

Synthesis, physicochemical characterization and antiproliferative activity of phosphino Ru(II) and Ir(III) complexes.

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Herein we present the synthesis of new complexes based on ruthenium(II) (Ru(η^6 -*p*-cymene)Cl₂PPh₂CH₂OH (RuPOH), Ru(η^6 -*p*-cymene)Cl₂P(*p*-OCH₃Ph)₂CH₂OH (RuMPOH)) and iridium(III) (Ir(η^5 -Cp*)Cl₂P(*p*-OCH₃Ph)₂CH₂OH (IrMPOH), Ir(η^5 -Cp*)Cl₂PPh₂CH₂OH (IrPOH) containing phosphine ligands with/without methoxy motif on phenyl rings (P(*p*-OCH₃Ph)₂CH₂OH (MPOH) and PPh₂CH₂OH (POH)). The complexes were characterized by mass spectrometry, NMR spectroscopy (1D: ¹H, ¹³C{¹H}, ³¹P{¹H}, 2D: HMQC, HMBC, COSY NMR) and elemental analysis. All the complexes were structurally identified by single-crystal X-ray diffraction analysis. The Ru(II) and Ir(III) complexes have typical piano-stool geometry with a η^6 -coordinated arene (Ru^{II} complexes) or η^5 -coordinated (Ir^{III} compounds) and three additional sites of ligation occupied by two chloride ligands and the phosphine ligand. Oxidation of NADH to NAD⁺ with high efficiency was catalyzed by complexes containing P(*p*-OCH₃Ph)₂CH₂OH (IrMPOH and RuMPOH). The catalytic property might have important future applications in biological and medical fields like production of reactive oxygen species (ROS). Furthermore redox activity of complexes was confirmed by cyclic voltamperometry. Biochemical assays demonstrated the ability of Ir(III) and Ru(II) complexes to induce significant cytotoxicity in various cancerous cell lines. Furthermore we found that RuPOH and RuMPOH selectively inhibit proliferation of skin cancer cells (WM266-4; IC₅₀, after 24h: *av.*48.3 μ M; after 72h: *av.*10.2 μ M) while Ir(III) complexes were found to be moderately active against prostate cancer cells (DU145).

Graphical Abstract

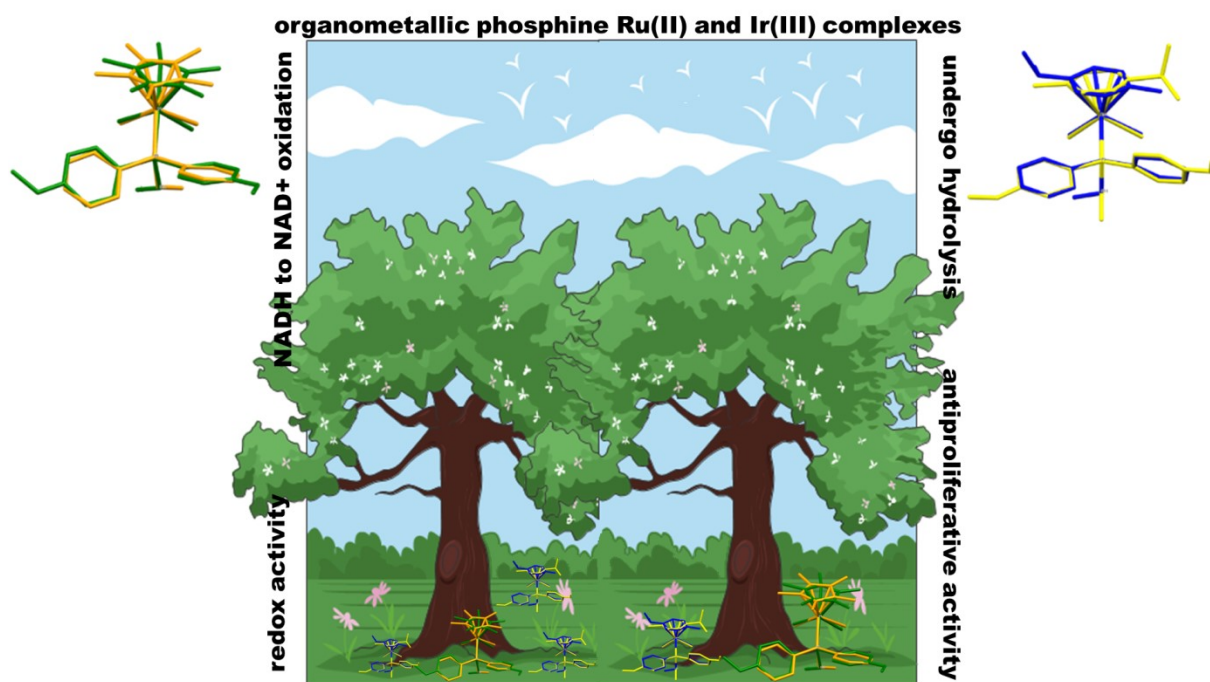


Table S1. Crystallographic experimental details.

Identification code	RuPOH	IrPOH	RuMPOH	IrMPOH
Crystal data				
Empirical formula	C ₂₃ H ₂₇ Cl ₂ OPRu	C ₂₃ H ₂₈ Cl ₂ IrOP	C ₂₅ H ₃₁ Cl ₂ O ₃ PRu	C ₂₅ H ₃₂ Cl ₂ IrO ₃ P
Formula weight	522.38	614.52	582.44	674.57
Temperature [K]	100(2)	293(2)	293(2)	100(2)
Wavelength [Å]	1.54184	0.71073	0.71073	1.54184
Crystal system, space group	Monoclinic, P 2 ₁	Monoclinic, P 2 ₁ /n	Monoclinic, P 2 ₁ /n	Monoclinic, P 2 ₁ /n
a, b, c [Å]	7.6065(1), 13.8332(1), 10.7154(1)	10.5605(1), 17.1832(2), 12.7618(1)	9.6600(1), 11.1867(1), 23.5077(3)	8.5562(2), 20.6545(6), 13.9418(2)
β [°]	105.490(1)	93.613(1)	92.598(1)	92.251(2)
V [Å ³]	1086.54(2)	2311.19(4)	2537.71(5)	2461.95(10)
Z	2	4	4	4
Radiation type	Cu K _α	Mo K _α	Mo K _α	Cu K _α
μ [mm ⁻¹]	8.881	6.089	0.916	13.313
Crystal size [mm]	0.40 x 0.25 x 0.10	0.20 x 0.20 x 0.20	0.50 x 0.30 x 0.20	0.10 x 0.05 x 0.01
Data collection				
Diffractometer	Rigaku (Cu) XtaLAB Synergy-DW VHF with a HyPix-Arc 150 detector	Rigaku OD SuperNova Dual source with an Atlas detector	Rigaku OD SuperNova Dual source with an Atlas detector	Rigaku (Cu) XtaLAB Synergy-DW VHF with a HyPix-Arc 150 detector
Absorption correction	Multi-scan (<i>CrysAlisPRO</i> ; Rigaku OD, 2020)	Multi-scan (<i>CrysAlisPRO</i> ; Rigaku OD, 2015)	Multi-scan (<i>CrysAlisPRO</i> ; Rigaku OD, 2015)	Multi-scan (<i>CrysAlisPRO</i> ; Rigaku OD, 2020)
No. of measured, independent and observed [I > 2σ(I)] reflections	9002/8981/3776	35098/6202/5185	37574/6823/5422	27876/5031/4572
R(int)	0.0360	0.0397	0.0388	0.0482
Completeness	99.7 %	99.6 %	99.8 %	99.6 %
Refinement				
Data / restraints / parameters	9002 / 1 / 261	6201 / 0 / 270	6823 / 0 / 297	5031 / 0 / 297
R[F ² > 2σ(F ²)], wR(F ²), S	0.0191, 0.0497, 1.201	0.0220, 0.0411, 1.075	0.0305, 0.0626, 1.069	0.0394, 0.1008, 1.095
Δρ _{max} , Δρ _{min} (e Å ⁻³)	0.376 and -0.427	0.850 and -0.614	0.350 and -0.470	1.426 and -2.097

Rigaku OD (2015). *CrysAlis PRO*. Rigaku Oxford Diffraction Ltd, Yarnton, Oxfordshire, England.

Rigaku OD (2020). *CrysAlis PRO*. Rigaku Oxford Diffraction, Yarnton, England.

Sheldrick, G. M. (2015a). *Acta Cryst.* **A71**, 3–8

Table S2. Hydrogen-bond geometry (Å, °)

D-H...A	d(D-H)	d(H...A)	d(D...A)	<(DHA)
RuPOH				
C11-H11A...Cl2	0.99	2.89	3.487(4)	120.0
O11-H11...Cl1	0.80(6)	2.30(6)	3.059(3)	159(5)
IrPOH				
C26-H26...Cl1	0.93	2.87	3.640(3)	140.5
C32-H32...Cl2	0.93	2.96	3.606(3)	127.6
O11B_a-H10B_a...Cl1	0.82	2.20	2.957(3)	153.0
O11A_b-H10A_b...Cl2	0.82	2.32	3.099(9)	159.7
RuMPOH				
C11-H11A...Cl1	0.97	2.76	3.336(2)	118.6
C11-H11B...Cl2	0.97	2.78	3.355(2)	118.4
C25-H25...O34_#1	0.93	2.47	3.313(3)	150.8
C27-H27C...Cl2_#2	0.96	2.97	3.870(3)	157.5
C33-H33...Cl2_#3	0.93	2.83	3.426(2)	122.6
O11-H11O...Cl1_#2	0.77(3)	2.50(3)	3.259(2)	171(3)
Symmetry transformations used to generate equivalent atoms: #1 -x-1/2, y+1/2, -z+3/2 #2 -x, -y, -z+1 #3 -x+1/2, y+1/2, -z+3/2				
IrMPOH				
C23-H23...Cl1#1	0.95	2.72	3.444(6)	133.5
C32-H32...Cl1	0.95	2.84	3.667(6)	145.4
C33-H33...Cl2#2	0.95	2.99	3.590(6)	122.6
O11-H11...Cl2	0.84	2.39	3.186(5)	157.2
Symmetry transformations used to generate equivalent atoms: #1 x-1/2,-y+1/2,z+1/2 #2 x-1/2,-y+1/2,z-1/2				

Table S3. Cumulative NMR data (298 K, δ [ppm], J [Hz]) for ligands (in CDCl₃) and iridium-complexes (in CD₂Cl₂).

