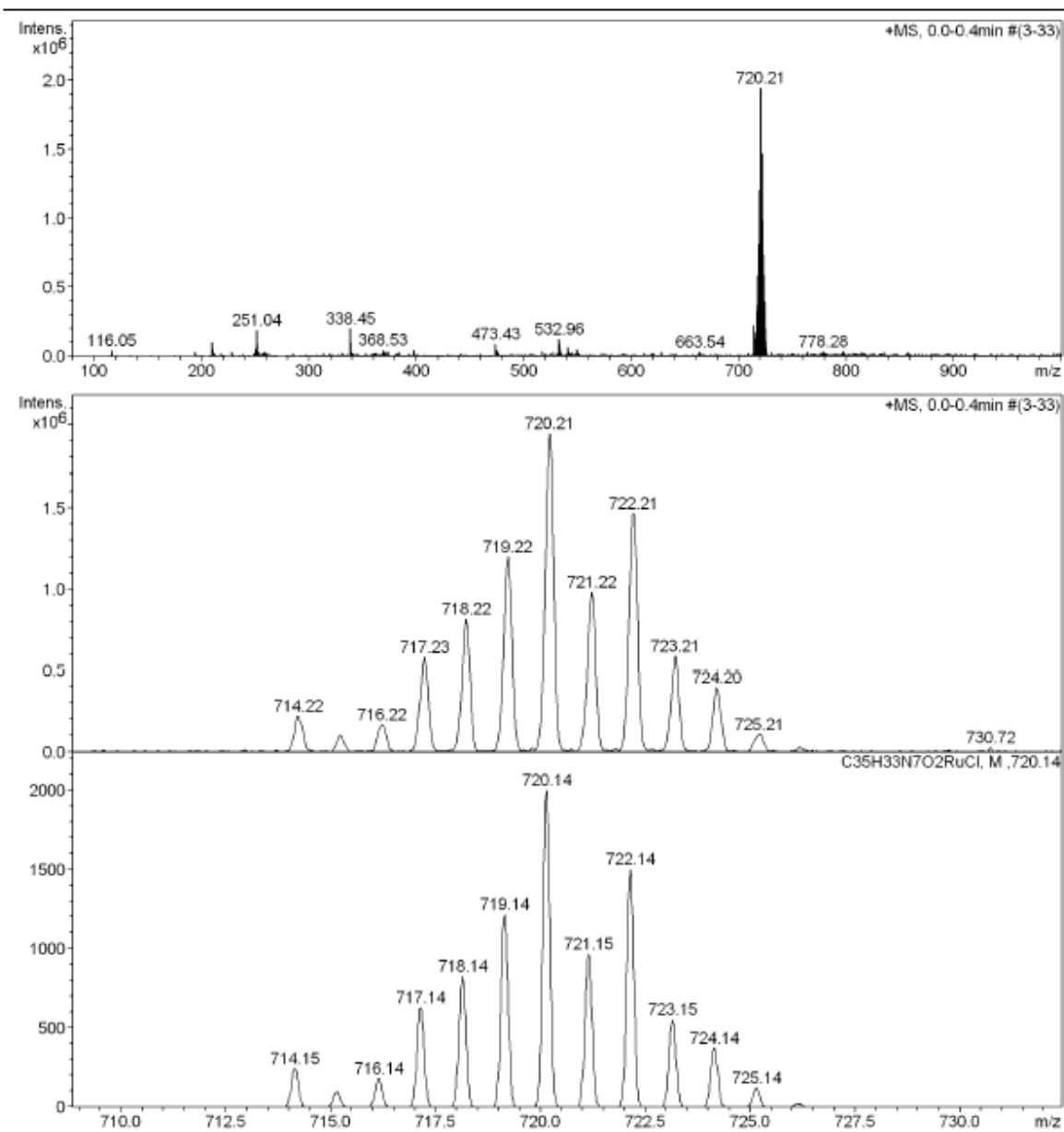


Figure S21. Positive ion mode ESI-mass spectrum of **1**.

