

Supplementary Materials

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NMR spectra of precursor platinum(II) complexes

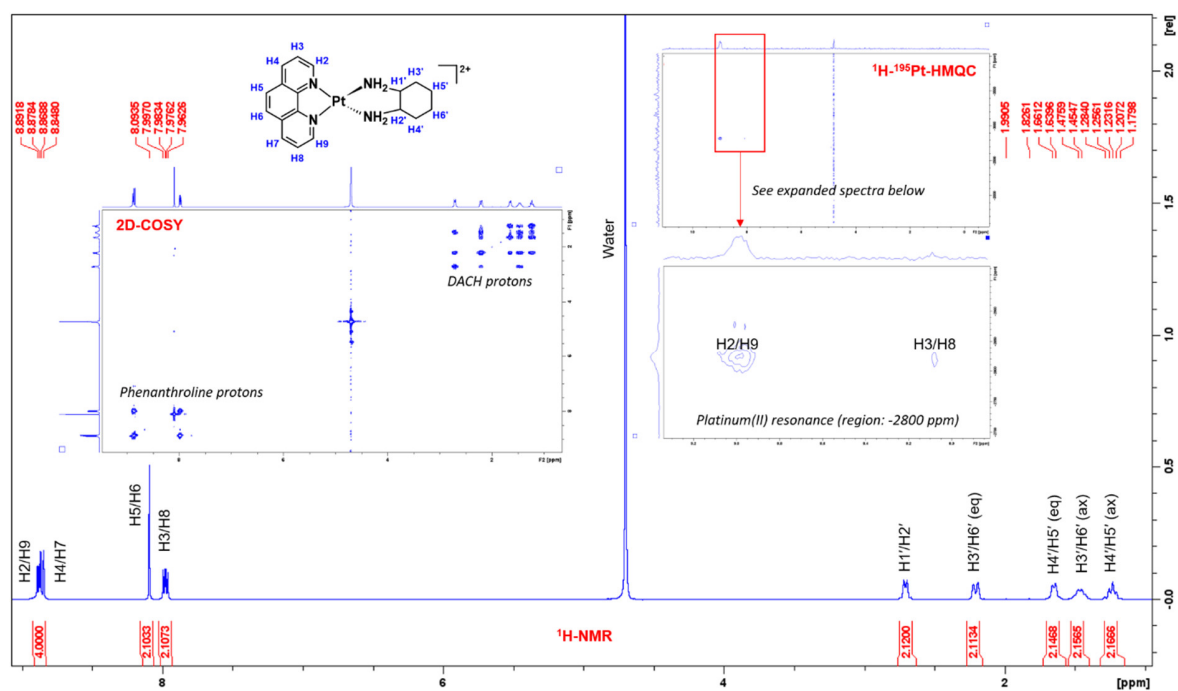


Figure S1: ¹H-NMR, 2D-COSY and ¹H-¹⁹⁵Pt-HMQC spectra of **PHENSS** in D₂O obtained at 298 K. Inset: structure of **PHENSS** with proton labeling system.

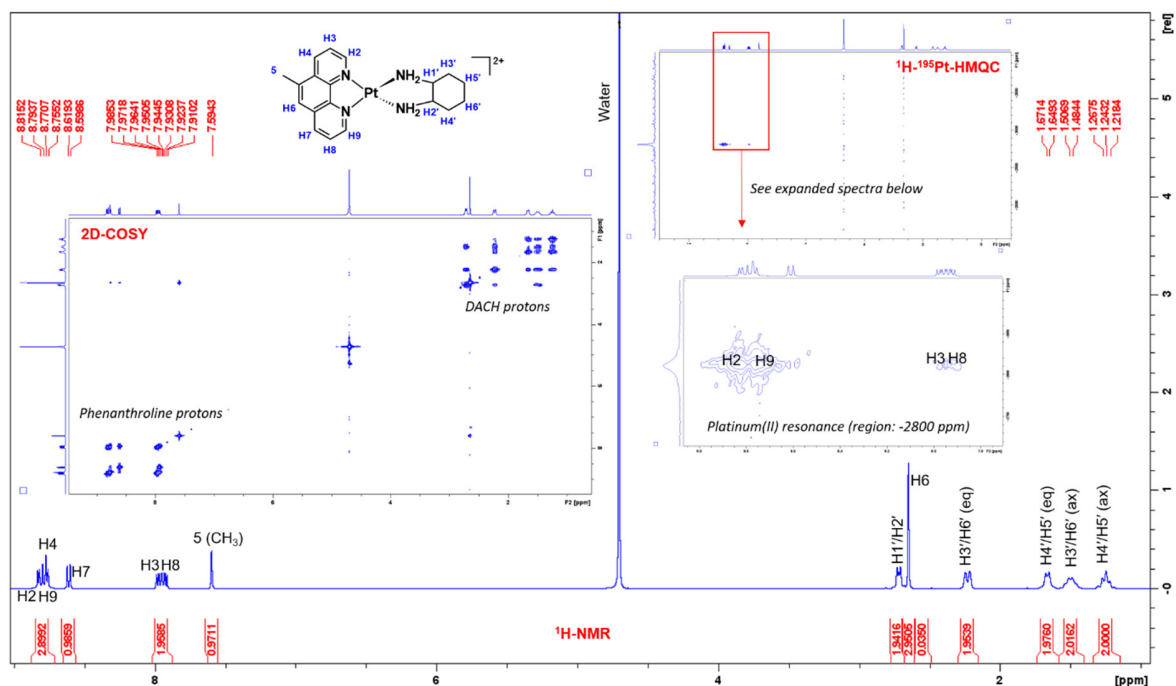


Figure S2: ^1H -NMR, 2D-COSY and ^1H - ^{195}Pt -HMQC spectra of **5MESS** in D_2O obtained at 298 K. Inset: structure of **5MESS** with proton labeling system.

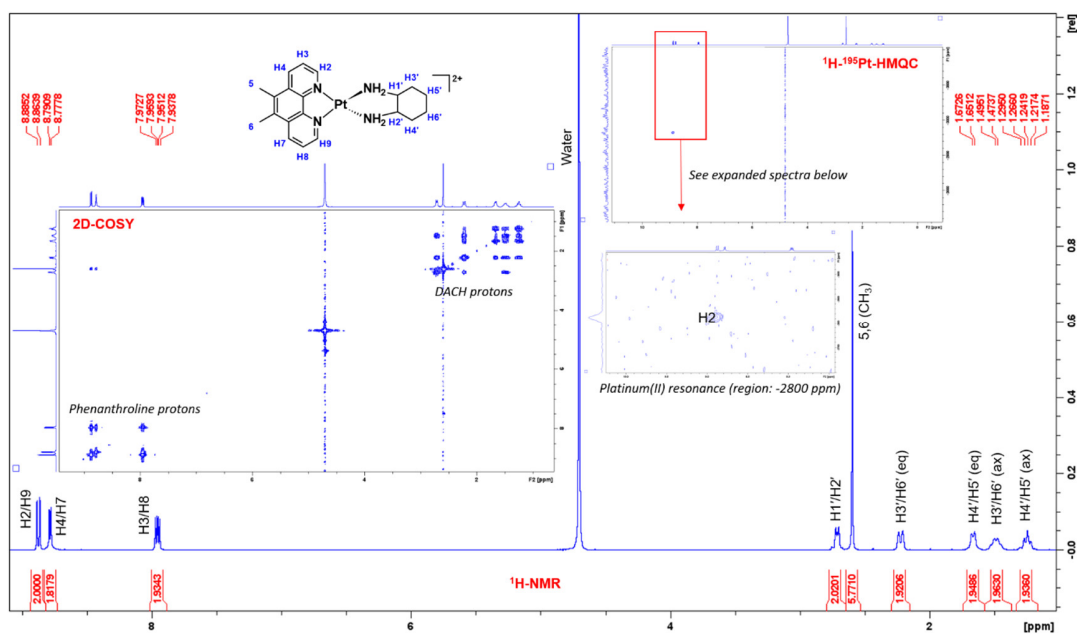


Figure S3: ^1H -NMR, 2D-COSY and ^1H - ^{195}Pt -HMQC spectra of **56MESS** in D_2O obtained at 298 K. Inset: structure of **56MESS** with proton labeling system.

NMR spectra of precursor platinum(IV) dihydroxy complexes

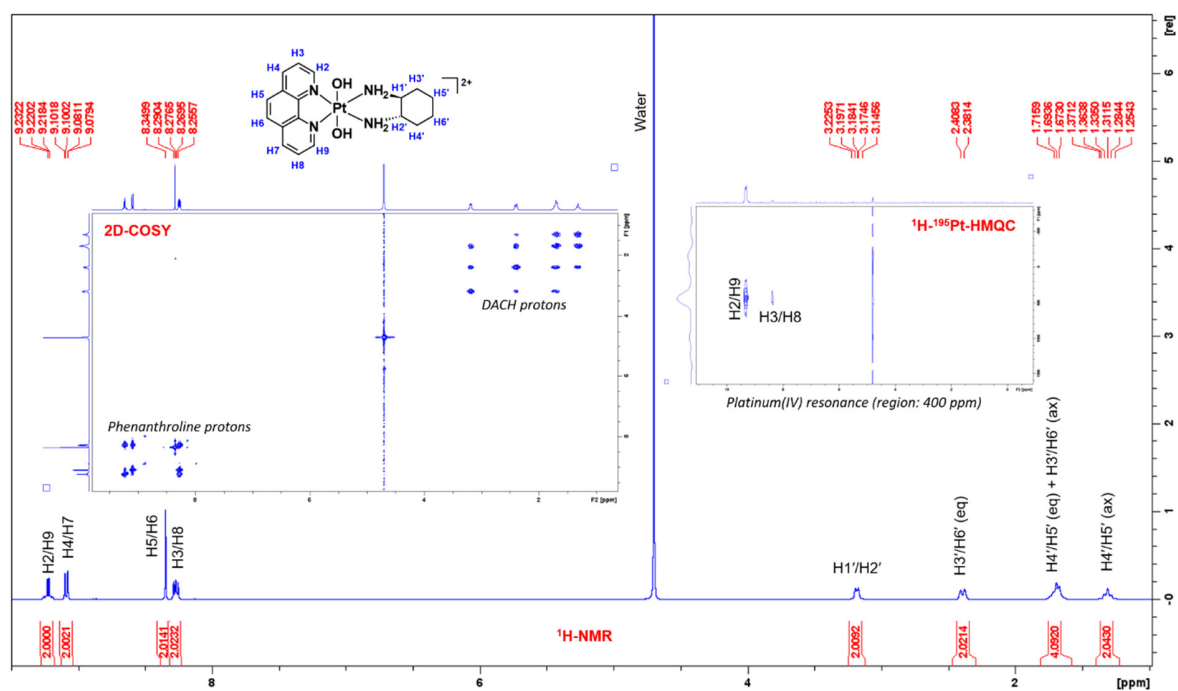


Figure S4: ^1H -NMR, 2D-COSY and ^1H - ^{195}Pt -HMQC spectra of **PHENSS(IV)(OH)₂** in D_2O obtained at 298 K. Inset: structure of **PHENSS(IV)(OH)₂** with proton labeling system.

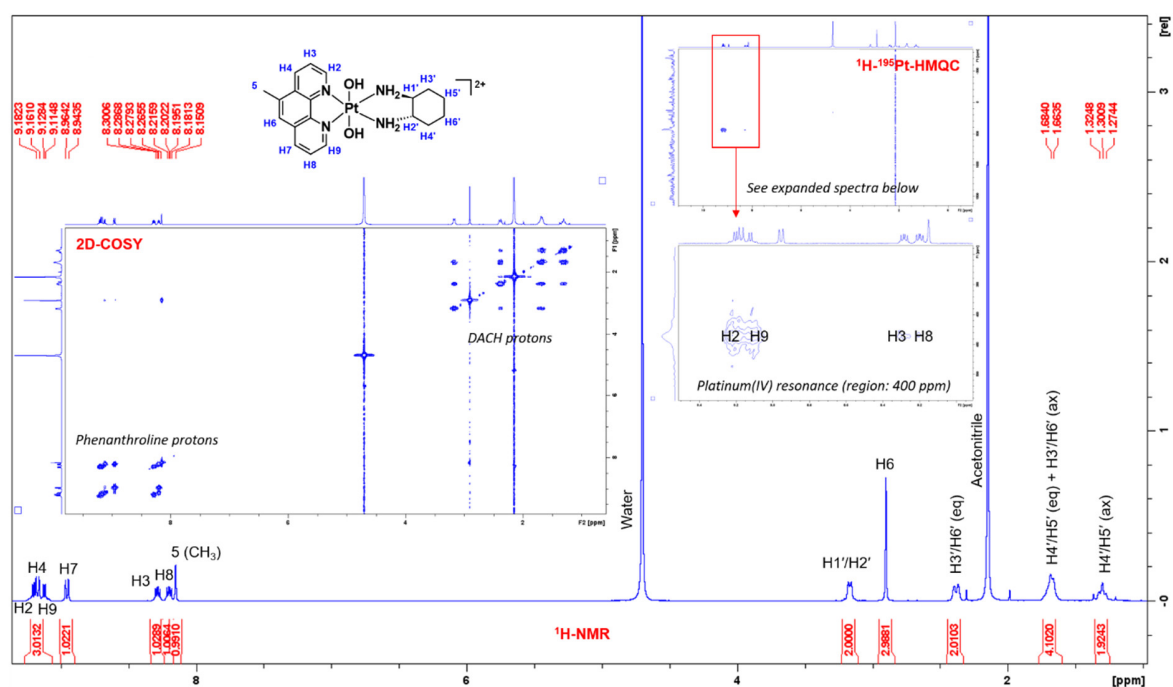


Figure S5: ^1H -NMR, 2D-COSY and ^1H - ^{195}Pt -HMQC spectra of **5MESS(IV)(OH)₂** in D_2O obtained at 298 K. Inset: structure of **5MESS(IV)(OH)₂** with proton labeling system.