	POHC ⁴⁰	MPOHC ⁴²	POH ⁴¹	MPOH ⁴²	[Ir(η^5 -Cp [*])Cl ₂] ₂ ^a	[Ru(η^6 -p-cymene)Cl ₂] ₂	IrPOH	IrMPOH	RuPOH	RuMPOH
³¹ P	-11.46	15.23	-9.33	-13.60			-7.15	-9.18	16.74	15.07
H ^{Ph(m)}	7.52-7.21	7.25-7.82	7.60-7.31	7.25-7.82			7.41-7.54 m	7.70 t (9.34)	7.41-7.58 m	7.77 t (9.07)
H ^{Ph(o)}							7.72-7.84 m	6.98 dd (8.80; 1.50)	7.79-7.93 m	6.98 dd (8.66; 1.40)
H ¹	4.32 d (7.7)	5.03 s	4.42 d (8.39)	4.27 d (9.01)			4.97 d (6.6)	4.86 s	4.64 s	4.54 s
H ^{-OH}		not observed	1.80 m	not observed			not observed	not observed	not observed	not observed
H ^{-OCH3}		3.87 s		3.74 s				3.85 s		3.86 s
C ^{Ph(i)}	135.65 d (12.0)	106.89 d (84.47)	135.65 d (12.0)	126.28 d (9.08)				118.82 d (55.4)	132.49 d (15.2)	123.14 d (47.23)
C ^{Ph(o)}	132.91 d (17.6)	115.63 d (7.54)	132.91 d (17.6)	114.41 d (7.27)			133.80 d (9.9)	113.93 d (10.9)	133.61 d (9.1)	114.15 s
C ^{Ph(m)}	128.39 d (5.6)	135.61 d (9.99)	128.39 d (5.6)	134.58 d (19.07)			128.15 d (9.9)	135.57 d (10.9)	128.55 d (9.1)	135.02 s
C ^{Ph(p)}	128.64 s	164.03 s	128.64 s	160.42 s			130.92 s	161.83 s	131.17 s	161.68 s
C ¹	62.46 d (13.0)	53.31 d (59.04)	62.46 d (13.0)	62.94 d (14.53)			64.25 d (39.1)	64.87 d (31.06)	63.28 d (40.1)	63.51 d (31.79)
C ^{-OCH3}		55.86 s		55.23 s				55.35 s		55.39 s
H ^{Cp*(CH3)}					0.96 s		1.40 d (2.19)	1.40 d (2.19)		
H ^{2,3}						1.29 d (6.9)			0.93 d (7.0)	0.97 d (6.98)
H ⁴						2.93 spt (7.0)			2.56 spt (7.0)	2.61 spt (7.00)
H ^{6,7,8,9}						5.42 dd (19.2; 6.0)			5.27 dd (19.6; 6.2)	5.25 s
H ¹¹						2.16 s			1.91 s	1.89 s
C ^{Cp*(CH3)}					9.38 s		8.08 s	8.35 s		
C ^{Cp*}					86.28 s		92.28 d (2.2)	92.34 d (2.72)		
C ^{2,3}						22.12 s			21.73 s	21.60 s
C ⁴						30.63 s			30.34 s	30.18 s
C ⁵						101.22 s			108.84 s	108.76 s
C ^{6,8}						80.54 s			86.42 d (5.4)	86.19 s
C ^{7,9}						81.32 s			89.74 d (3.6)	89.31 s
C ¹⁰						96.75 s			95.60 s	95.14 s
C ¹¹						18.89 s			17.70 s	17.57 s

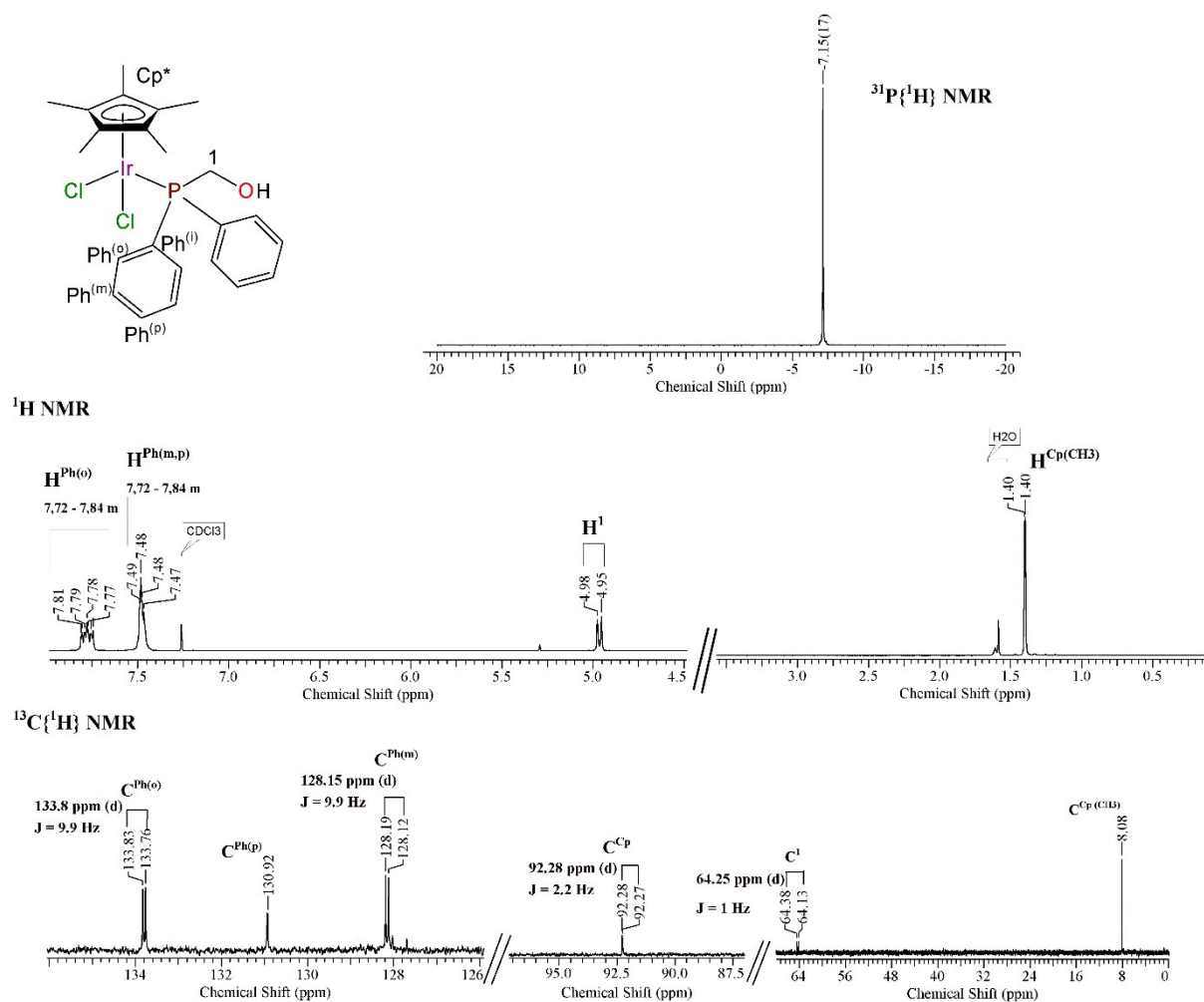


Figure S1. ^1H , $^{13}\text{C}\{^1\text{H}\}$ and $^{31}\text{P}\{^1\text{H}\}$ NMR spectra for IrPOH (298 K, CDCl_3).

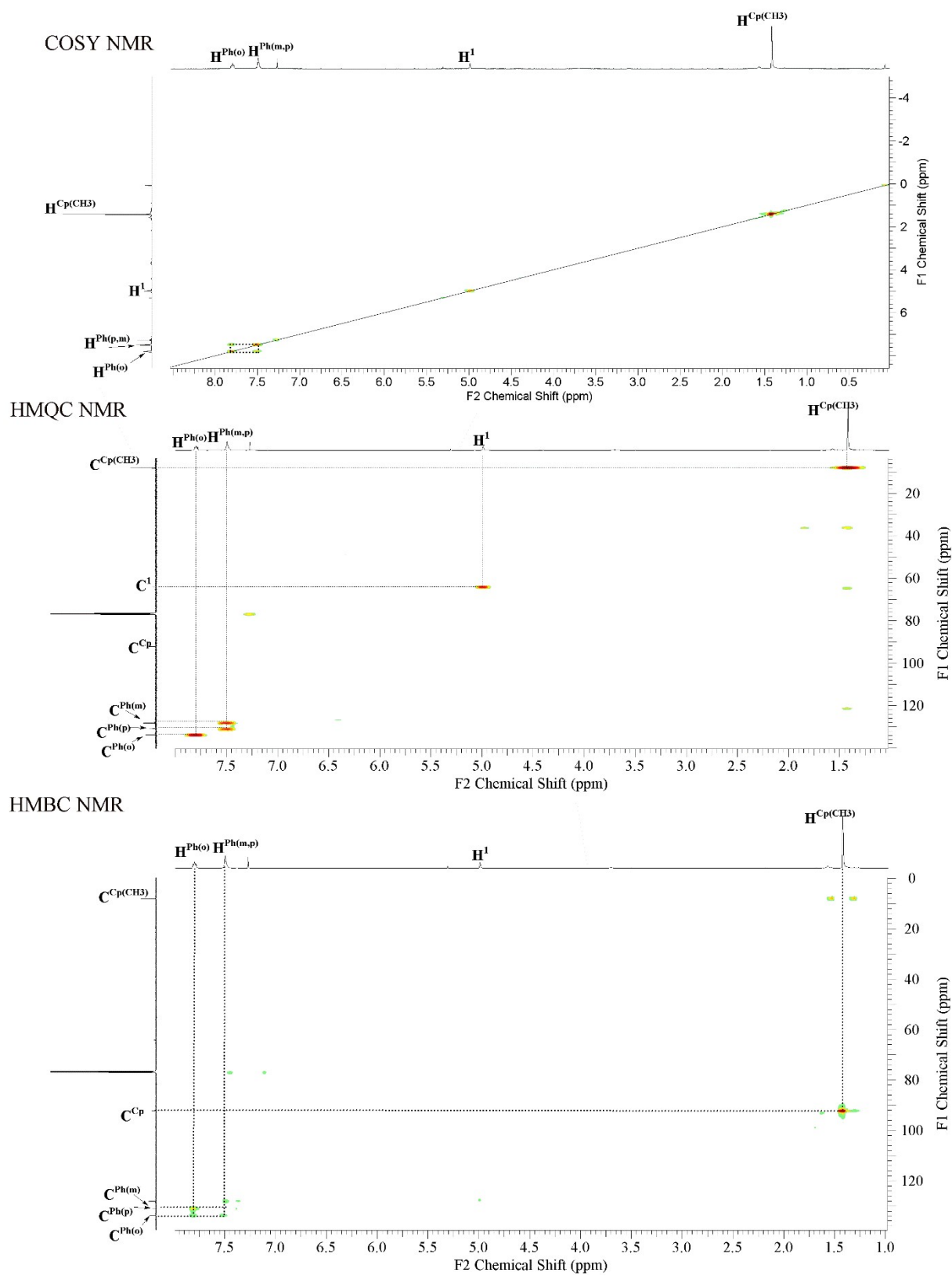


Figure S2. HMQC, HMBC, COSY NMR spectra for **IrPOH** (298 K, CDCl_3).

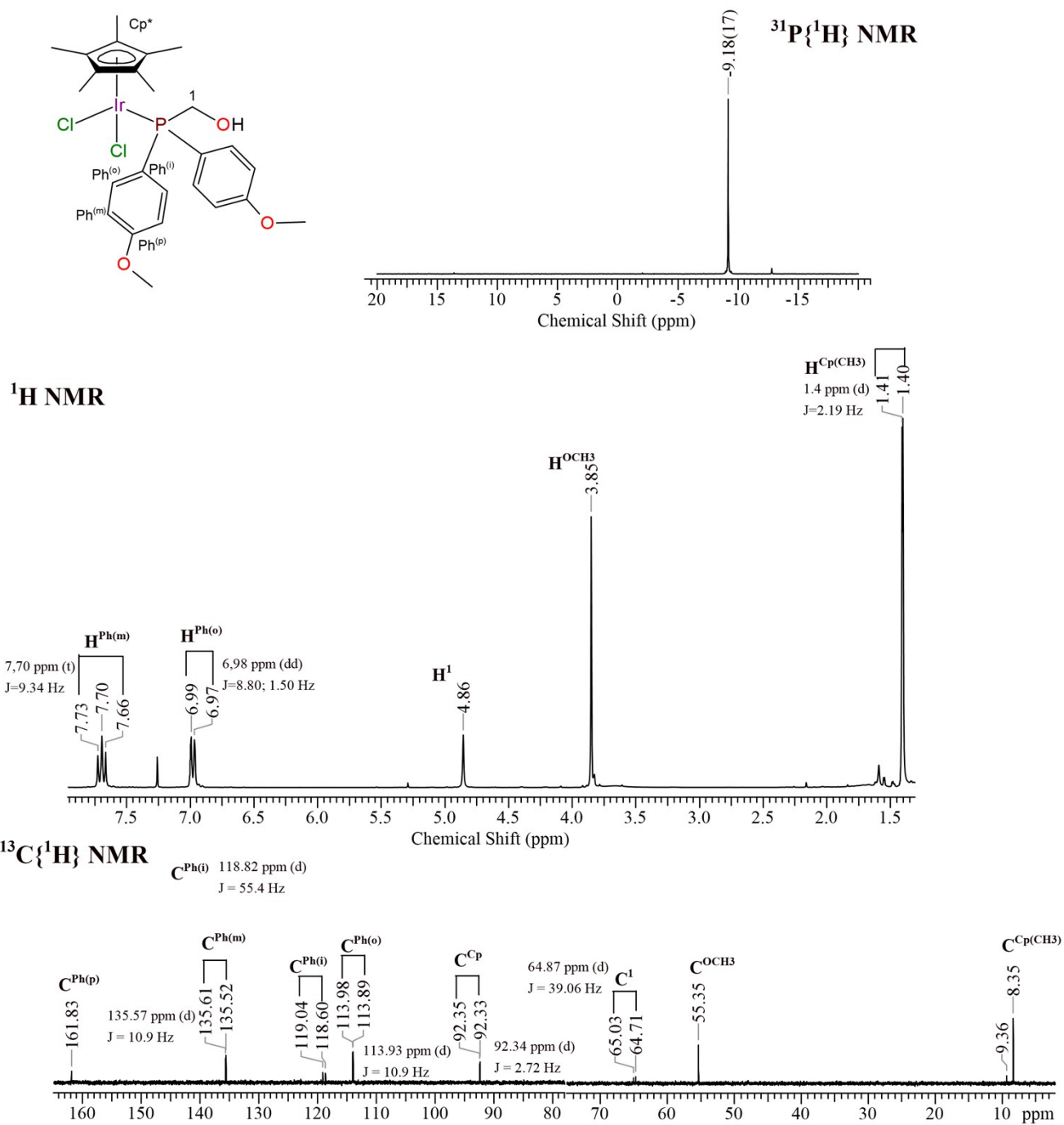


Figure S3. ^1H , $^{13}\text{C}\{^1\text{H}\}$ and $^{31}\text{P}\{^1\text{H}\}$ NMR spectra for IrMPOH (298 K, CDCl_3).