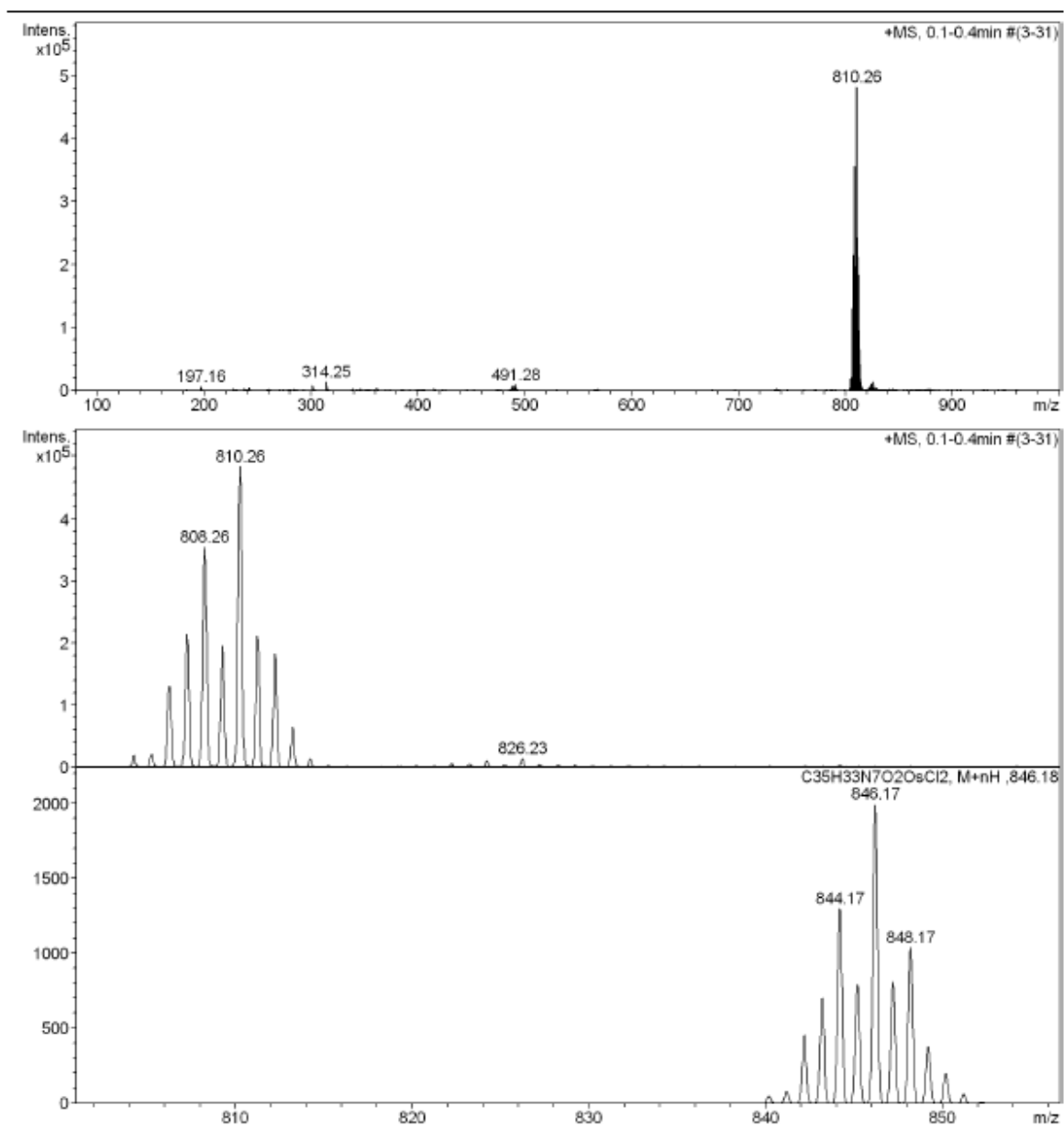


Figure S22. Positive ion mode ESI-mass spectrum of **2**.