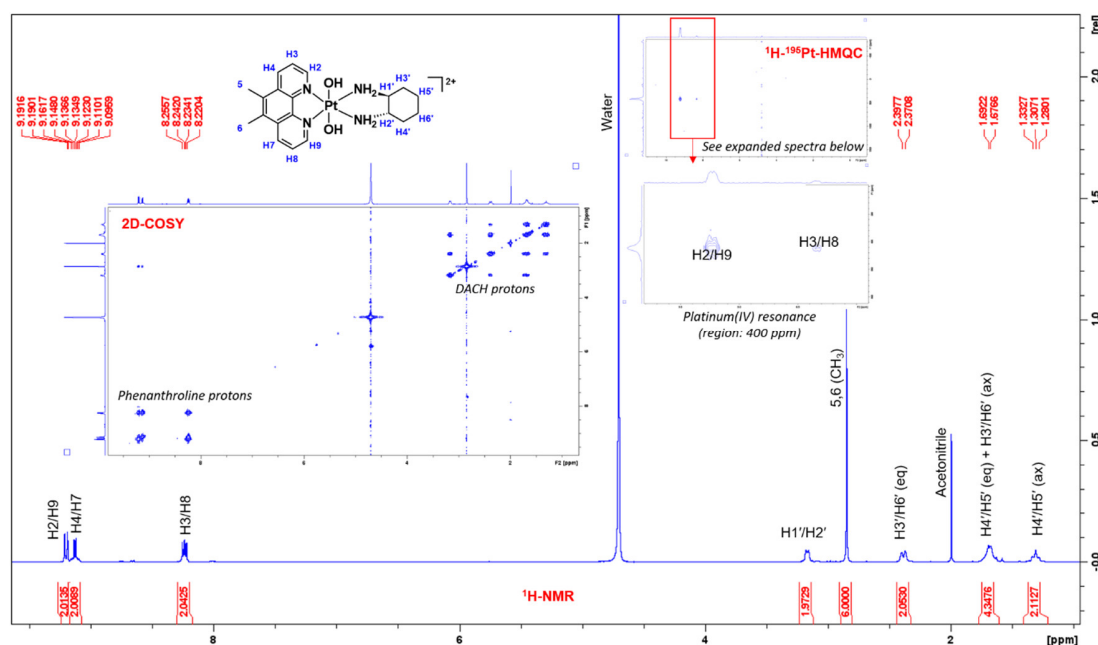


Figure S6: ^1H -NMR, 2D-COSY and ^1H - ^{195}Pt -HMQC spectra of **56MESS(IV)(OH)₂** in D_2O obtained at 298 K. Inset: structure of **56MESS(IV)(OH)₂** with proton labeling system.

HPLC chromatogram of CLB anhydride

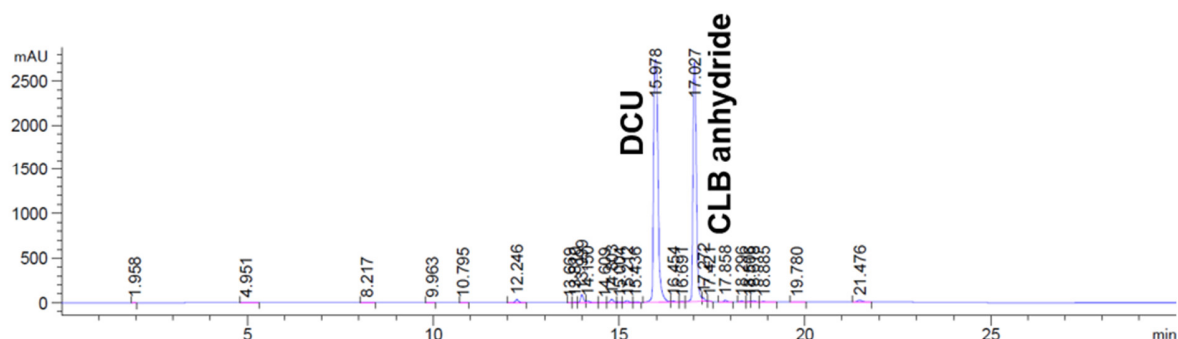


Figure S7: HPLC chromatogram of CLB anhydride (at 17 min) with DCU by-product (at 15.9 min), within the collection wavelength of 254 nm obtained at 298 K, by an Agilent ZORBAX RX-C₁₈ column (100 × 4.6 mm, 3.5 μm pore size).

HPLC chromatograms of platinum(IV)-CLB complexes

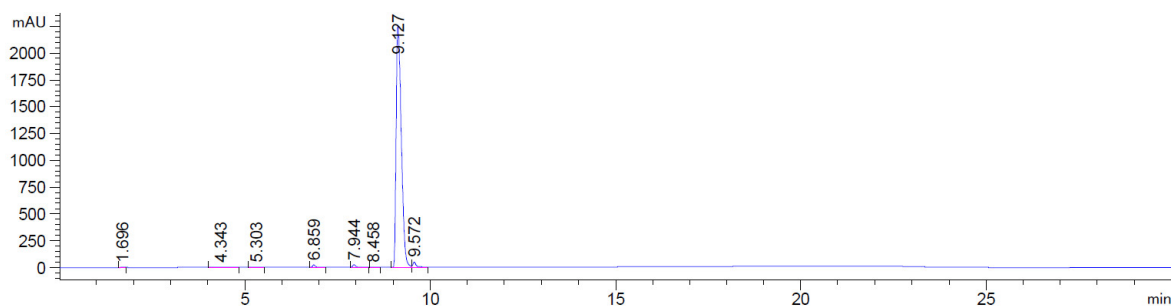


Figure S8: HPLC chromatogram of **PCLB** within the collection wavelength of 254 nm obtained at 298 K, by a Phenomenex Onyx™ Monolithic C₁₈-reverse phase column (100 × 4.6 mm, 5 μm pore size).

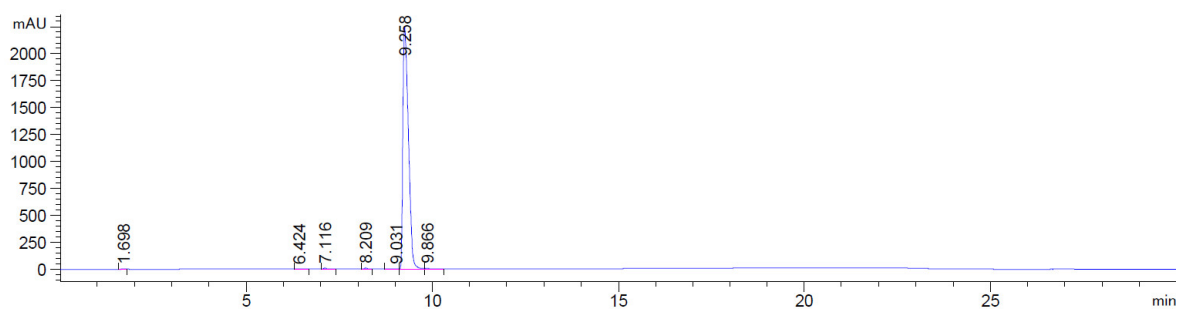


Figure S9: HPLC chromatogram of **5CLB** within the collection wavelength of 254 nm obtained at 298 K, by a Phenomenex Onyx™ Monolithic C₁₈-reverse phase column (100 × 4.6 mm, 5 μm pore size).

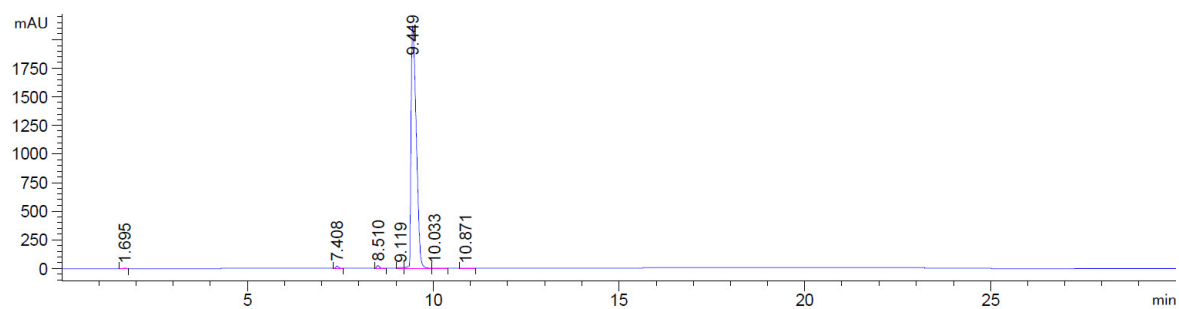


Figure S10: HPLC chromatogram of **56CLB** within the collection wavelength of 254 nm obtained at 298 K, by a Phenomenex Onyx™ Monolithic C₁₈-reverse phase column (100 × 4.6 mm, 5 μm pore size).

NMR spectra of platinum(IV)-CLB complexes

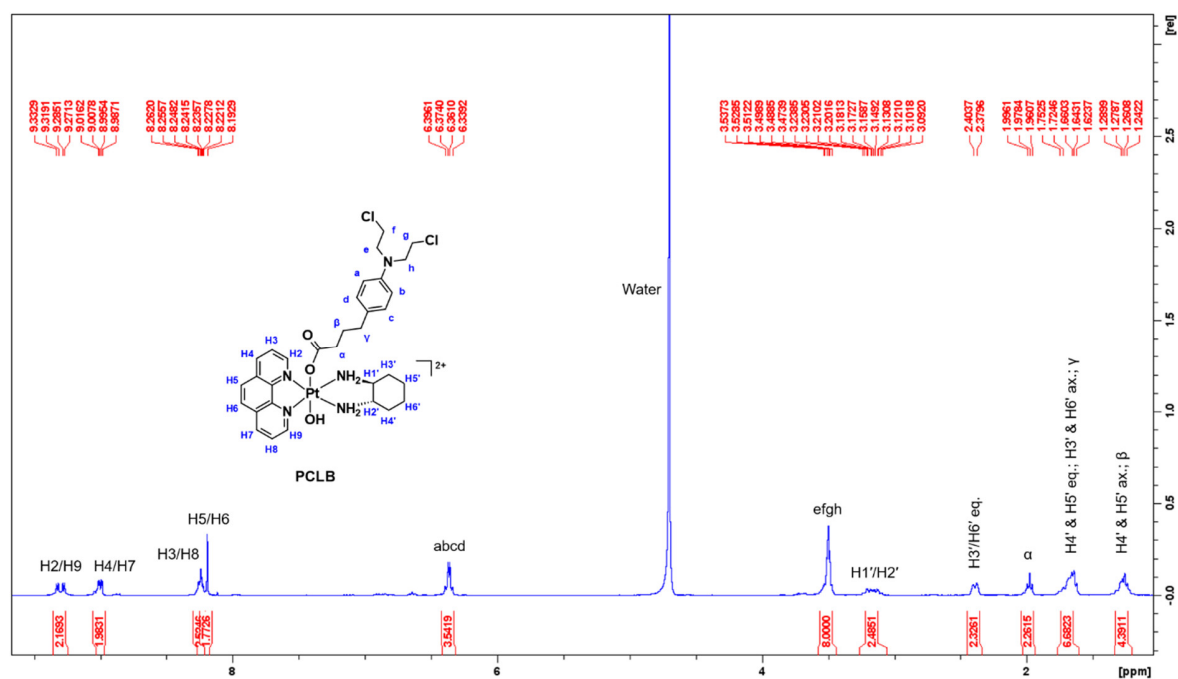
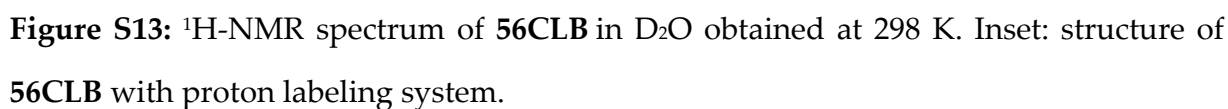


Figure S11: ^1H -NMR spectrum of PCLB in D_2O obtained at 298 K. Inset: structure of PCLB with proton labeling system.