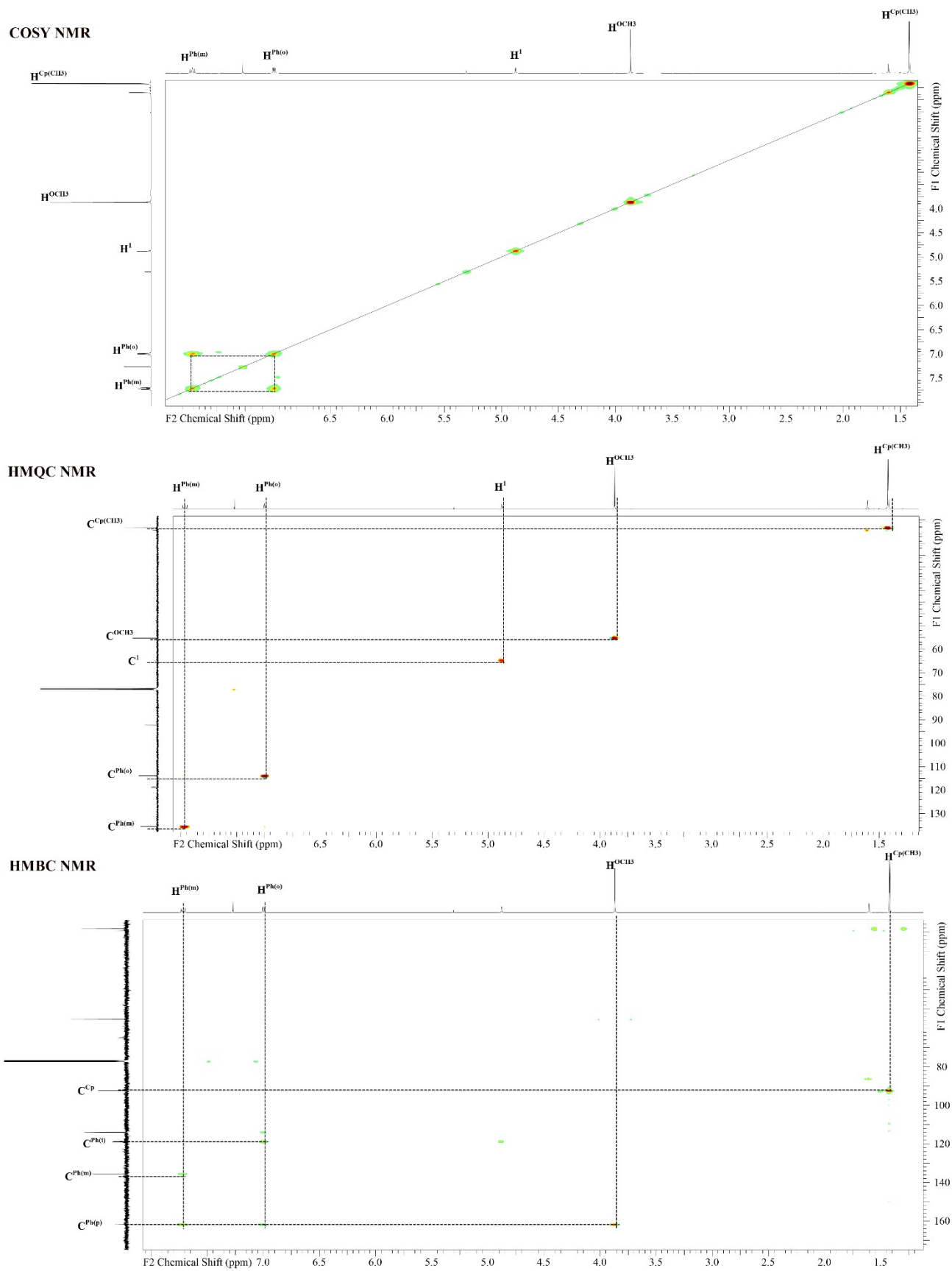
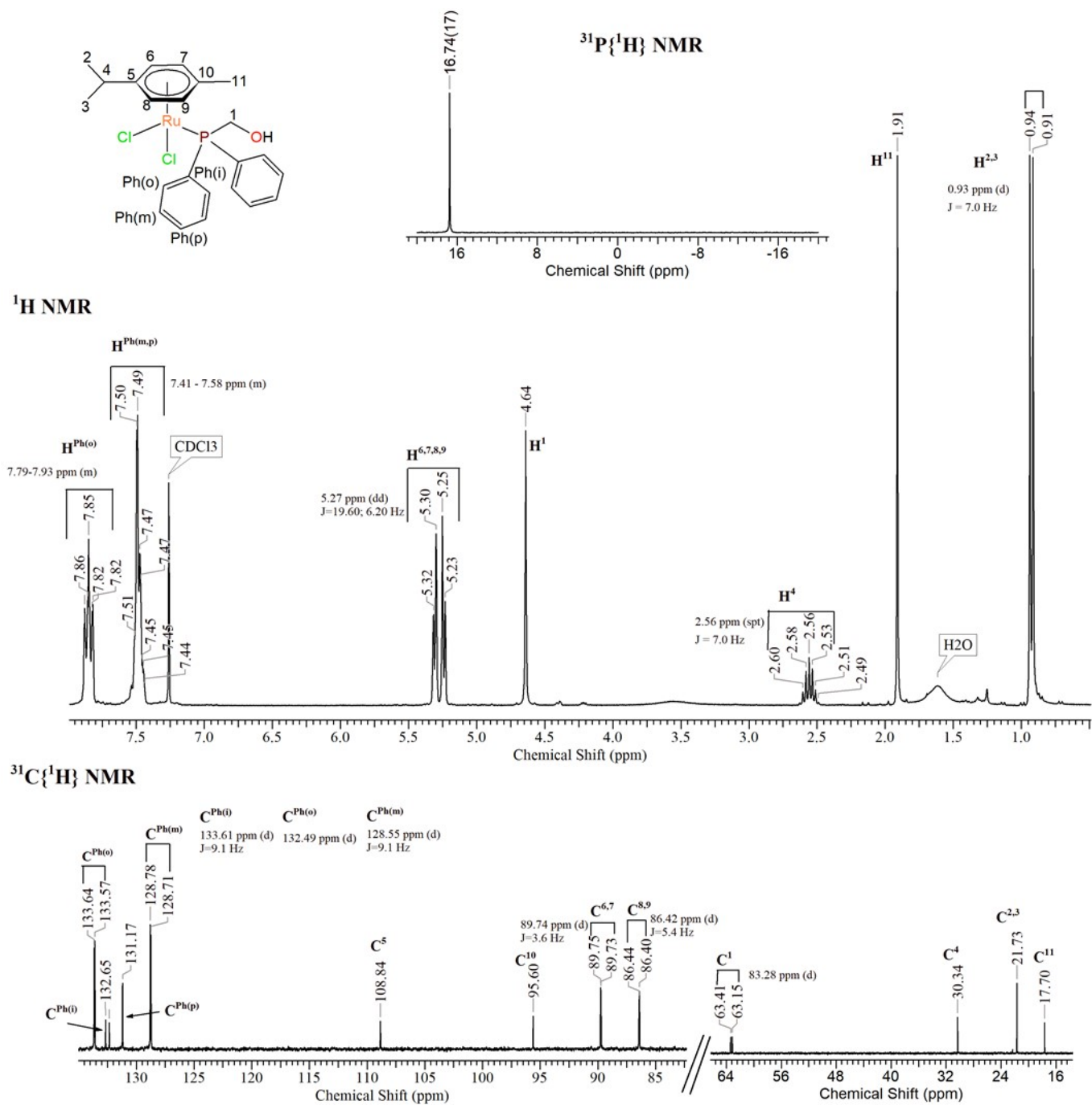


Figure S4. HMQC, HMBC, COSY NMR spectra for **IrMPOH** (298 K, CDCl_3).



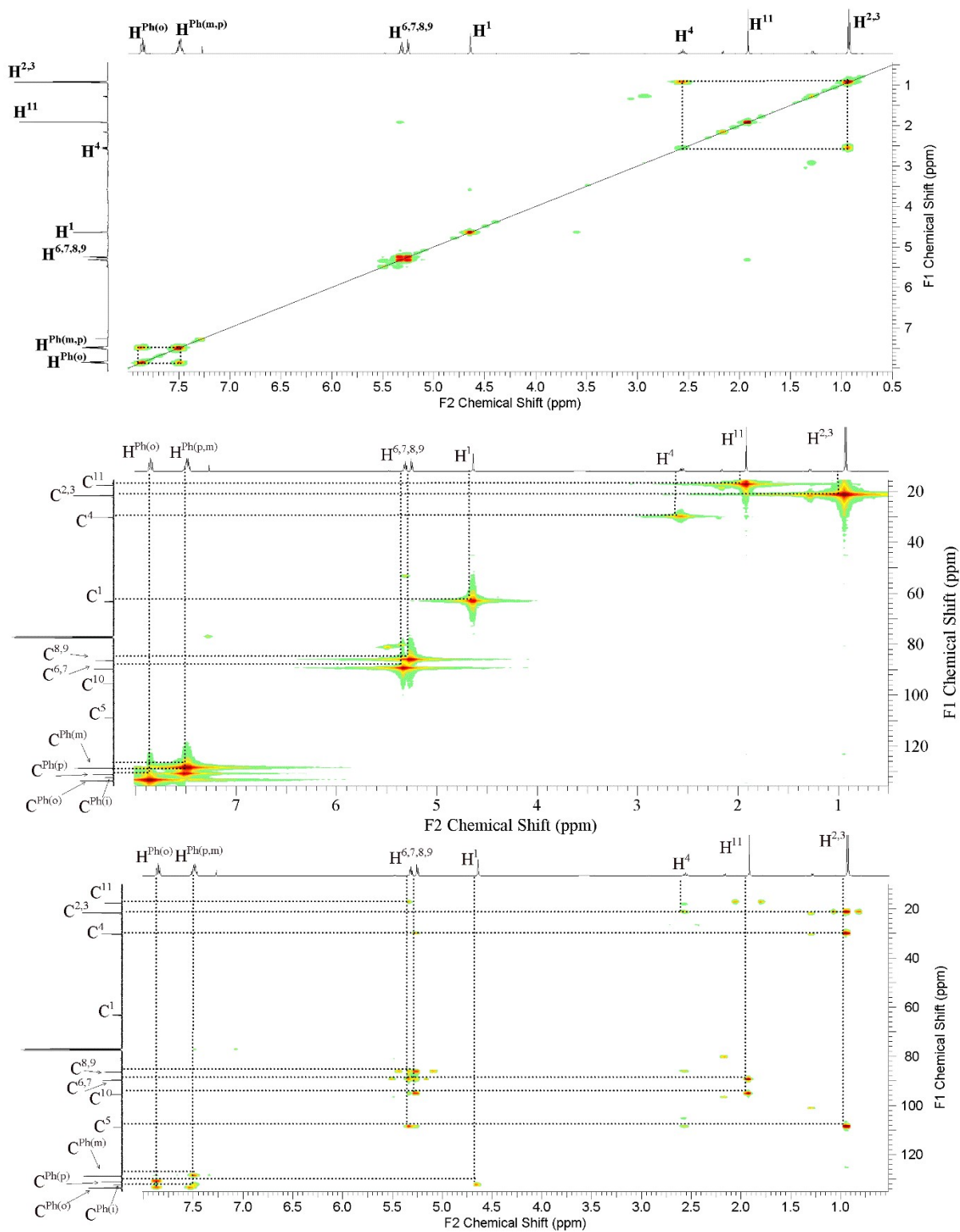


Figure S6. HMQC, HMBC, COSY NMR spectra for **RuPOH** (298 K, CDCl₃).