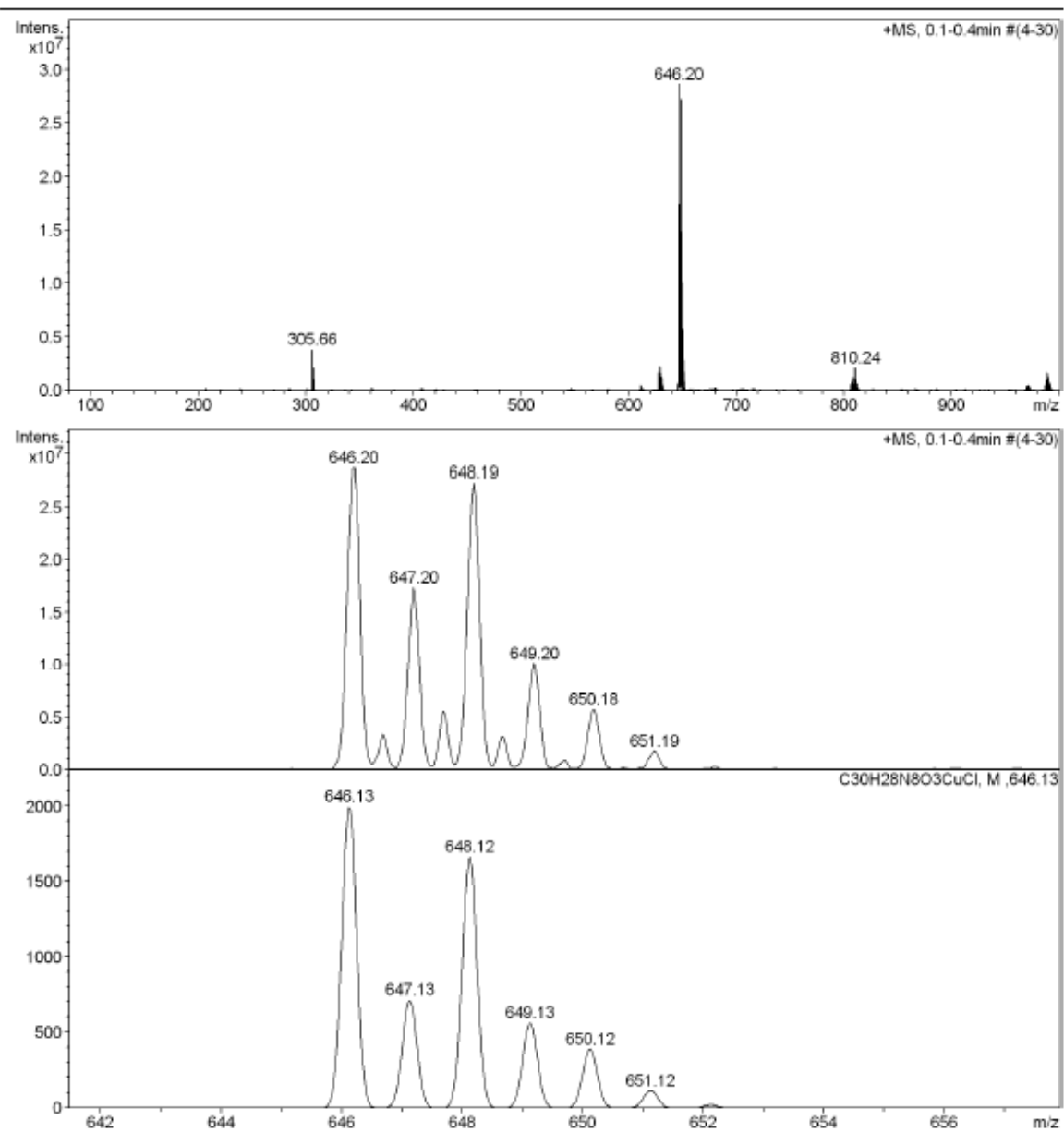


Figure S23. Positive ion mode ESI-mass spectrum of **3**.

Molecular Docking

Table S3. The binding affinities as predicted by the scoring functions for the Pim1 kinase. **LI7** is the co-crystallized ligand. Root-mean-square deviation – RMSD from the co-crystallized ligand (heavy atoms) in Å.

Ligands	GS	CS	ChemPLP	ASP
B	51.6	32.3	69.6	40.0
A	44.9	33.7	66.5	35.8
LI7	50.5	31.4	58.0	28.4
RMSD:	0.9020	3.1517	0.5348	6.7059

Table S4. The molecular descriptors as calculated by QikProp and their corresponding Known Drug Indexes 2a and 2b (KDI_{2a/2b}).

	RB	MW(g/mol)	HD	HA	Log P	PSA (Å ²)	KDI _{2A}	KDI _{2B}
B	3	390.4	3	4.5	2.4	162.3	4.92	0.20
A	2	293.3	2	3.5	2.0	98.3	5.19	0.40

Table S5. Definition of lead-like, drug-like and Known Drug Space (KDS) in terms of molecular descriptors. The values given are the maxima for each descriptor for the volumes of chemical space used.

	Lead-like Space	Drug-like Space	Known Drug Space
Molecular weight (g mol ⁻¹)	300	500	800
Lipophilicity (Log P)	3	5	6.5
Hydrogen bond donors (HD)	3	5	7
Hydrogen bond acceptors (HA)	3	10	15
Polar surface area (Å ²) (PSA)	60	140	180
Rotatable bonds (RB)	3	10	17

Table S6. The single point and corrected zero-point vibrational energies (ZPE) of **B** in hartrees (a.u.).

Systems	Energy/a.u.	ZPE/a.u. ^a
B	−1362.24233	0.31450644
B^{•+}	−1361.9785	0.31396035
B^{•−}	−1362.2934	0.31071621
Fragment 1	−985.962007	0.24248527
Fragment 2	−376.173285	0.06496719

Corrected according to Wong¹