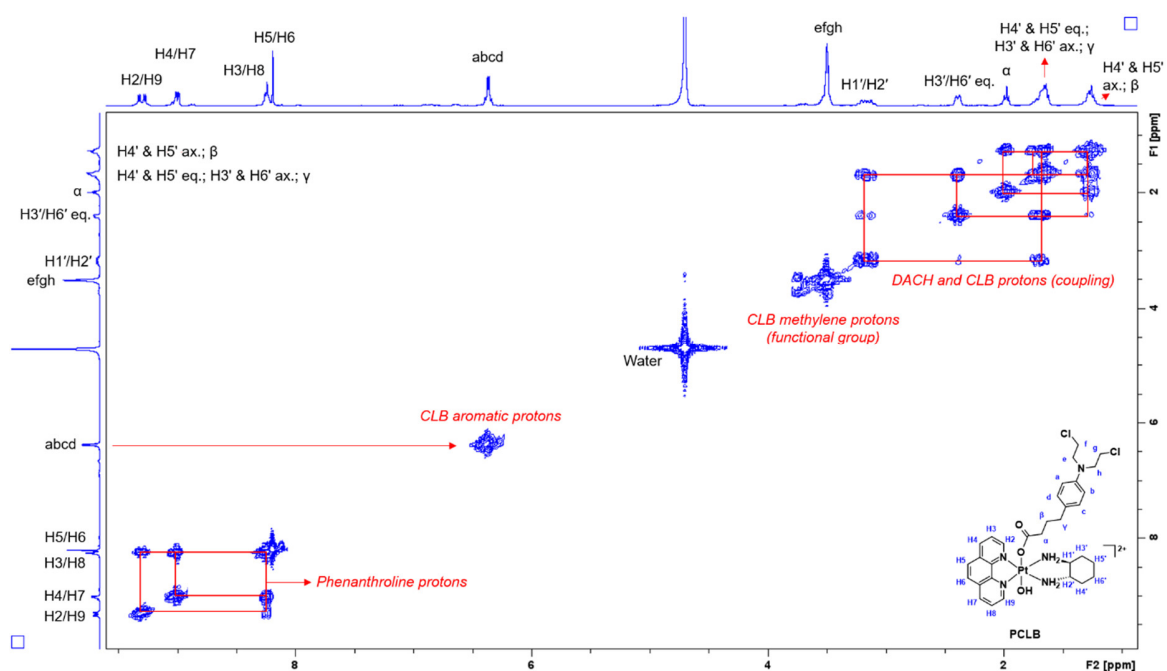


Figure S14: 2D-COSY spectrum of **PCLB** in D₂O obtained at 298 K. Inset: structure of **PCLB** with proton labeling system.

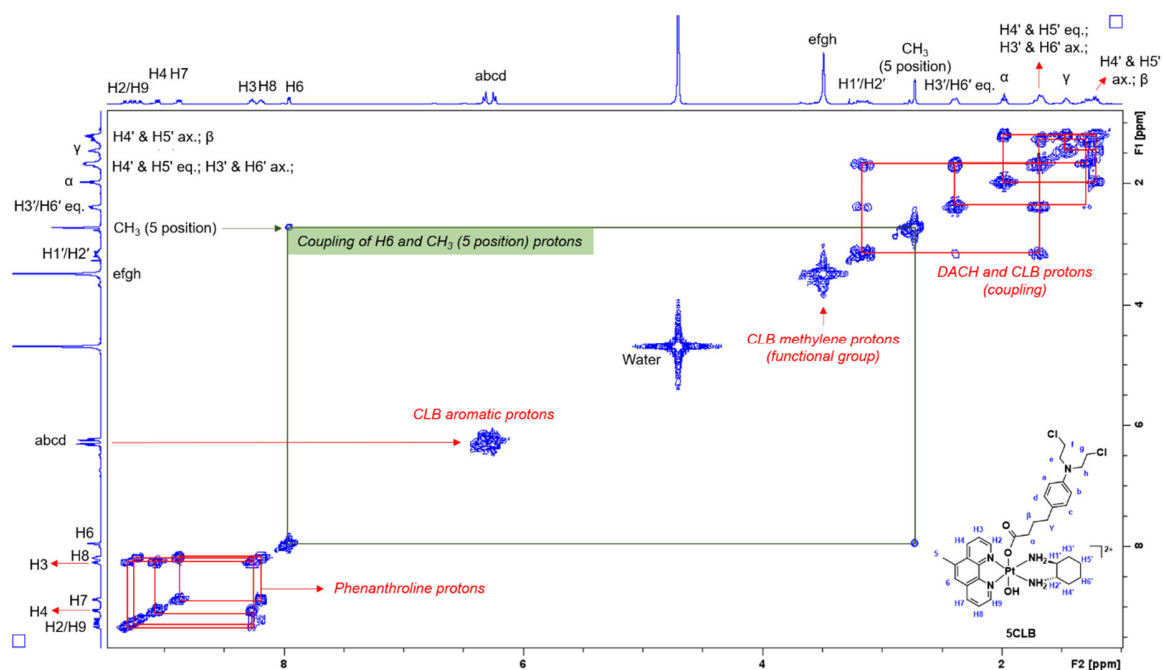


Figure S15: 2D-COSY spectrum of **5CLB** in D₂O obtained at 298 K. Inset: structure of **5CLB** with proton labeling system.

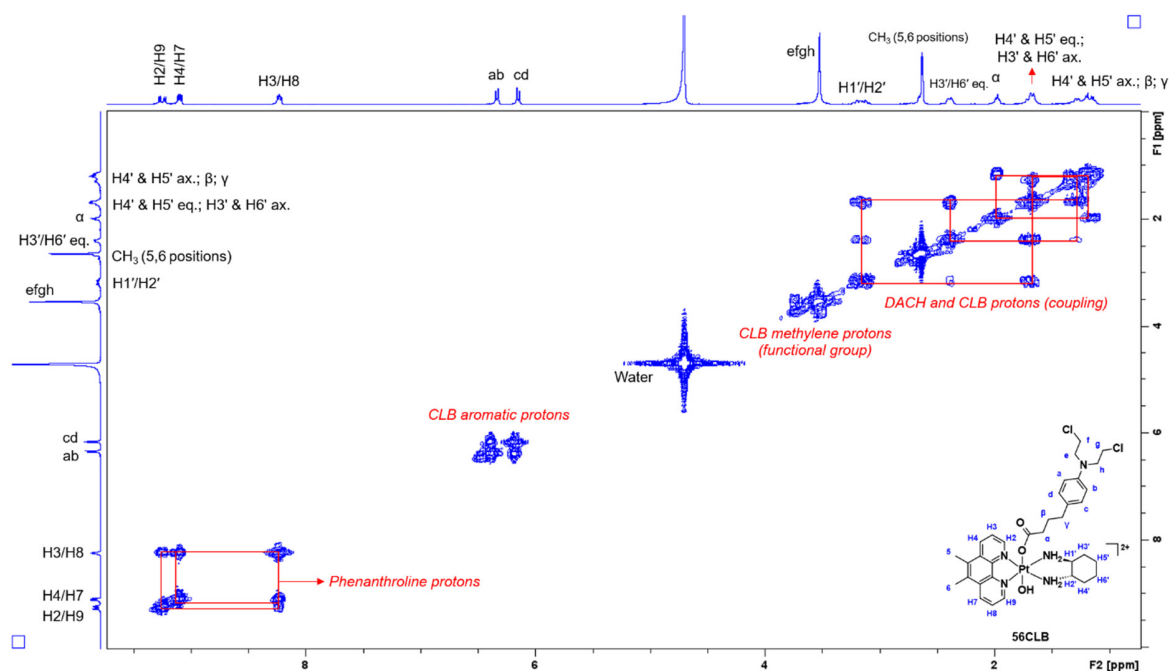


Figure S16: 2D-COSY spectrum of **56CLB** in D_2O obtained at 298 K. Inset: structure of **56CLB** with proton labeling system.

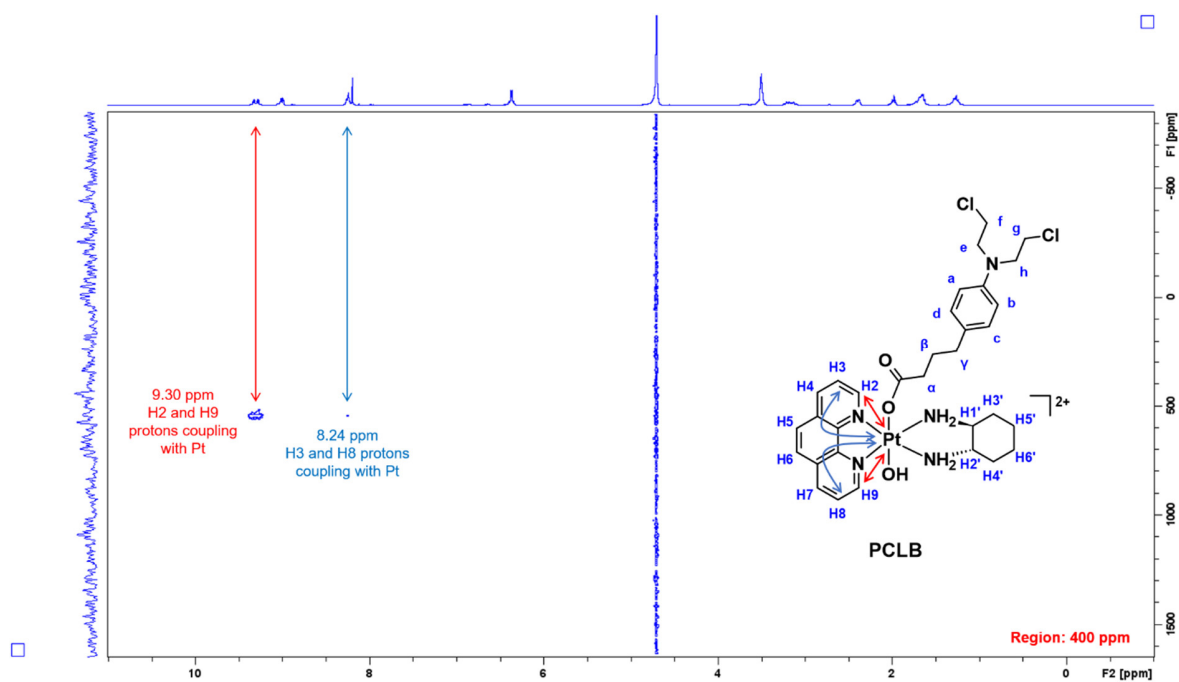


Figure S17: ^1H - ^{195}Pt -HMQC spectrum of **PCLB** in D_2O obtained at 298 K. Region: 400 ppm. Inset: structure of **PCLB** with proton labelling system and arrows that indicate coupling.

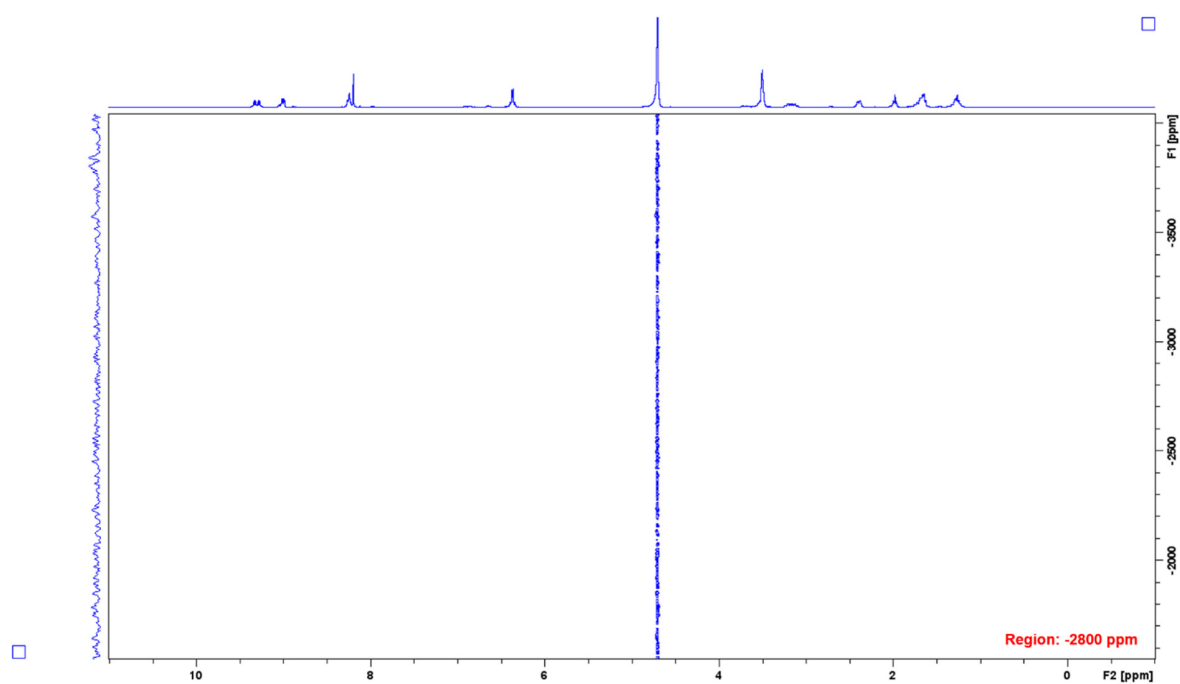


Figure S18: ^1H - ^{195}Pt -HMQC spectrum of **PCLB** in D_2O obtained at 298 K. Region: -2800 ppm. Absence of platinum(II) resonance.