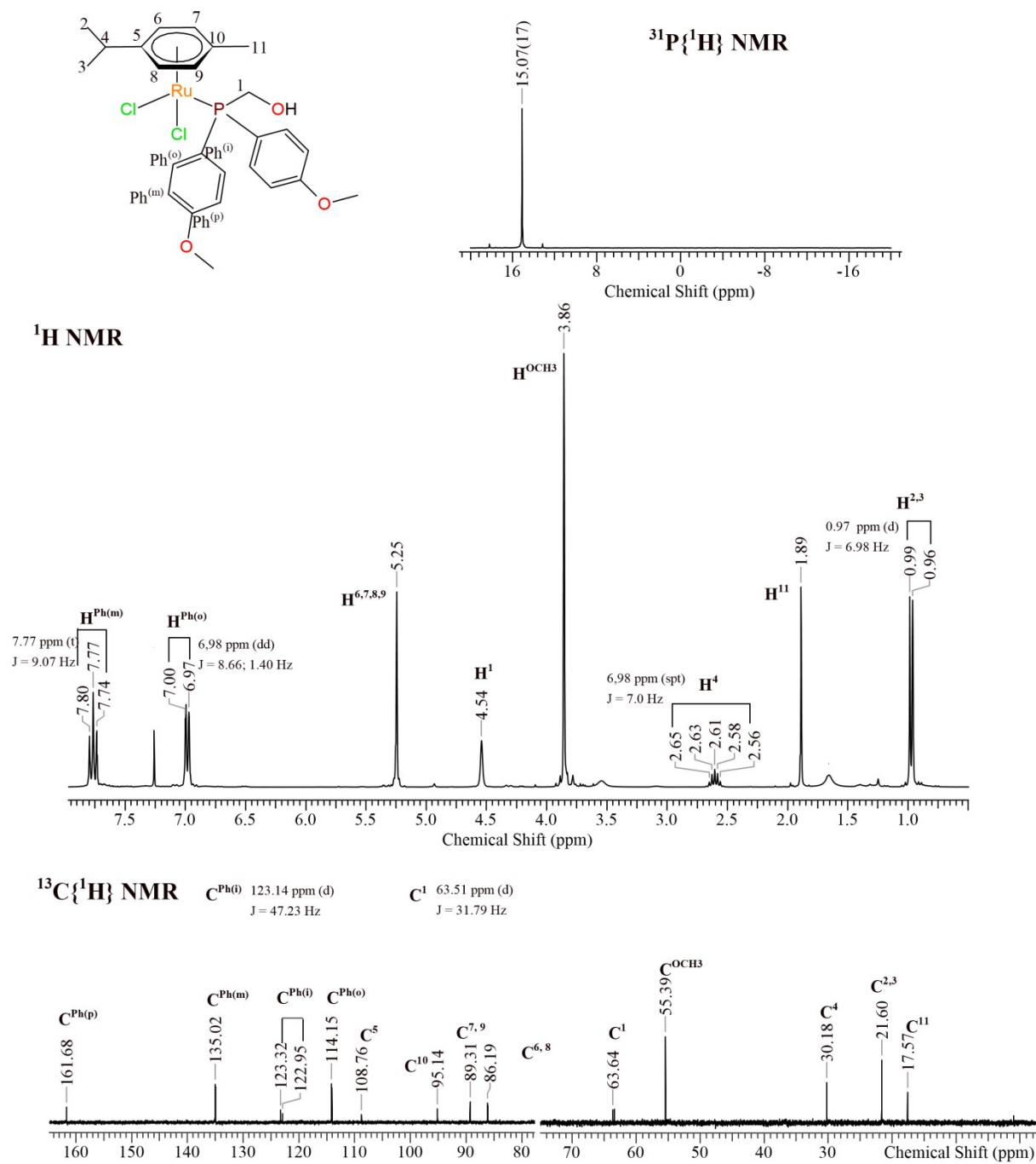


Figure S7. ^1H , $^{13}\text{C}\{^1\text{H}\}$ and $^{31}\text{P}\{^1\text{H}\}$ NMR spectra for **RuMPOH** (298 K, CDCl_3).

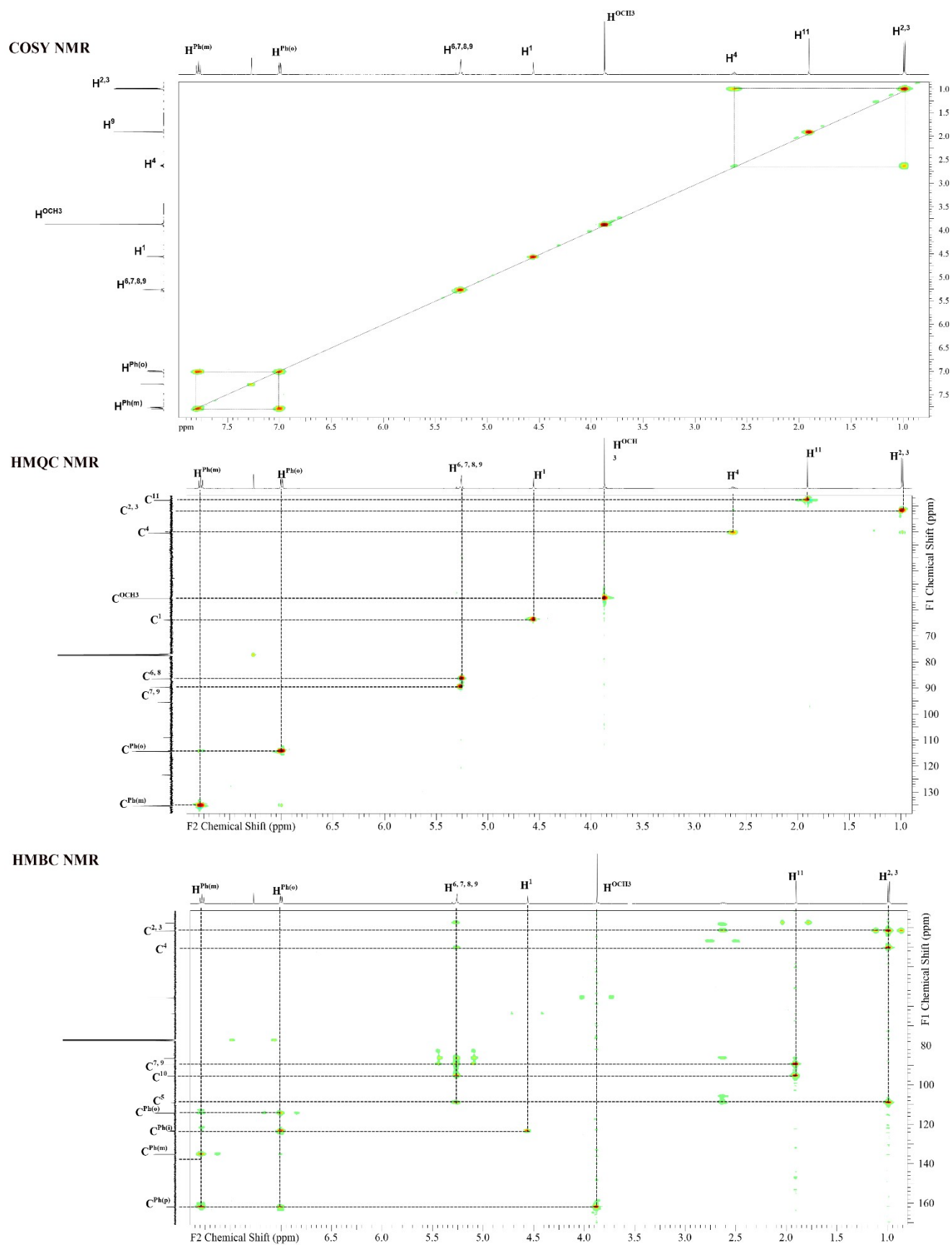


Figure S8. HMQC, HMBC, COSY NMR spectra for **RuMPOH** (298 K, CDCl_3).

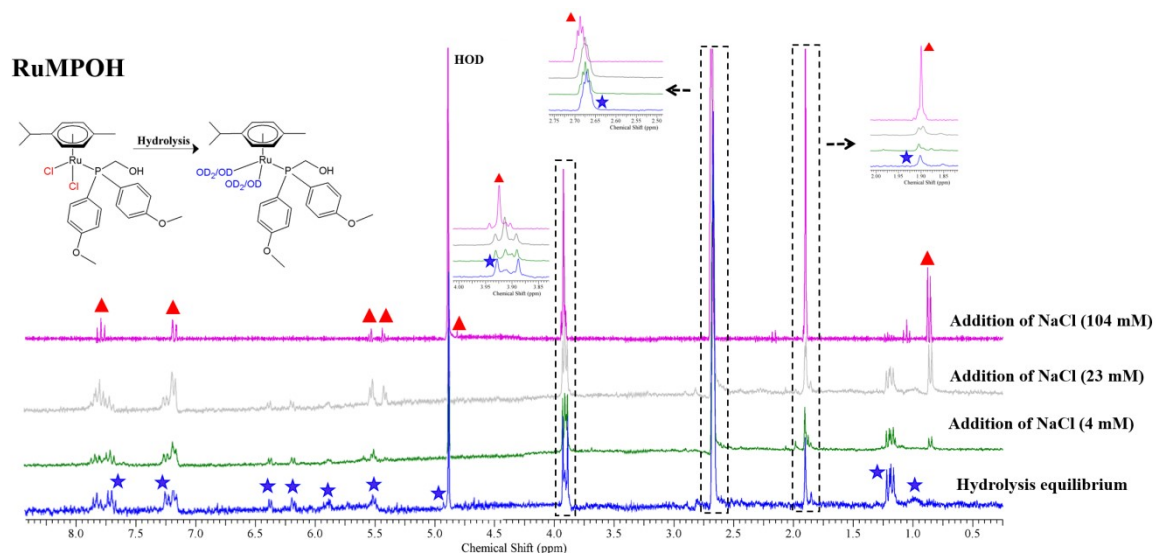


Figure S9. Confirmation of hydrolysis of **RuMPOH** by ^1H NMR in 20% $\text{DMSO-d}_6/80\%$ D_2O (v/v) at 298 K. From bottom to top: ^1H NMR spectrum of an equilibrium solution of **RuMPOH** (1 mM); spectrum recorded 10 min after addition of NaCl (final concentration, 4 mM) to the equilibrium solution of **RuMPOH**; final concentration of NaCl, 23mM; final concentration of NaCl, 104mM – respectively. The peaks for the chlorido complex **RuMPOH** (red triangle) increased in intensity while peaks for the aqua complex decreased (blue star).

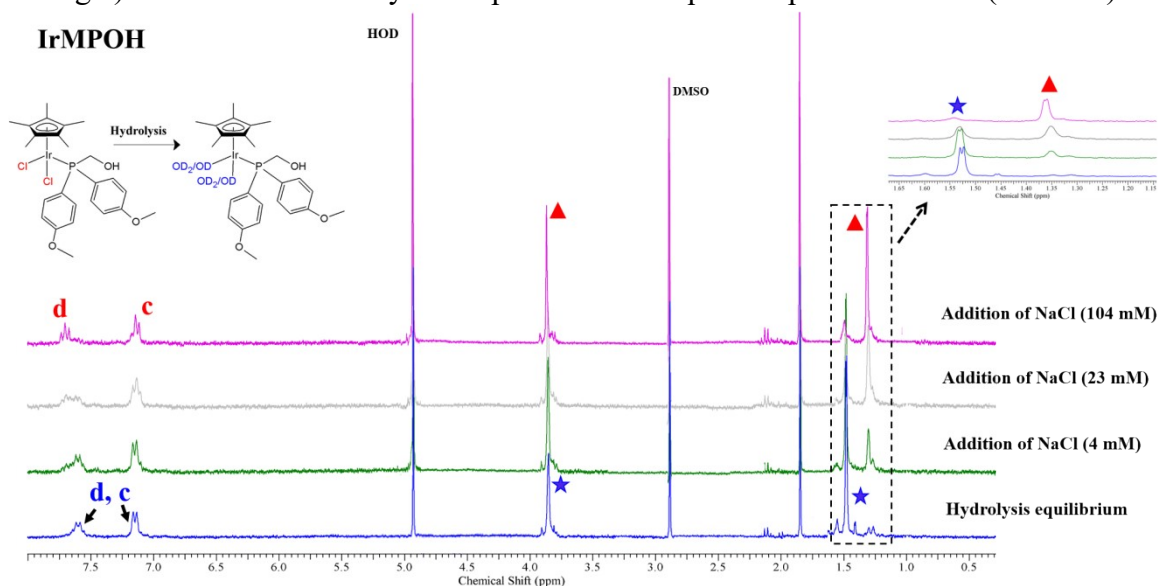


Figure S10. Confirmation of hydrolysis of **IrMPOH** by ^1H NMR in 20% $\text{DMSO-d}_6/80\%$ D_2O (v/v) at 298 K. From bottom to top: ^1H NMR spectrum of an equilibrium solution of **IrMPOH** (1 mM); spectrum recorded 10 min after addition of NaCl (final concentration, 4 mM) to the equilibrium solution of **IrMPOH**; final concentration of NaCl, 23mM; final concentration of NaCl, 104mM – respectively. The peaks for the chlorido complex **IrMPOH** (red triangle) increased in intensity while peaks for the aqua complex decreased (blue star).

Ir PK010 CHCl3 MeOH bis Z 637_181116110705 #1 RT: 0,00 AV: 1 NL: 4,15E4
T: ITMS + c ESI Full ms [155,00-1500,00]

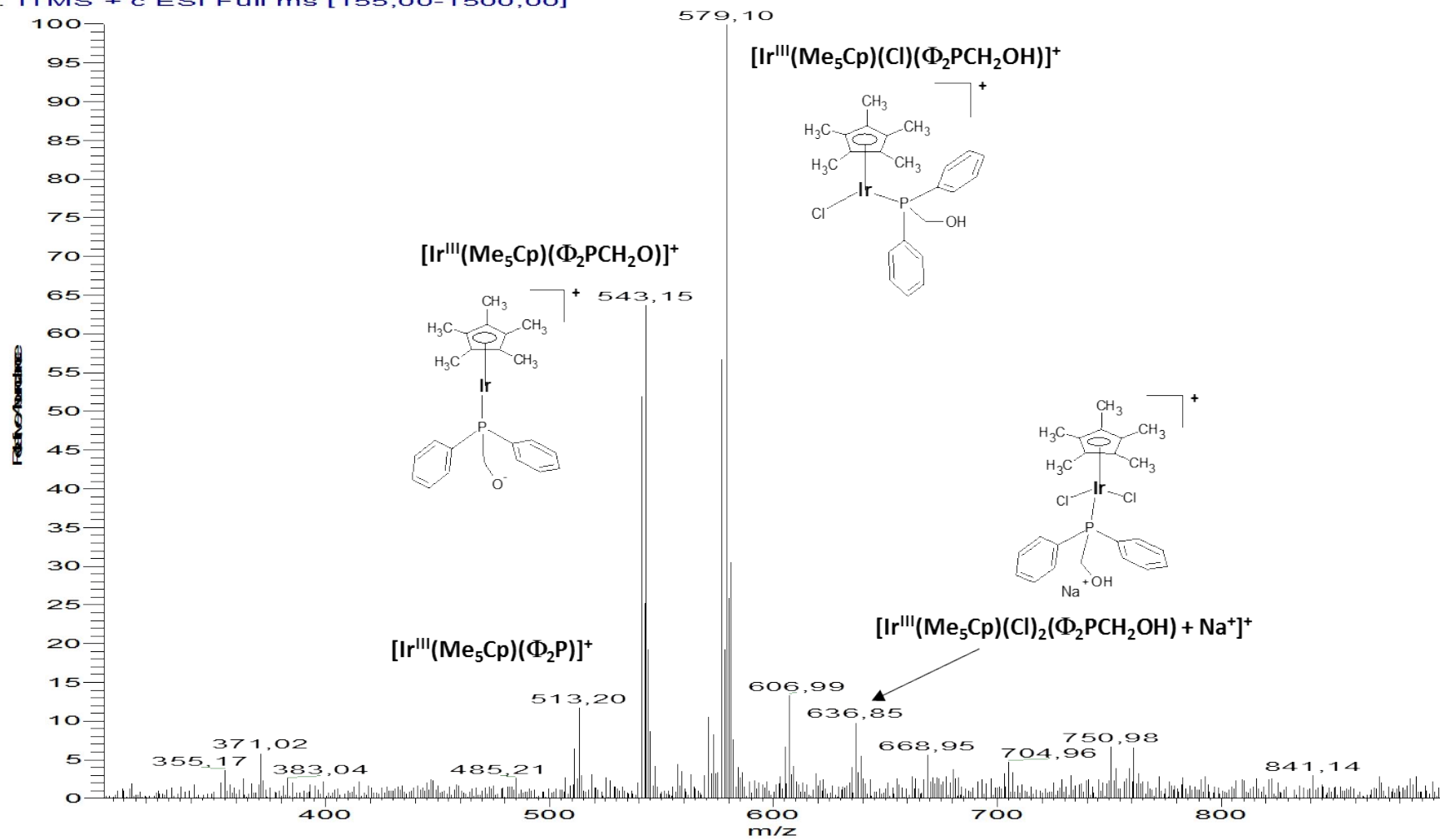


Figure S11. Full ESI(+)-MS spectrum of IrPOH

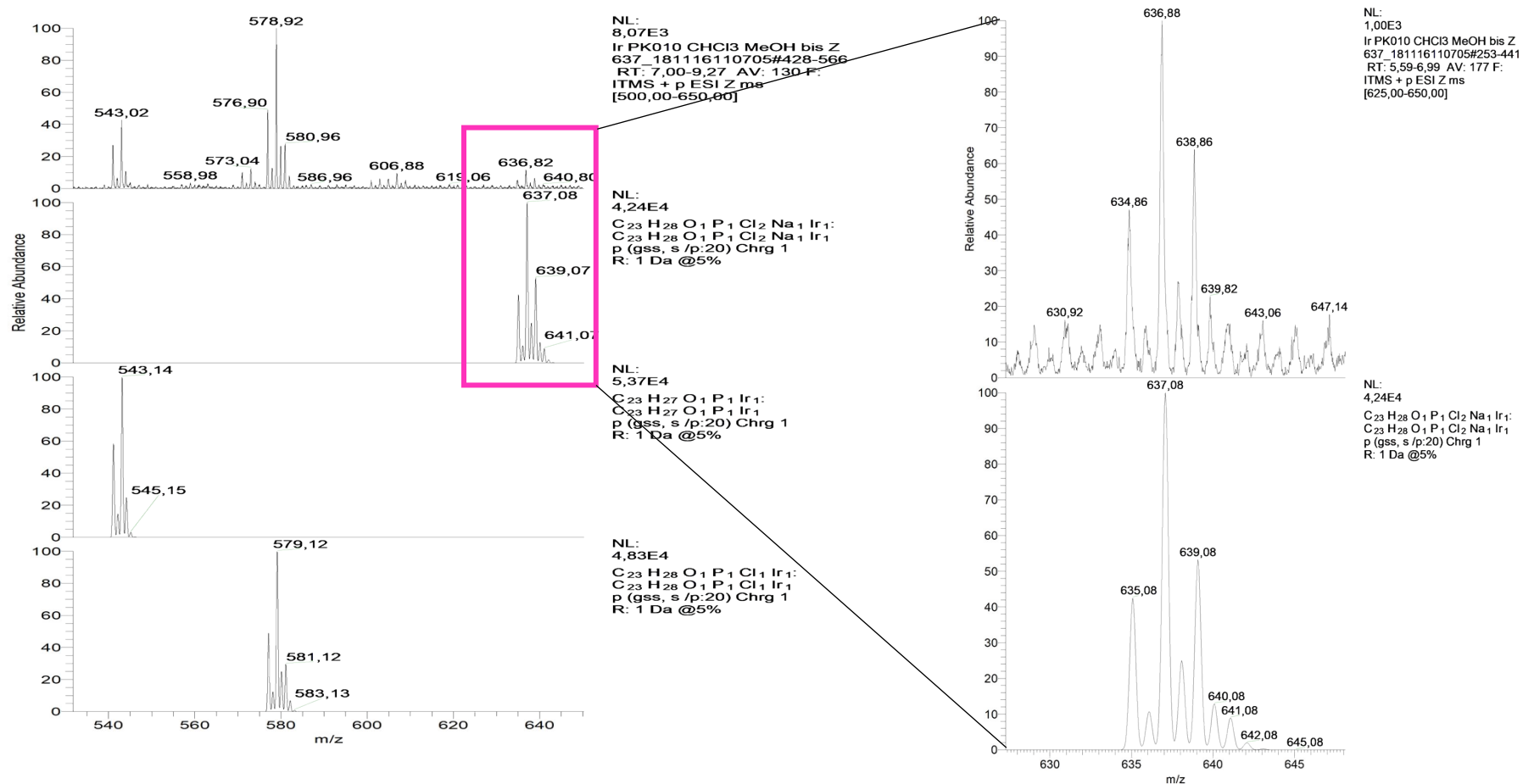


Figure S12. Comparison of ion peaks (experimental (top traces) vs calculated (bottom traces)) in the 630 – 650 m/z region of IrPOH. Note the different profile of the cluster centered at m/z 637 ([Ir(Me₅-Cp)(Cl)₂(Ph₂PCH₂OH) + Na]⁺, containing the IrCl₂ moiety) with respect to that centered at m/z 579 ([Ir(Me₅-Cp)(Cl)(Ph₂PCH₂OH)]⁺, containing the IrCl moiety), and to that centered at m/z 543 ([Ir(Me₅-Cp)(Ph₂PCH₂O)]⁺, containing the Ir ion only).