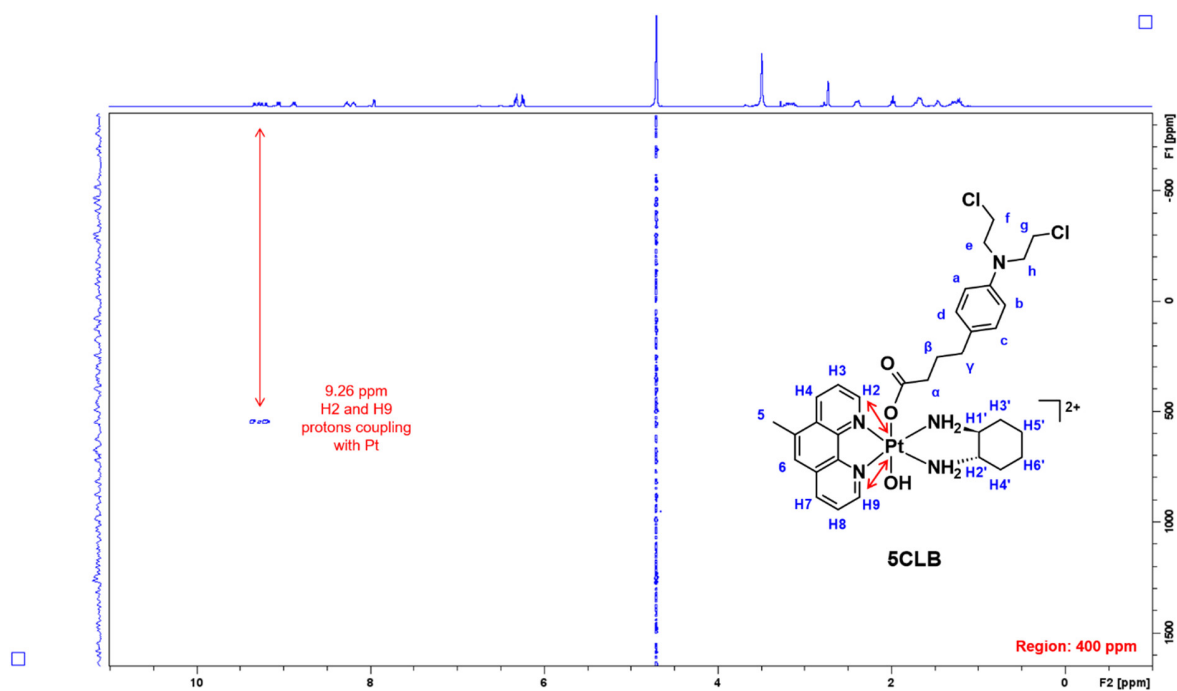


Figure S19: ^1H - ^{195}Pt -HMQC spectrum of **5CLB** in D_2O obtained at 298 K. Region: 400 ppm. Inset: structure of **5CLB** with proton labelling system and arrow that indicate coupling.

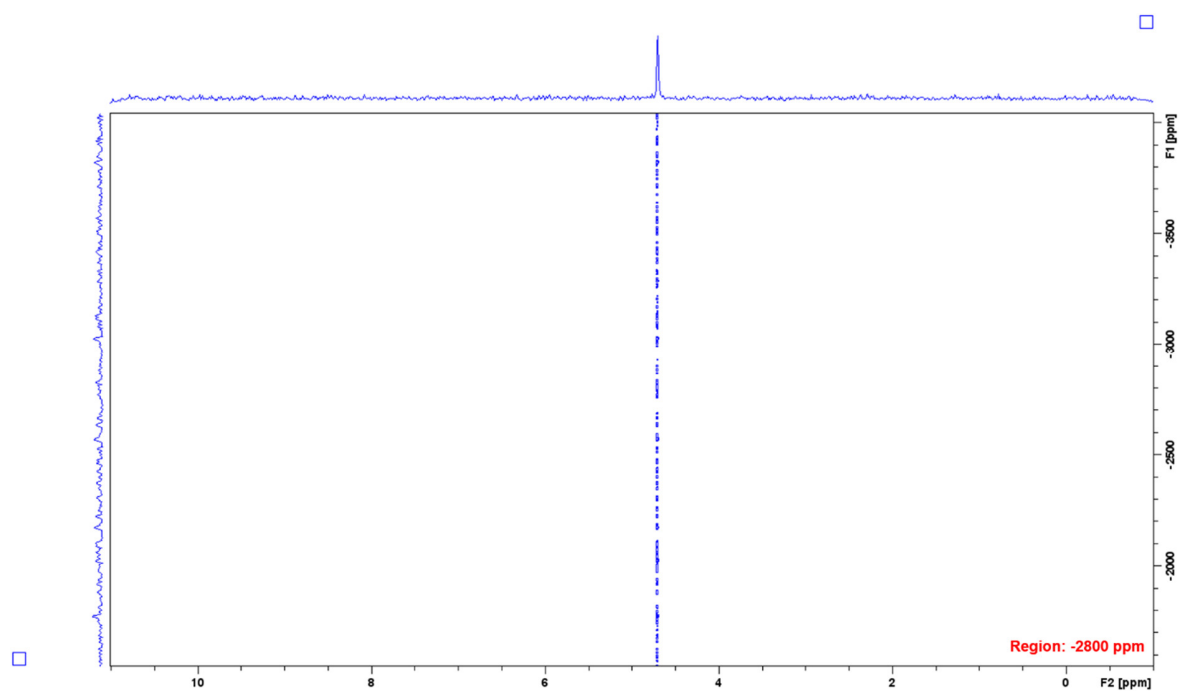


Figure S20: ^1H - ^{195}Pt -HMQC spectrum of **5CLB** in D_2O obtained at 298 K. Region: -2800 ppm. Absence of platinum(II) resonance.

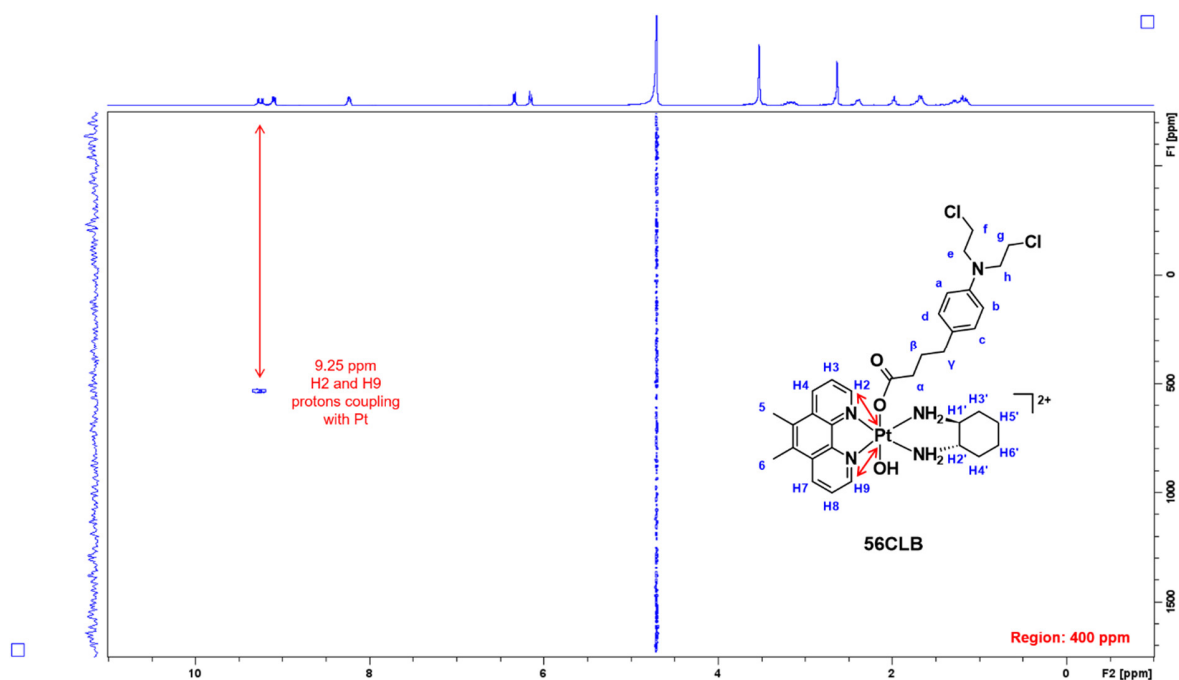


Figure S21: ^1H - ^{195}Pt -HMQC spectrum of **56CLB** in D_2O obtained at 298 K. Region: 400 ppm. Inset: structure of **56CLB** with proton labelling system and arrows that indicate coupling.

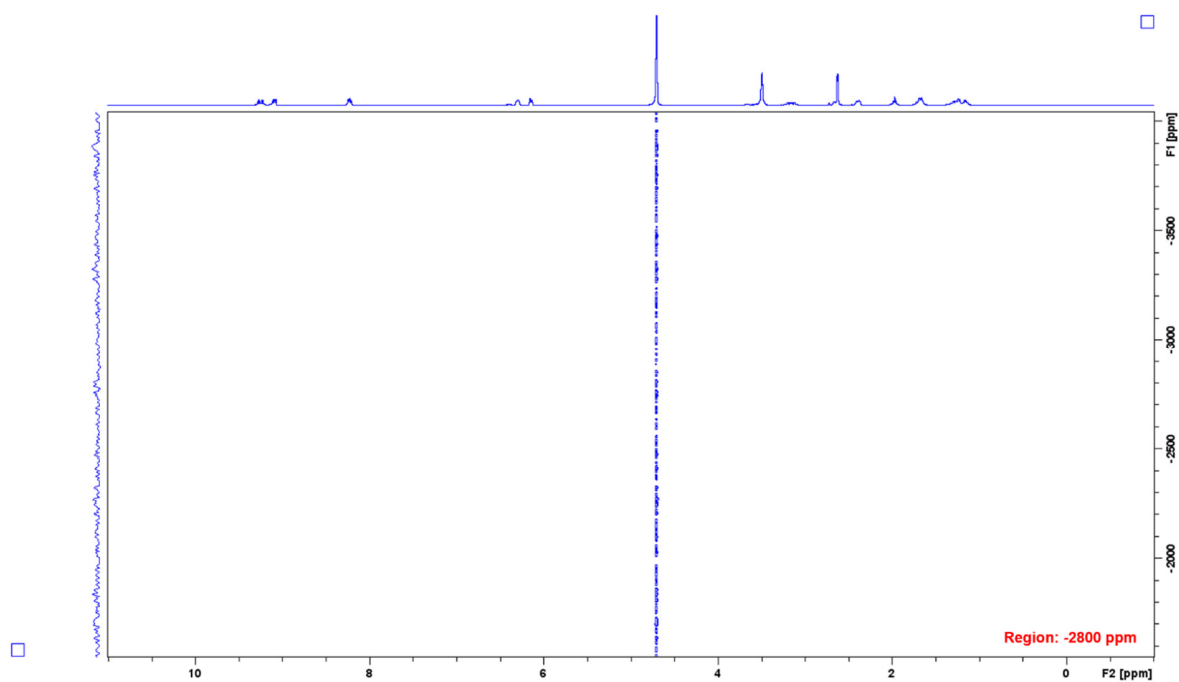


Figure S22: ^1H - ^{195}Pt -HMQC spectrum of **56CLB** in D_2O obtained at 298 K. Region: -2800 ppm. Absence of platinum(II) resonance.

UV spectra of platinum(IV)-CLB complexes

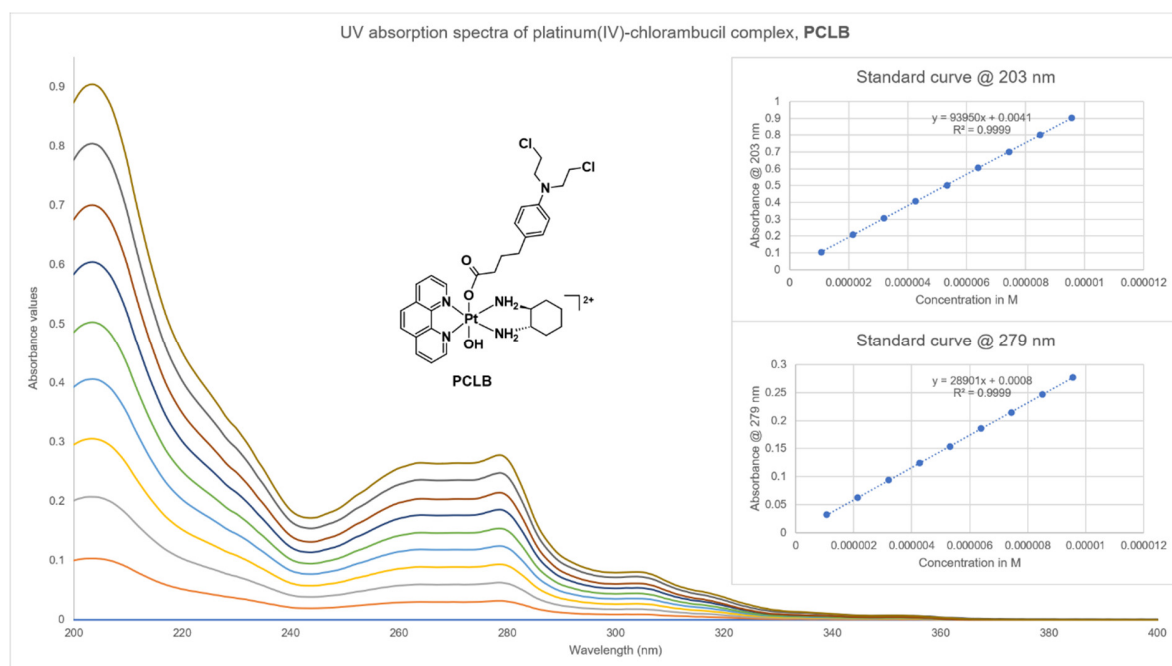


Figure S23: UV spectra of **PCLB** obtained at 298 K. Inset: structure of **PCLB** and generated plot curves within the wavelengths 203 and 279 nm at 298 K.