Ir PK012 CHCl3 MeOH bis Z 697_181116110705 #1 RT: 0,00 AV: 1 NL: 8,45E4
T: ITMS + c ESI Full ms [50,00-2000,00]

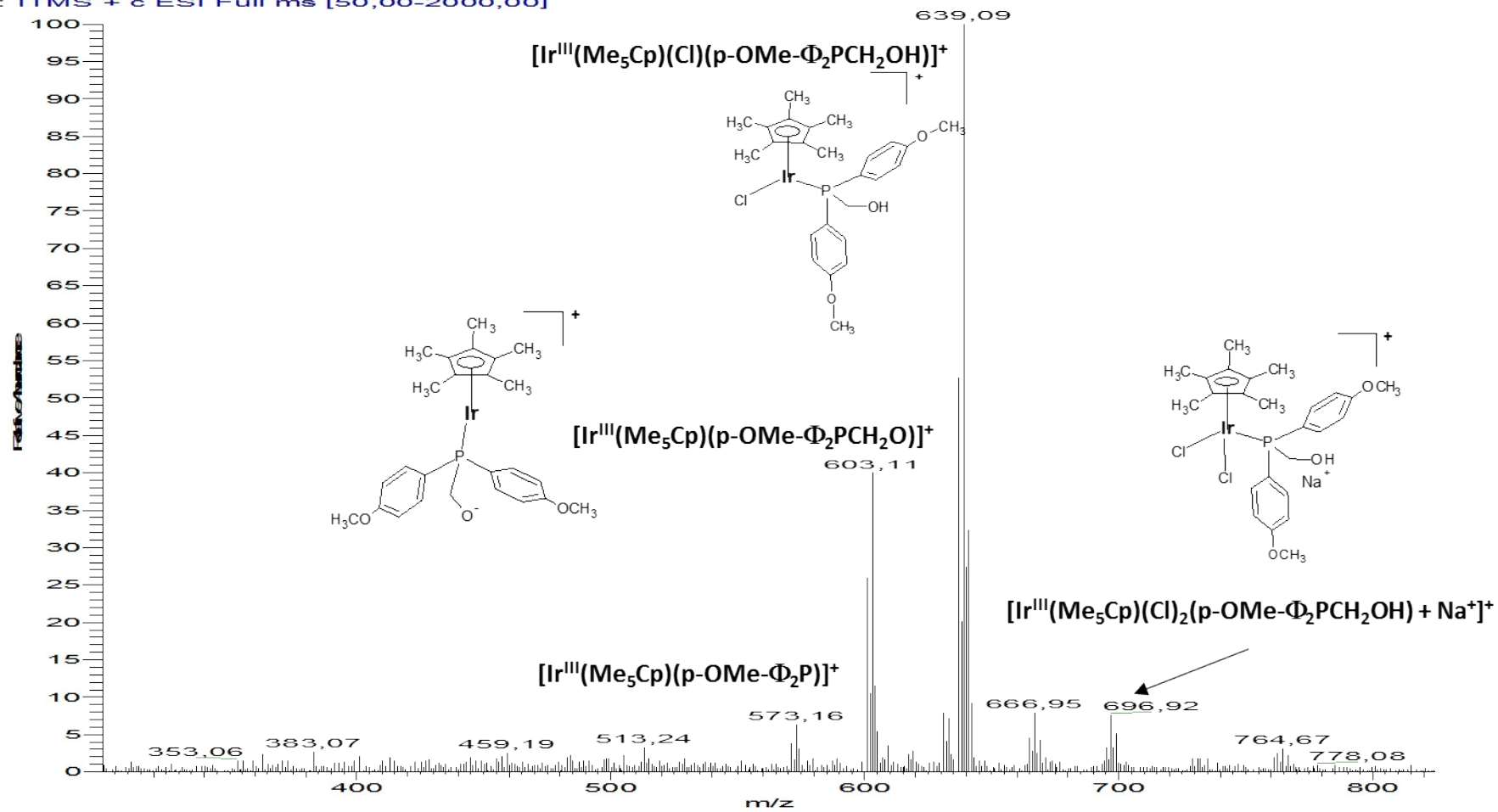


Figure S13. Full ESI(+)-MS spectrum of IrMPOH.

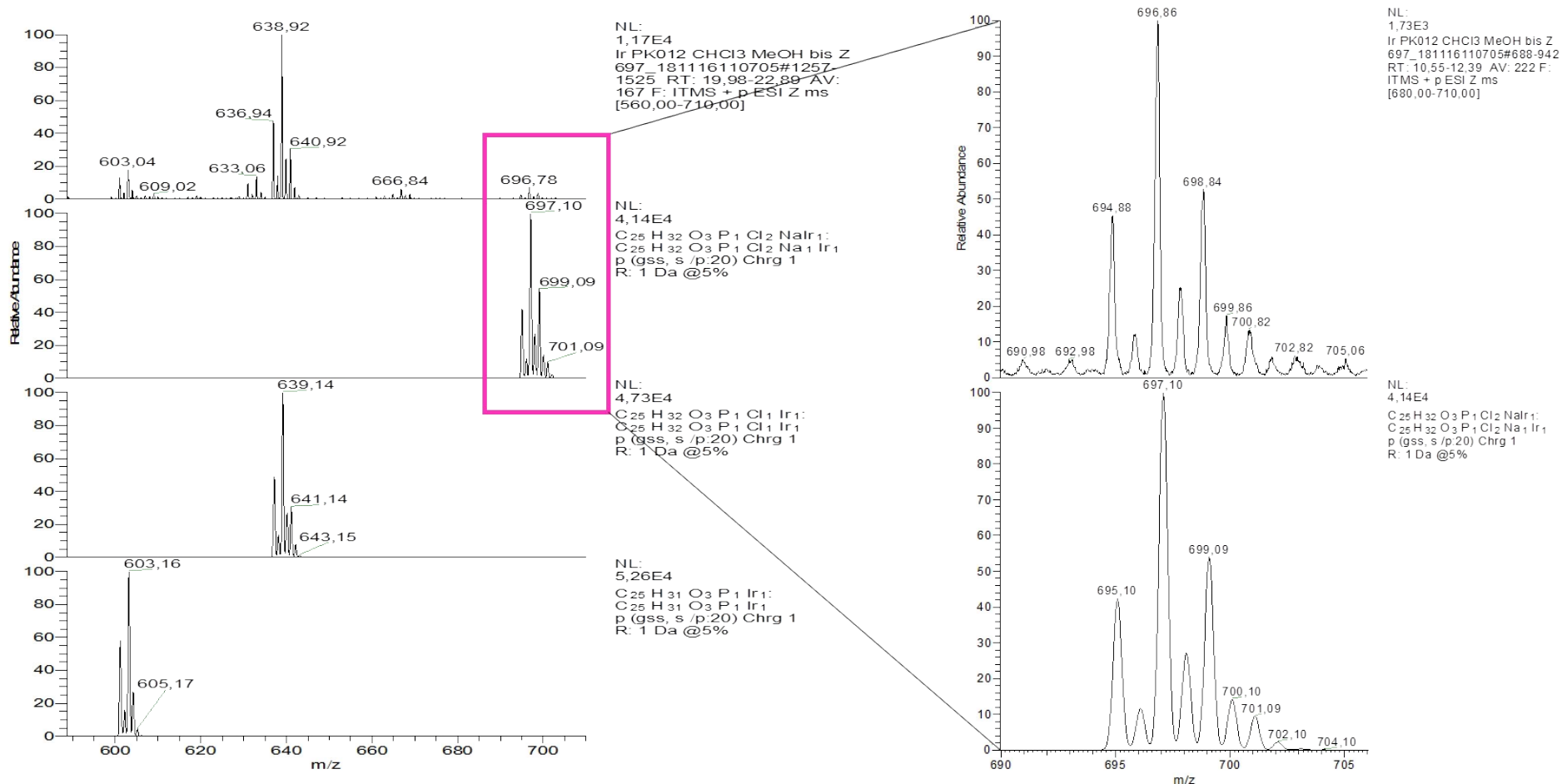


Figure S14 Comparison of high abundant cluster ion peaks (experimental vs calculated) in the 690 – 710 m/z region of **IrMPOH**. Note the different profile of the cluster centered at m/z 639 (containing the IrCl moiety) with respect to the other two profiles centered at m/z 603 and 697 (containing Ir only).

Ru PK009 CHCl3 MeOH bis Z 545_181116110705 #939 RT: 11,16 AV: 1 NL: 1,17E5
T: ITMS + c ESI Full ms [110,00-1000,00]

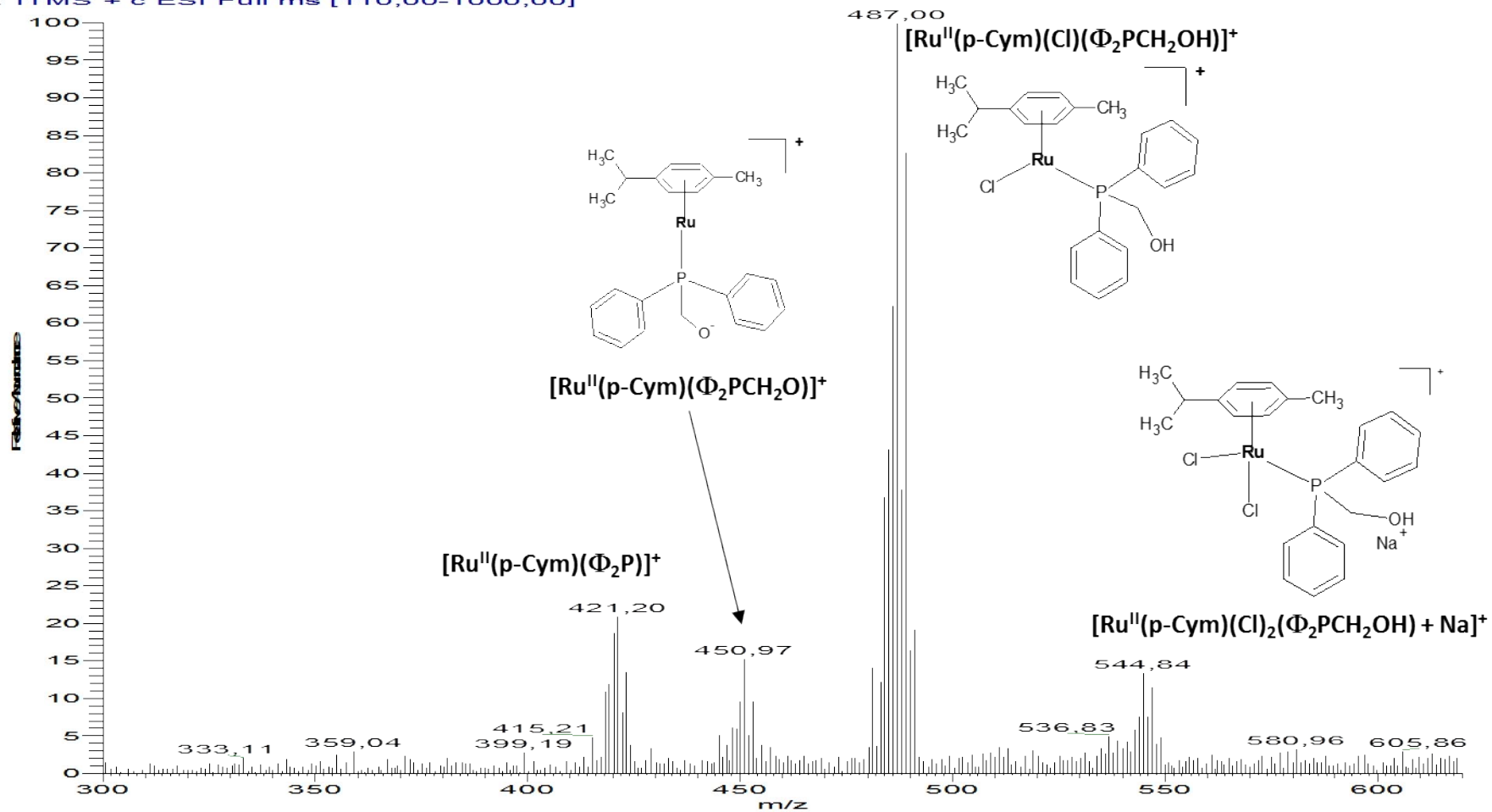


Figure S15. Full ESI(+)-MS spectrum of RuPOH.

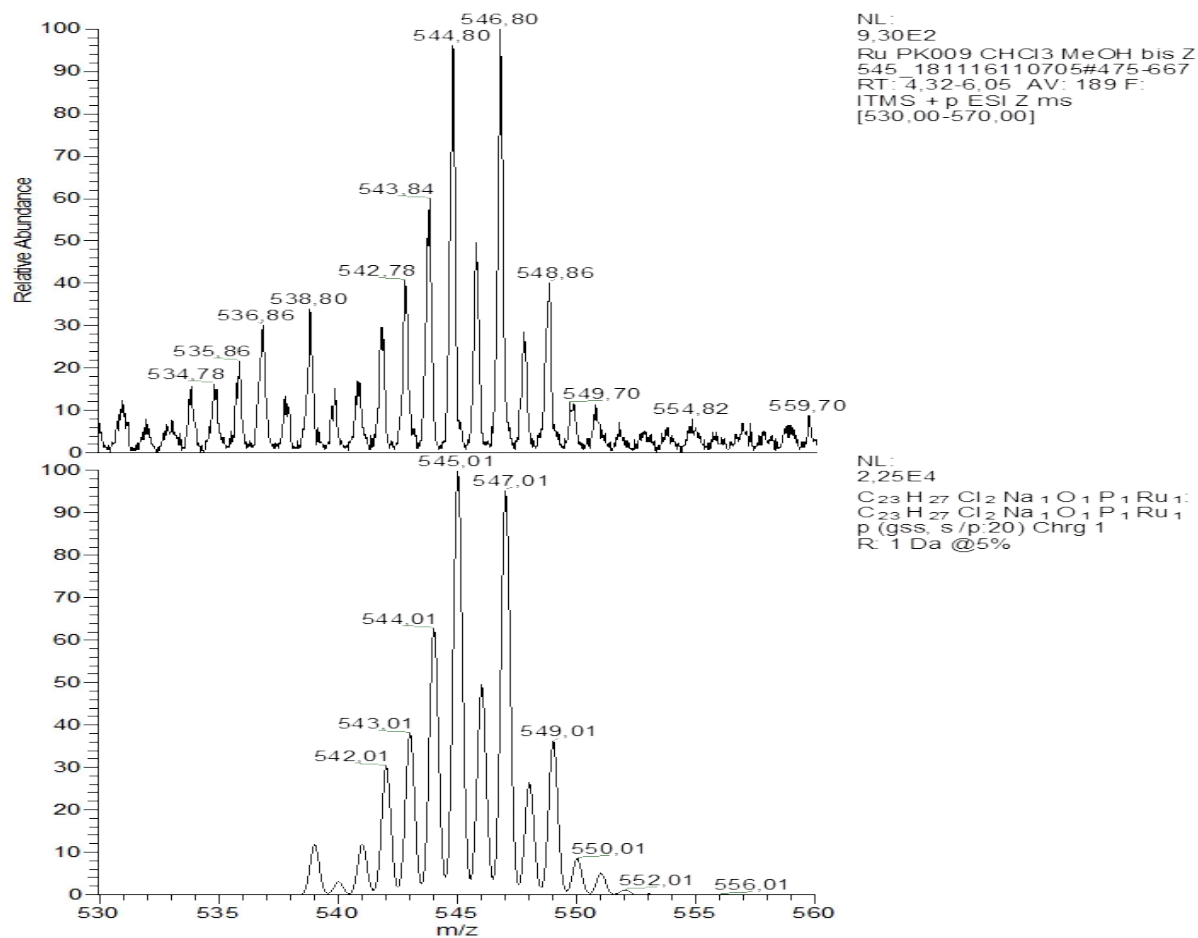


Figure S16. Comparison of the cluster ion peak of **RuPOH** centered at m/z 545 (experimental (top) vs calculated (bottom)) corresponding to the sodiated $[M + Na]^+$ ion.

Ru PK009 CHCl3 MeOH #1156-1582 RT: 7,02-10,35 AV: 73 NL: 1,36E3
F: ITMS - c ESI Full ms [50,00-1400,00]

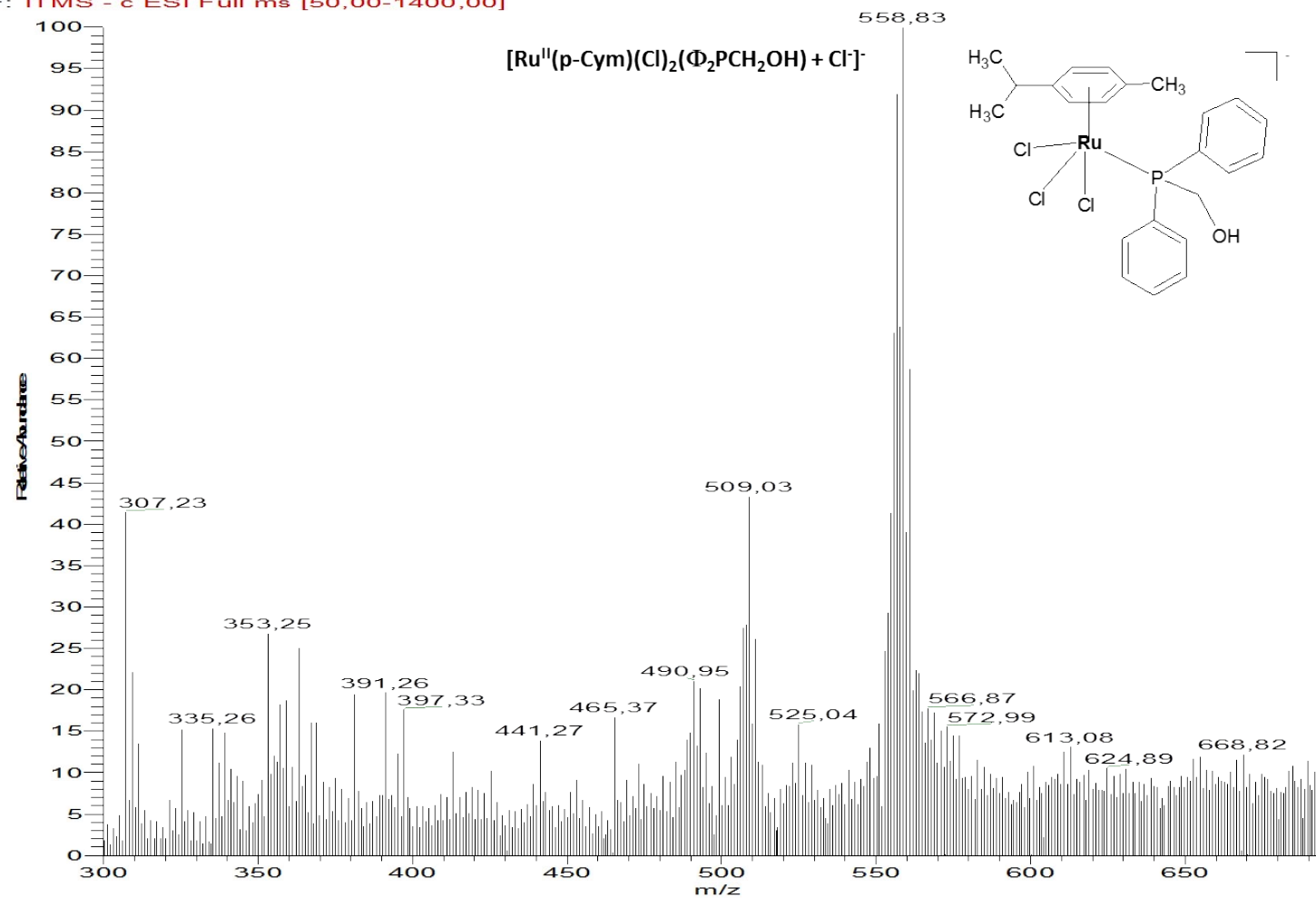


Figure S17. Full ESI(+)-MS spectrum of RuPOH.

Ru PK011 CHCl3 MeOH bis Z 605 181116110705 #610 RT: 6,92 AV: 1 NL: 3,86E4
T: ITMS + c ESI Full ms [130,00-1000,00]

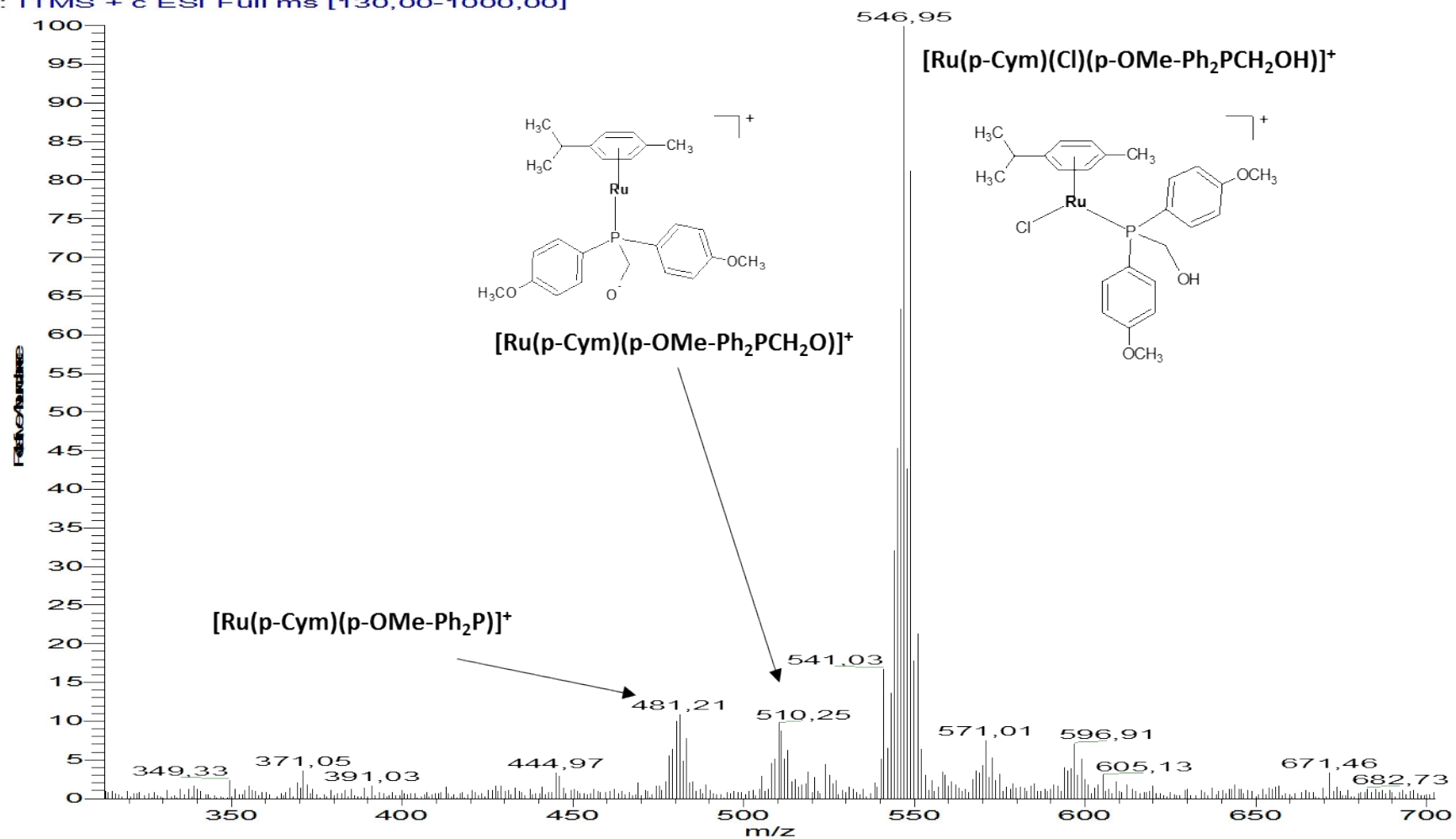


Figure S18. Full ESI(-)MS spectrum of RuMPOH.