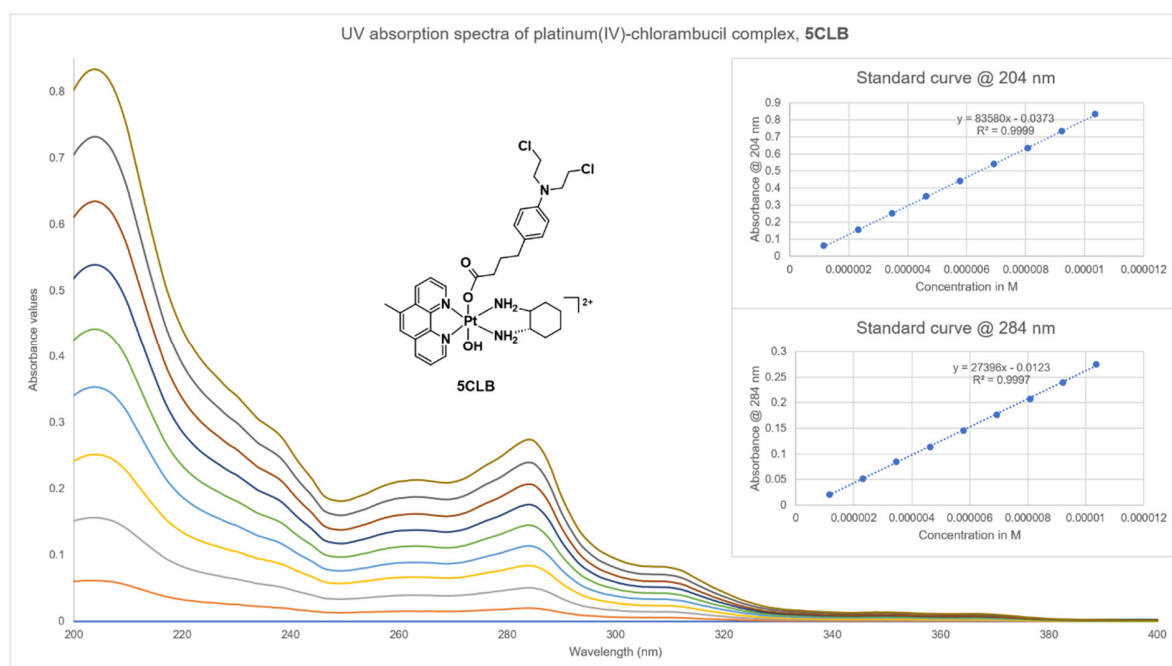


Figure S24: UV spectra of **5CLB** obtained at 298 K. Inset: structure of **5CLB** and generated plot curves within the wavelengths 204 and 284 nm at 298 K.

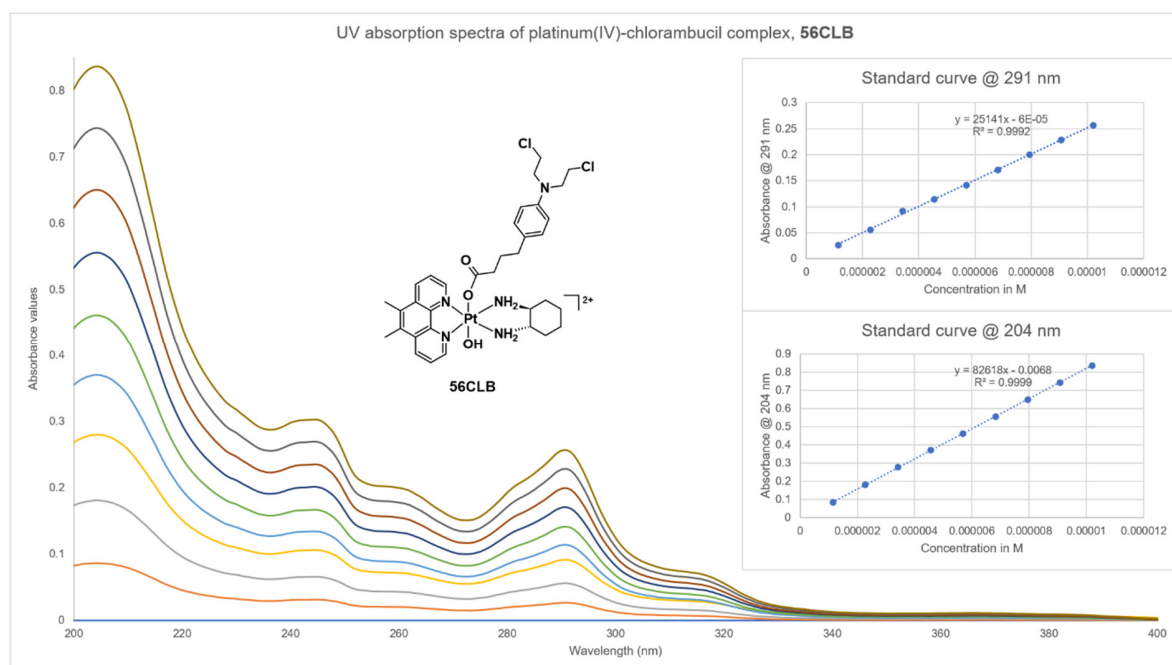


Figure S25: UV spectra of **56CLB** obtained at 298 K. Inset: structure of **56CLB** and generated plot curves within the wavelengths 204 and 291 nm at 298 K.

CD spectra of platinum(IV)-CLB complexes

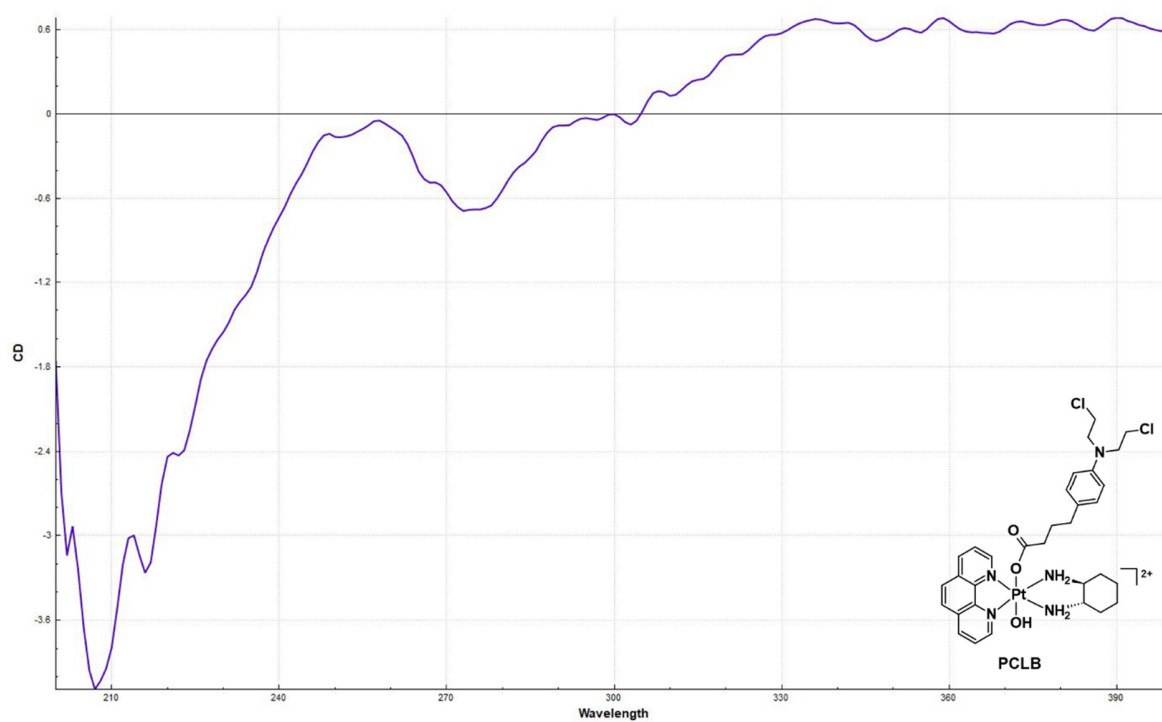


Figure S26: CD spectrum of **PCLB** in d.i.H₂O obtained at 298 K. Inset: structure of **PCLB**.

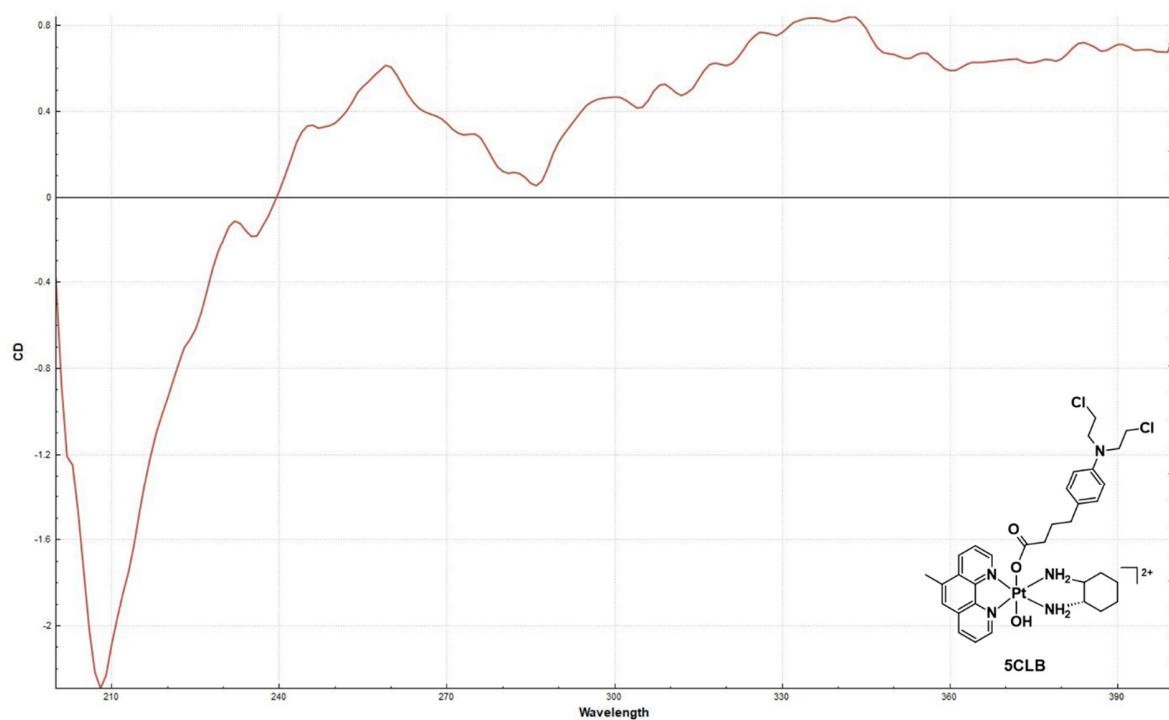


Figure S27: CD spectrum of **5CLB** in d.i.H₂O obtained at 298 K. Inset: structure of **5CLB**.

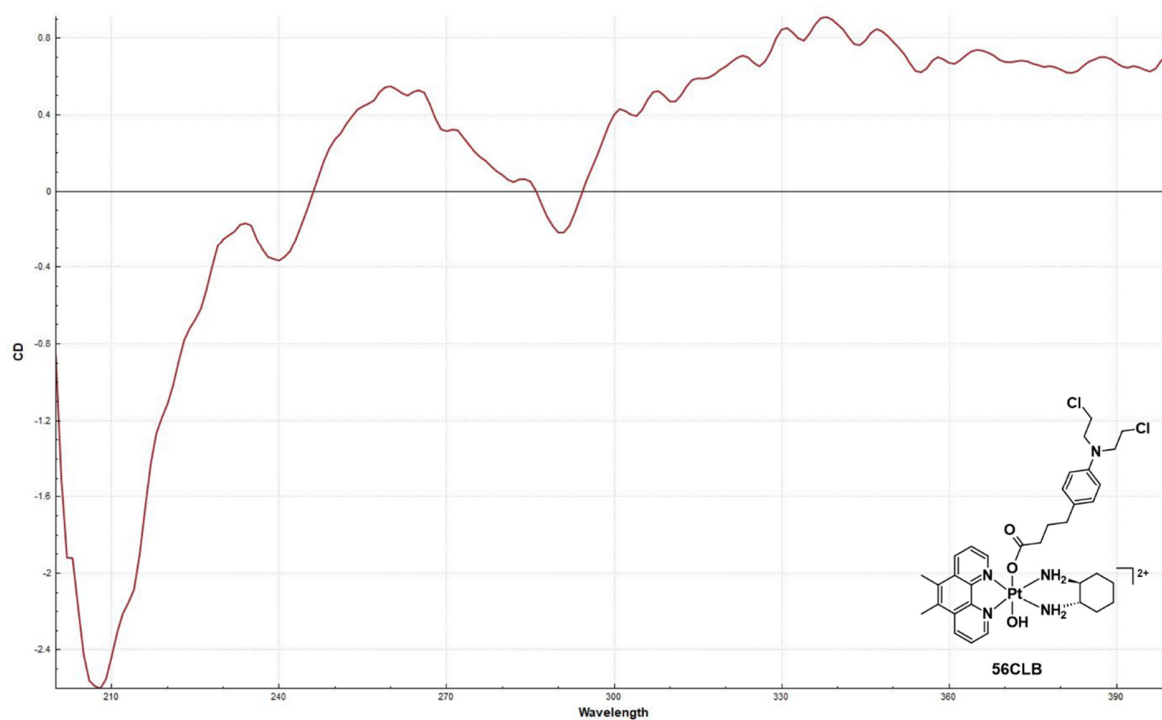


Figure S28: CD spectrum of **56CLB** in d.i.H₂O obtained at 298 K. Inset: structure of **56CLB**.

ESI-MS spectra of platinum(IV)-CLB complexes

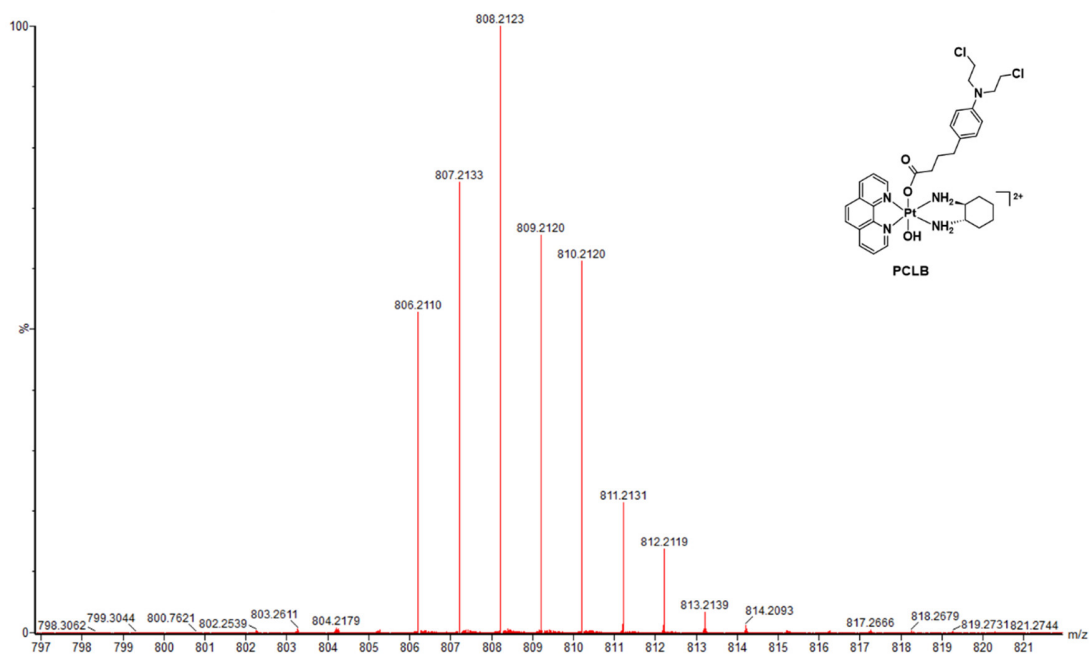


Figure S29: ESI-MS spectrum of **PCLB** in d.i.H₂O obtained at 298 K, using a Waters SYNAPT G2-Si quadrupole time-of-flight (QTOF) HDMS. Inset: structure of **PCLB**.

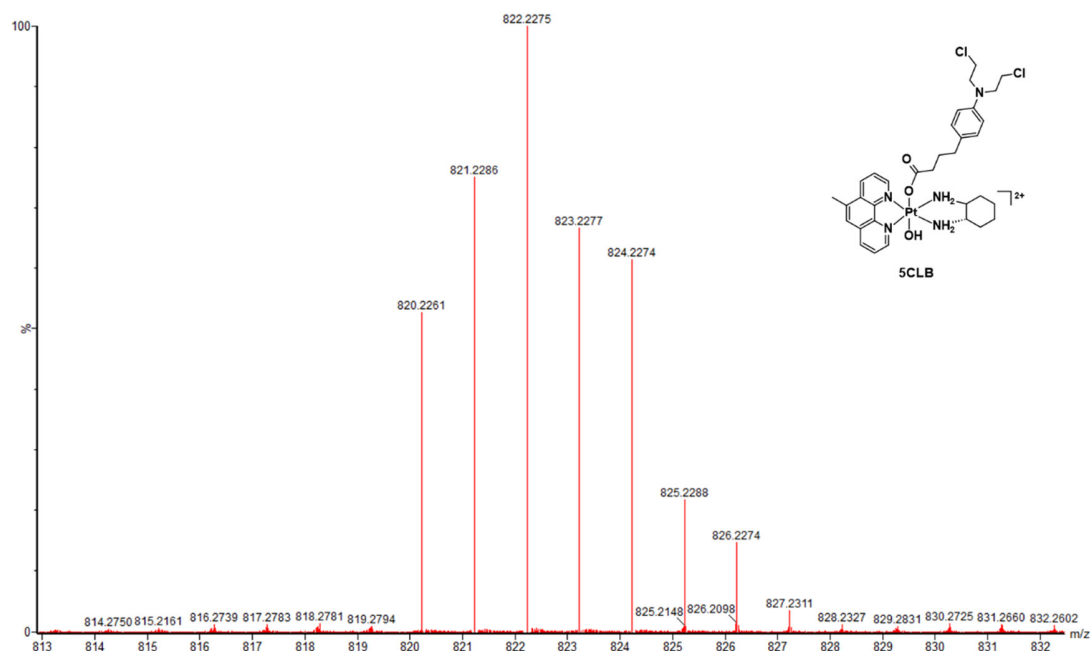


Figure S30: ESI-MS spectrum of **5CLB** in d.i.H₂O obtained at 298 K, using a Waters SYNAPT G2-Si quadrupole time-of-flight (QTOF) HDMS. Inset: structure of **5CLB**.

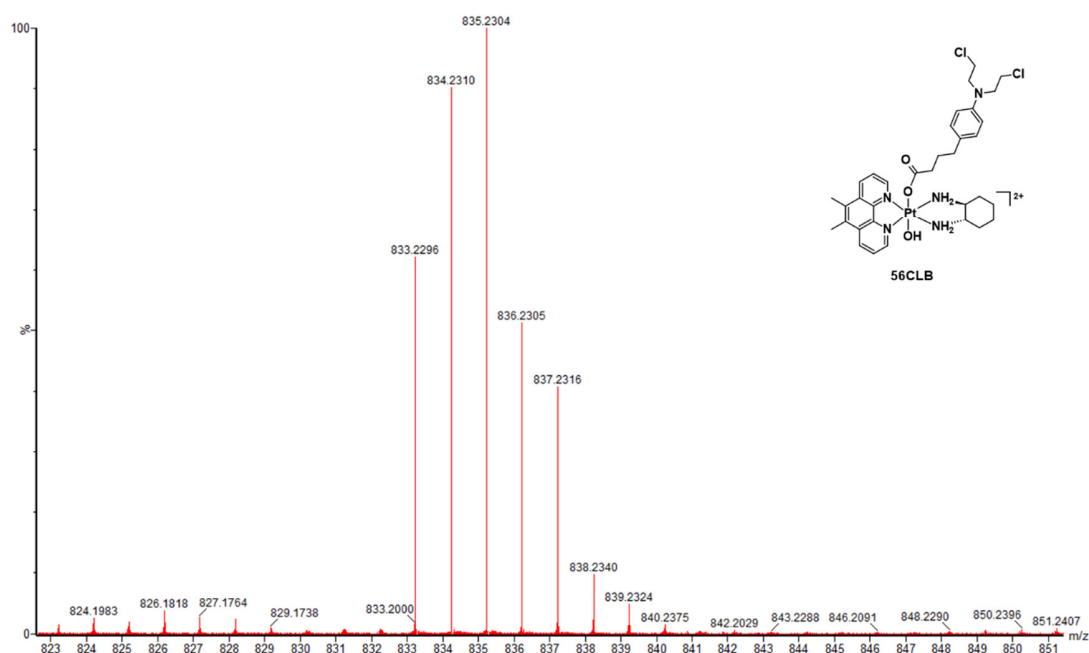


Figure S31: ESI-MS spectrum of **56CLB** in d.i.H₂O obtained at 298 K, using a Waters SYNAPT G2-Si quadrupole time-of-flight (QTOF) HDMS. Inset: structure of **56CLB**.

Stability experiments

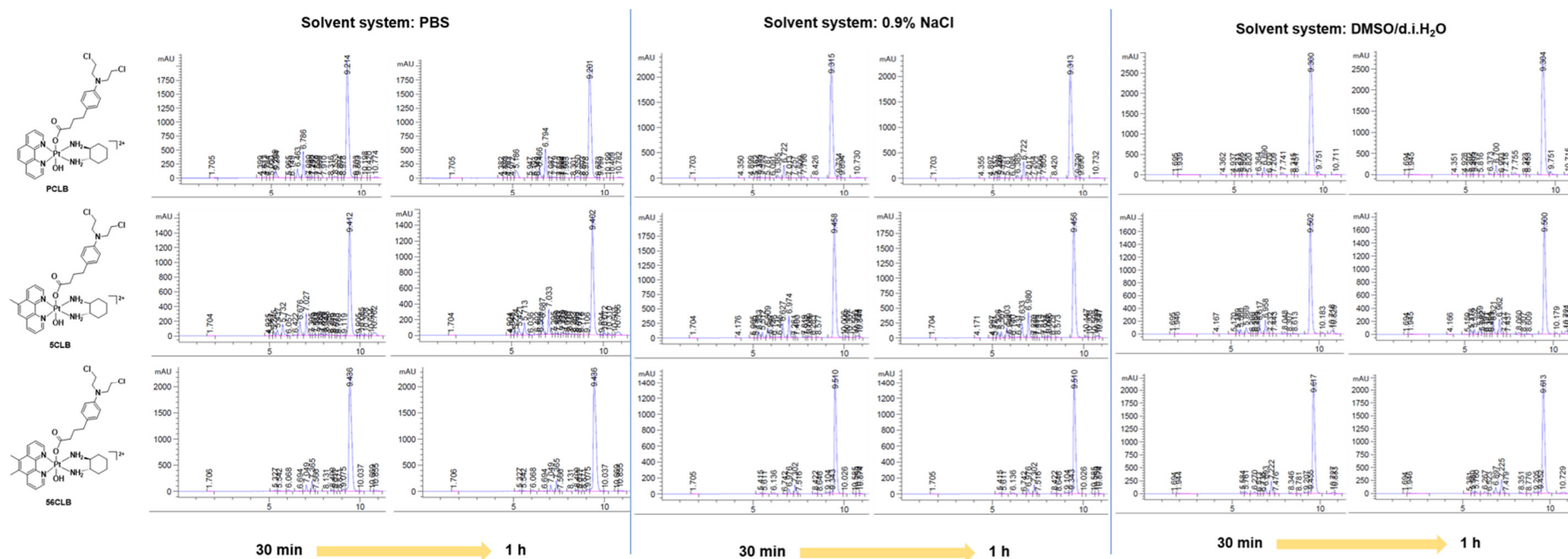


Figure S32: Combined HPLC chromatograms of **PCLB**, **5CLB** and **56CLB** in three solvent systems, obtained at 298 K. Inset: structures of **PCLB**, **5CLB** and **56CLB**.