Ru PK011 CHCl3 MeOH #498-534 RT: 8,82-9,27 AV: 29 NL: 8,27E2
F: ITMS - c ESI Full ms [70,00-1300,00]

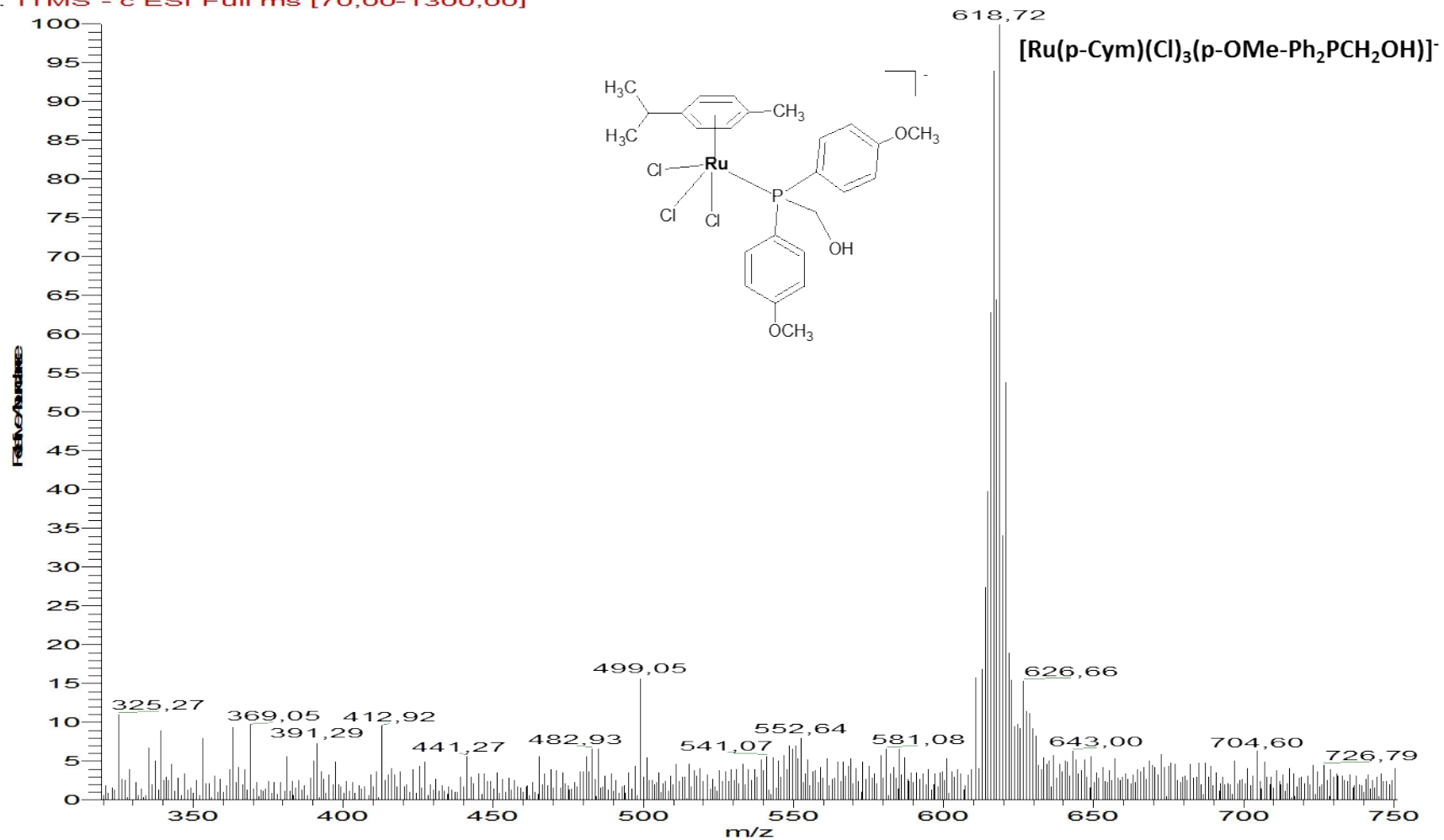


Figure S19. Full ESI(-)MS spectrum of RuMPOH.

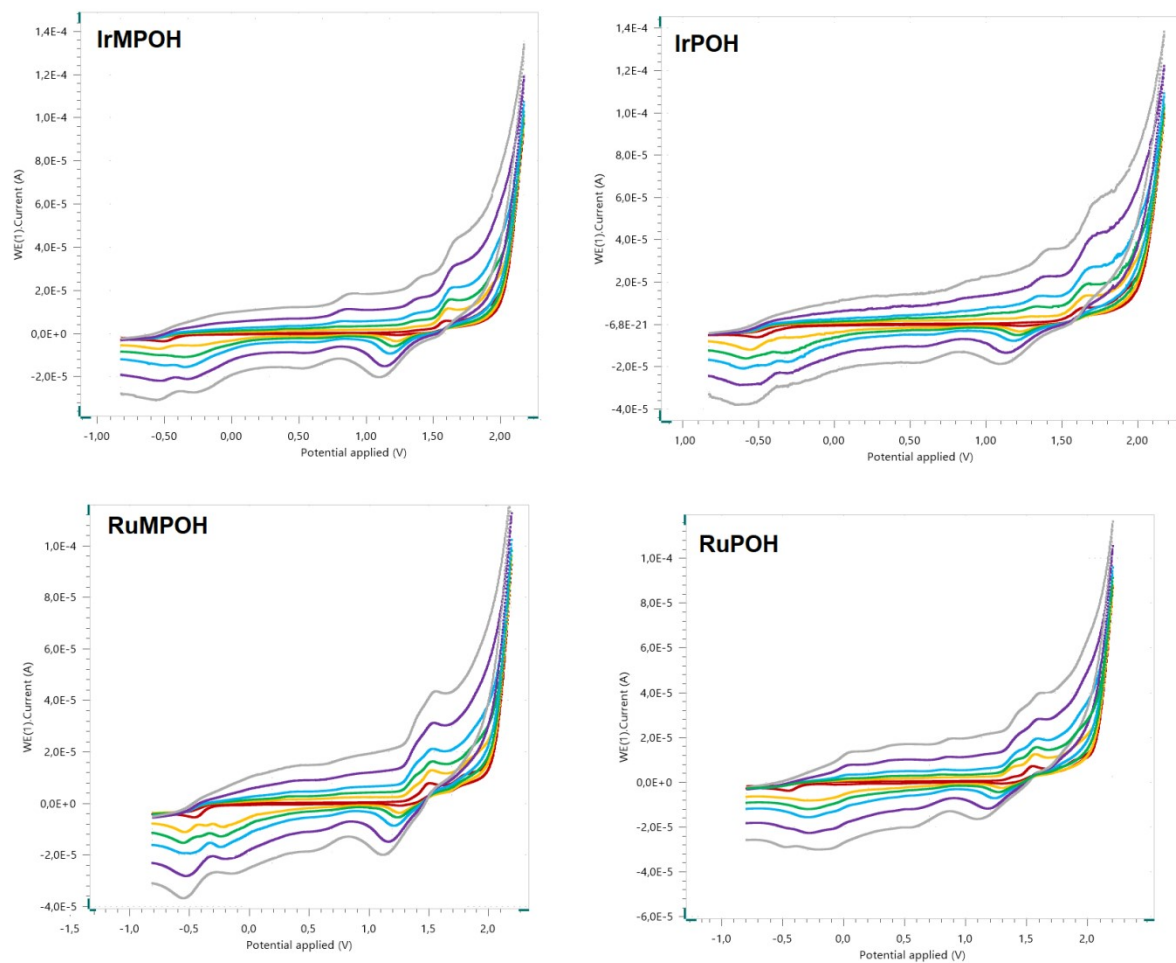


Figure S20. Cyclic voltammetric trace of **RuPOH**, **RuMPOH**, **IrPOH** and **IrMPOH** (1 mM) as a function of scan rate, recorded with recorded with 0.1 M tetrabutyl ammonium perchlorate (TBAP) as supporting electrolyte in DMF solution. Scan rates 1 – 100 (mV s^{-1}). Potential (V) versus $\text{Fc}^{0/+}$.

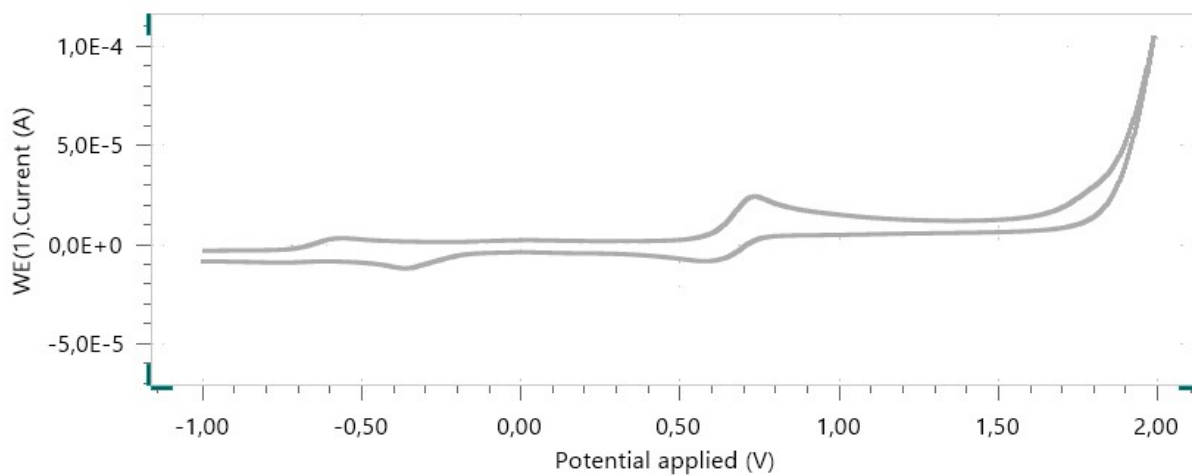


Figure S21. CV voltammograms for ferrocene in DMF in the range of potentials from -01 V to 1.2 V. Scan rate: 10 mV s⁻¹.

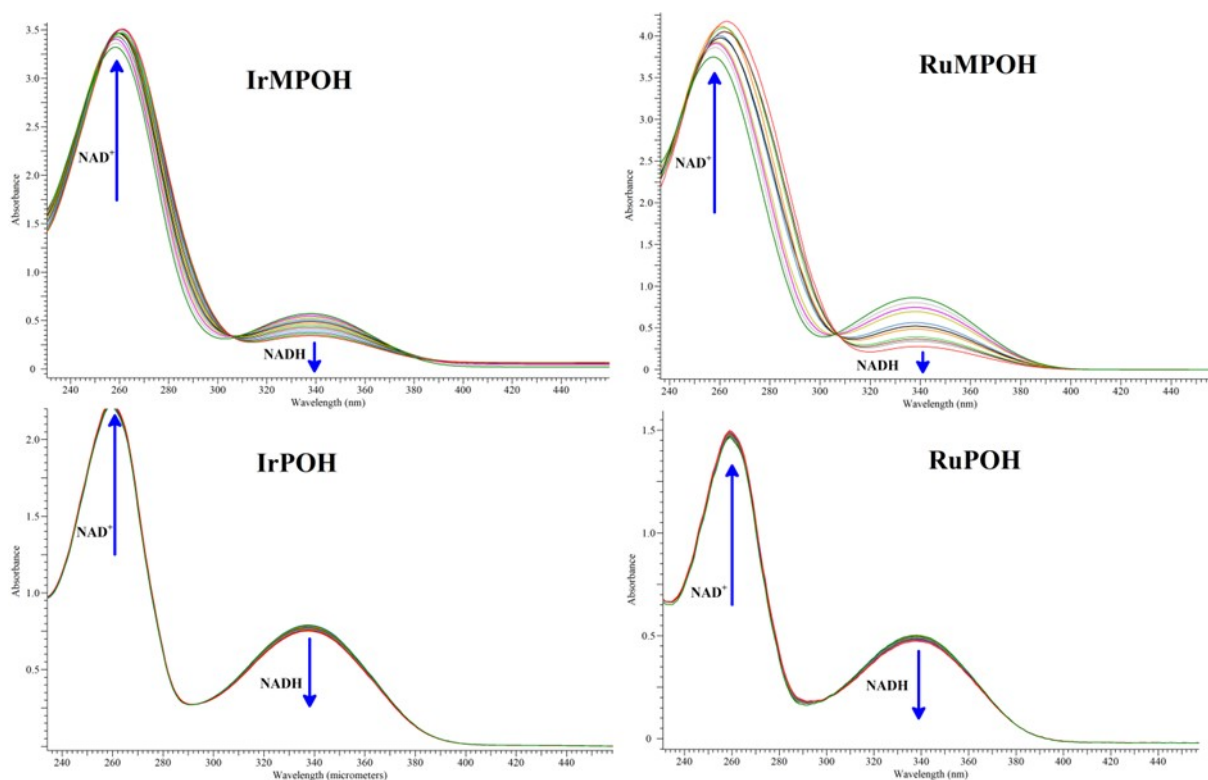


Figure S22. UV/Vis spectra of NADH (100 μ M) were determined after incubated with RuPOH, RuMPOH, IrPOH, IrMPOH (1 μ M) in CH₃OH/H₂O (1 : 9, v/v) at 298 K for 8 h. The arrows show the absorbance change over time.