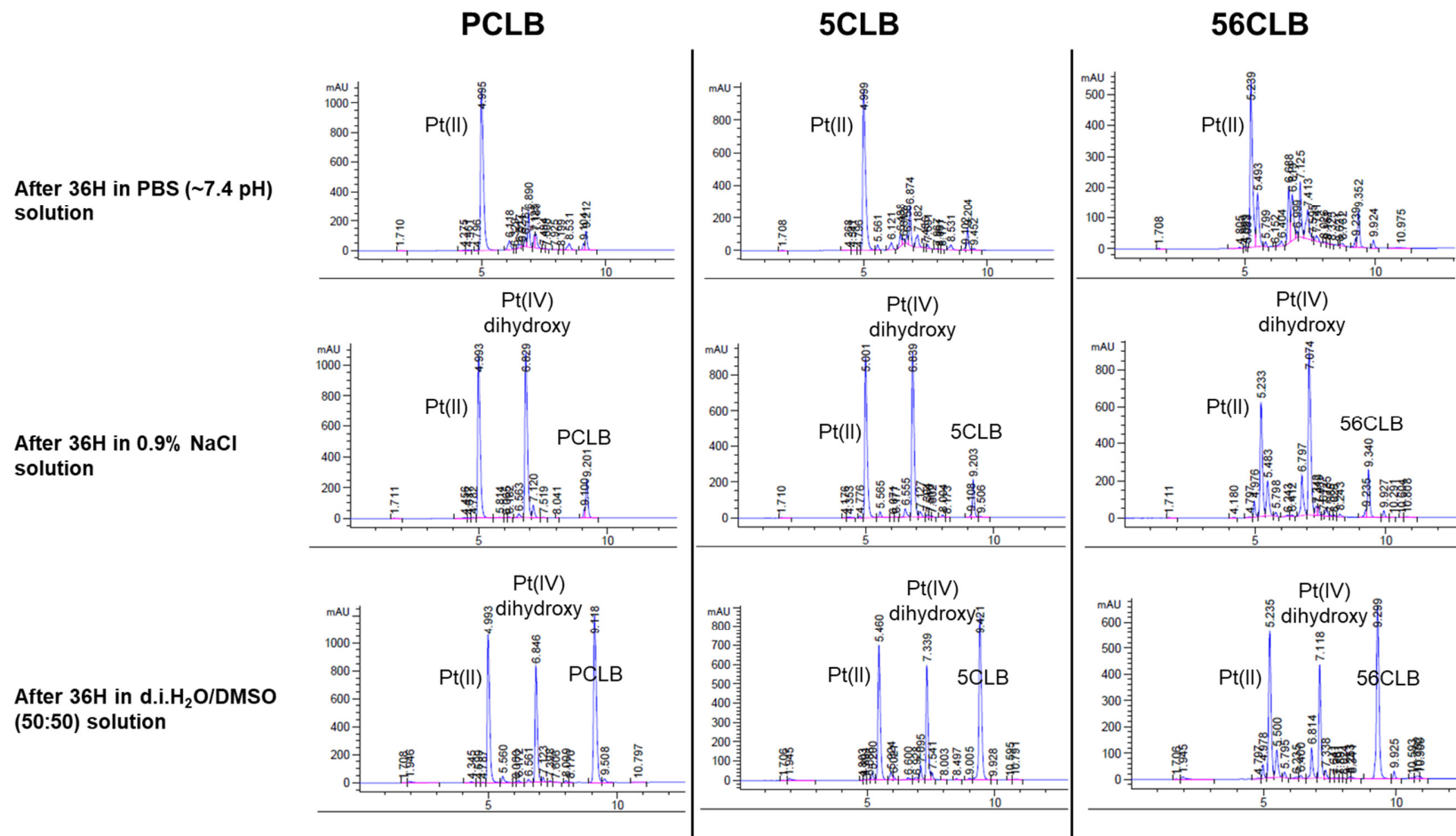


Figure S33: Combined HPLC chromatograms of **PCLB**, **5CLB** and **56CLB** in three solvent systems, obtained at 298 K at 36 h. Inset: structures of **PCLB**, **5CLB** and **56CLB**.

Lipophilicity

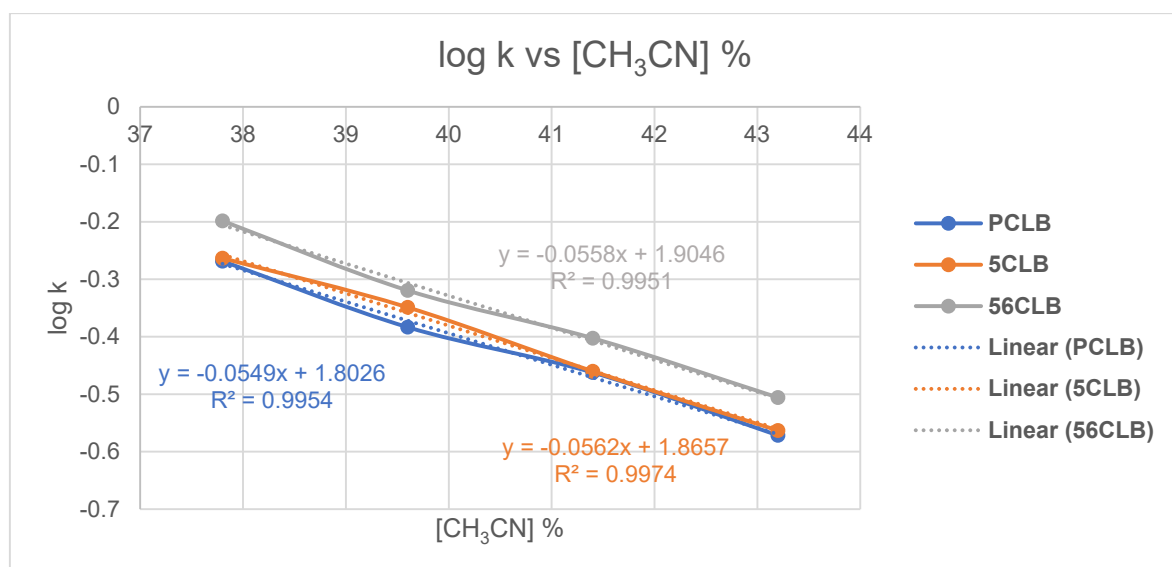


Figure S34: Generated plot curve of log *k* *versus* concentration of organic solvent, CH₃CN to determine the chromatographic lipophilicity index, log *k*_w of the platinum(IV)-CLB complexes.

Reduction reaction experiments

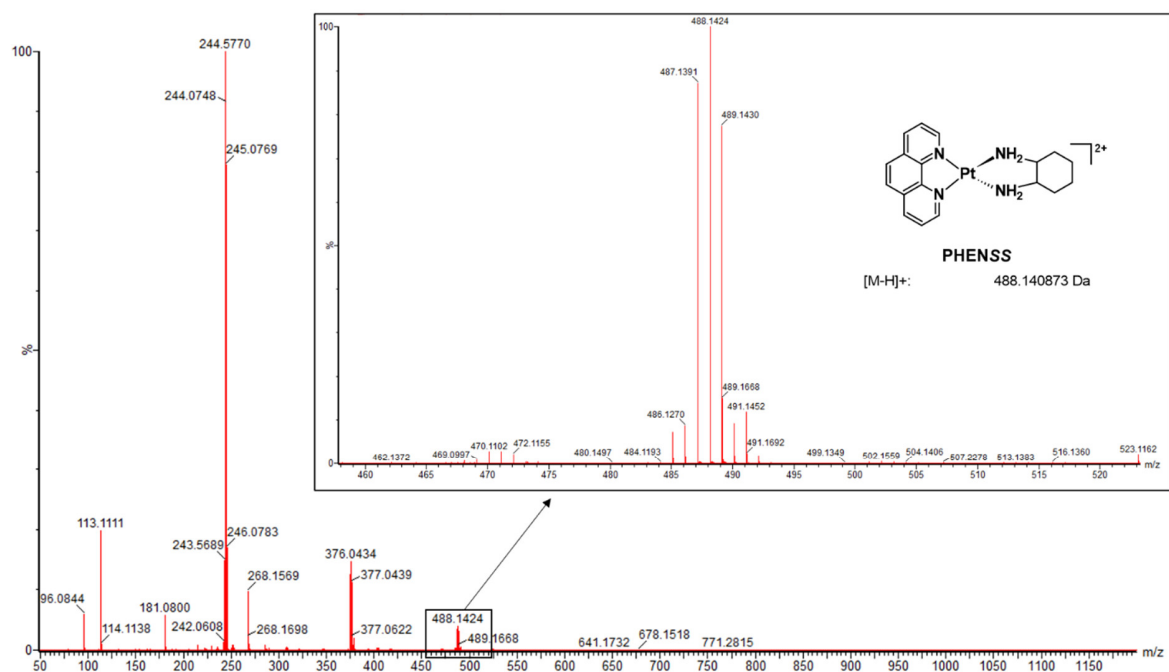


Figure S35: ESI-MS spectra of the reduced **PCLB** prodrug, highlighting the formation of the corresponding platinum(II) congener, **PHENSS**.

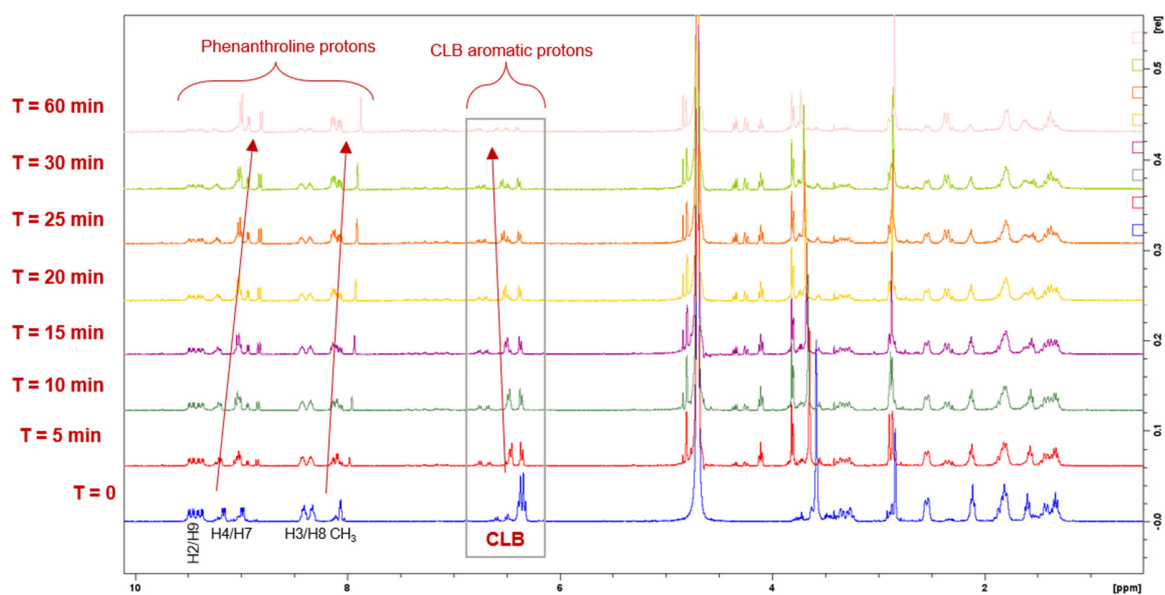


Figure S36: ^1H -NMR spectra of **5CLB** with PBS and AsA in D_2O at 310.15 K, in different time intervals, highlighting the movement of resonances from the phenanthroline protons and the aromatic protons of the CLB ligand. **T** represents time in min.

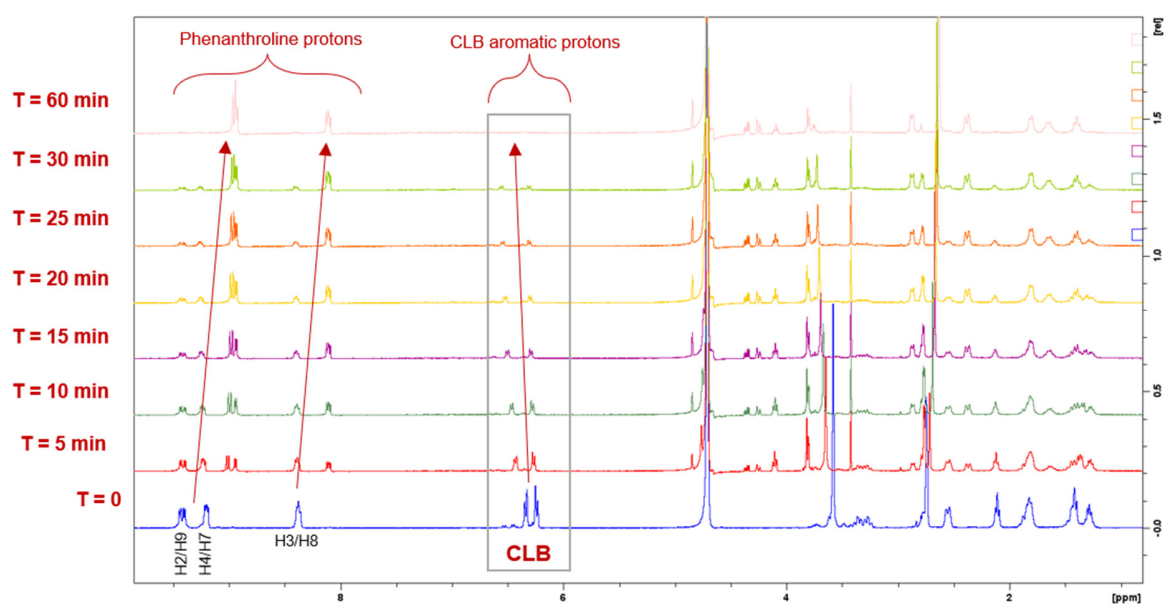


Figure S37: ^1H -NMR spectra of **56CLB** with PBS and AsA in D_2O at 310.15 K, in different time intervals, highlighting the movement of resonances from the phenanthroline protons and the aromatic protons of the CLB ligand. **T** represents time in min.

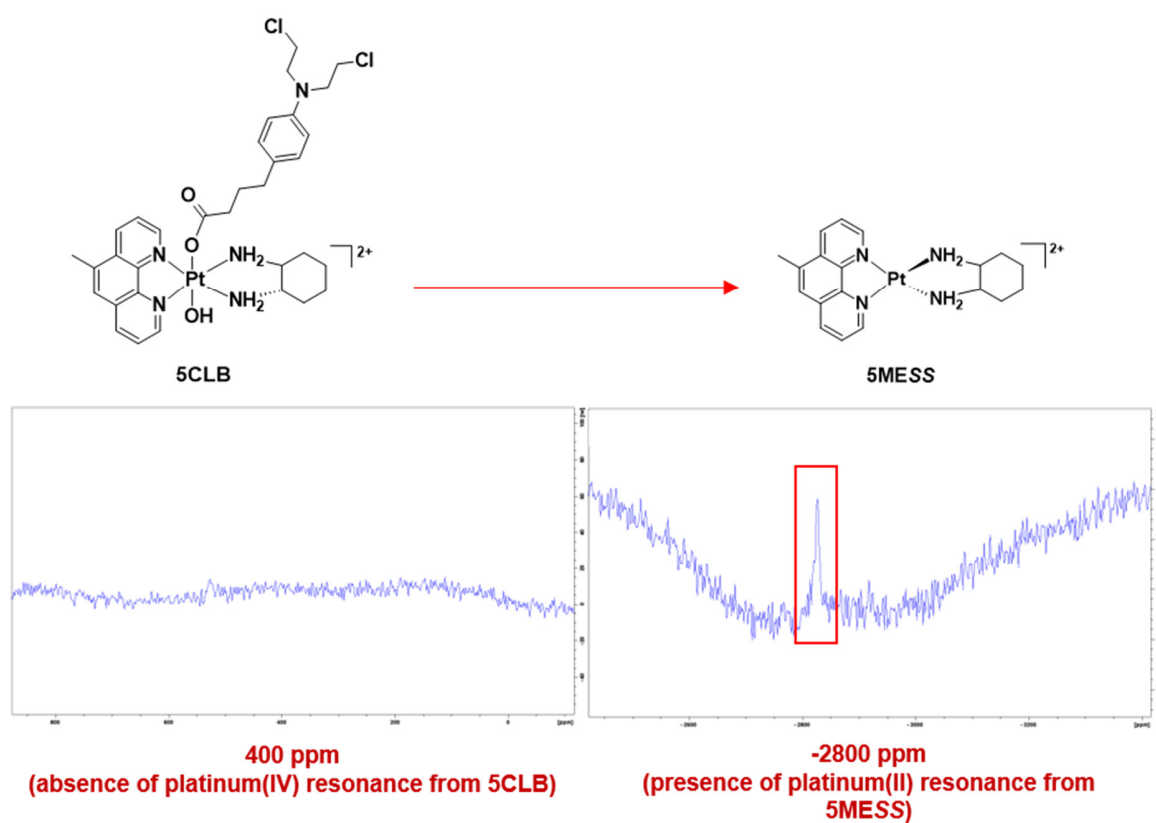


Figure S38: 1D- ^{195}Pt NMR spectra of **5CLB** with PBS and AsA in D_2O at 310.15 K, within the regions of 400 and -2800 ppm, highlighting the complete reduction of the complex after 1 h. Above: Structures of **5CLB** and its platinum(II) scaffold, **5MESS**.

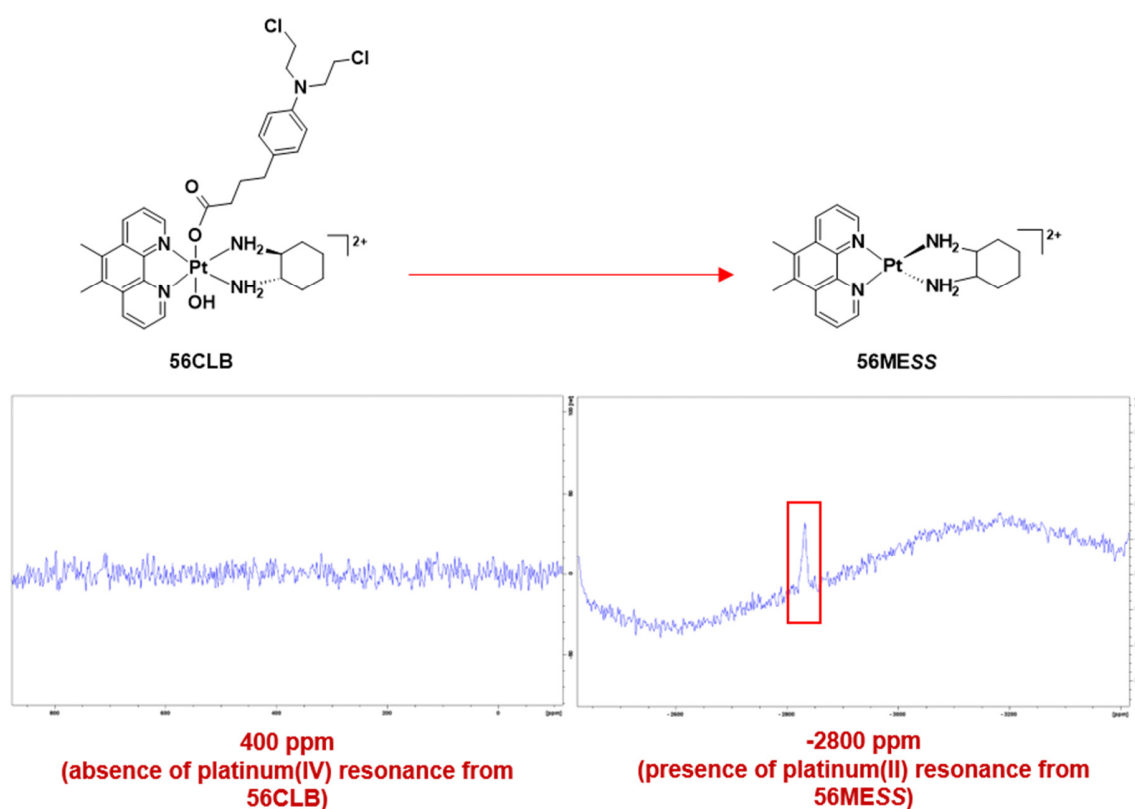


Figure S39: $1D-^{195}\text{Pt}$ NMR spectra of **56CLB** with PBS and AsA in D_2O at 310.15 K, within the regions of 400 and -2800 ppm, highlighting the complete reduction of the complex after 1 h. Above: Structures of **56CLB** and its platinum(II) scaffold, **56MESS**.

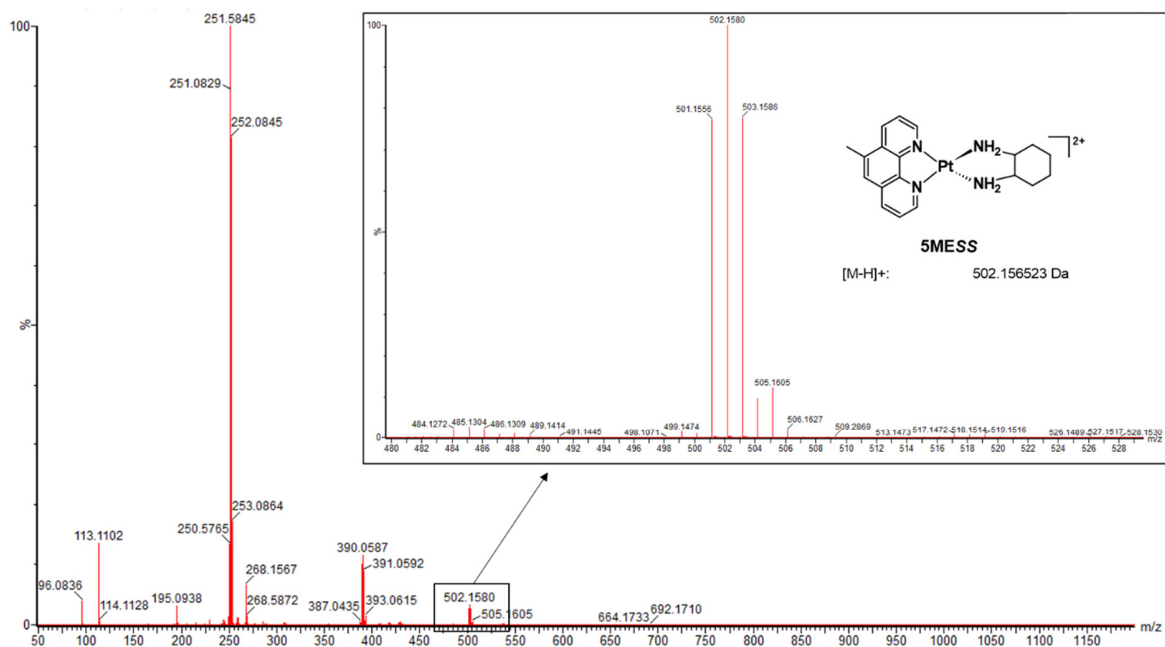


Figure S40: ESI-MS spectra of the reduced **5CLB** prodrug, highlighting the formation of the corresponding platinum(II) congener, **5MESS**.

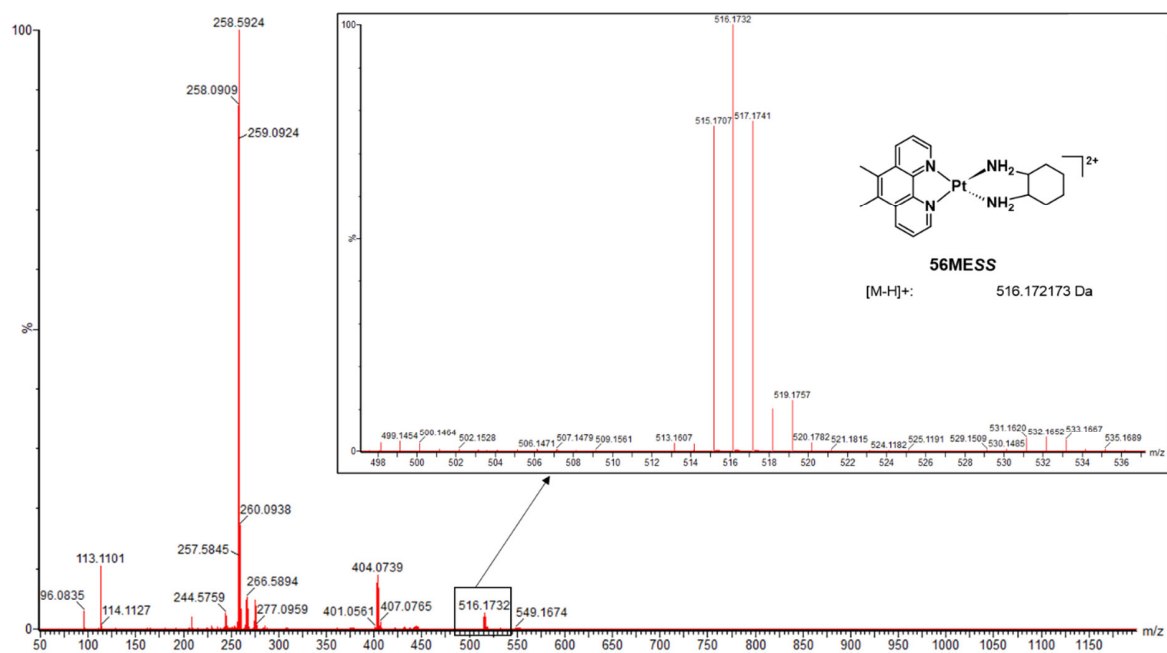


Figure S41: ESI-MS spectra of the reduced **56CLB** prodrug, highlighting the formation of the corresponding platinum(II) congener, **56MESS**.

ROS experiments

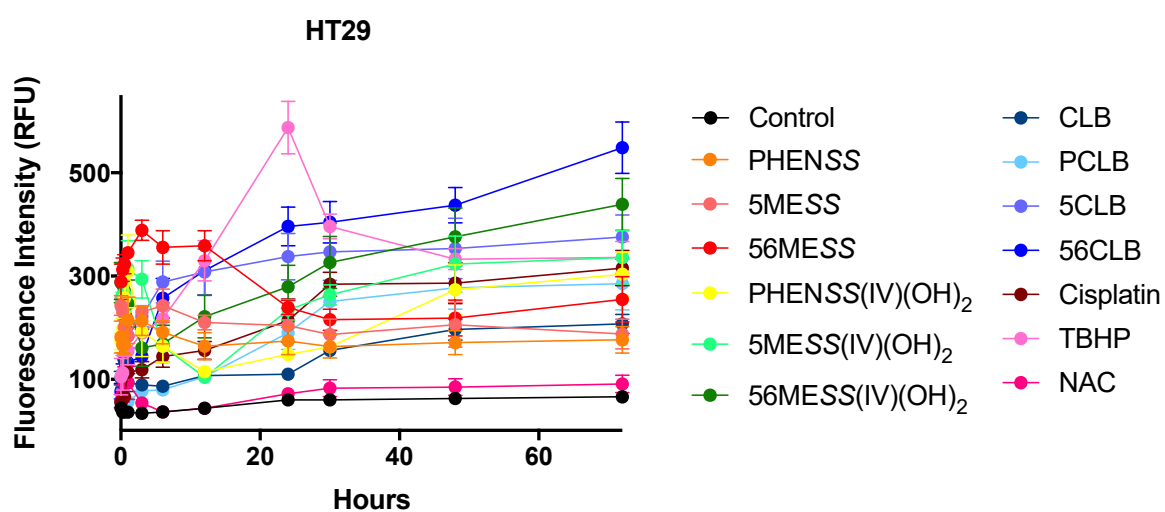


Figure S42: ROS production upon treatment with **PCLB**, **5CLB**, **56CLB**, cisplatin, together with precursor platinum(II) and (IV) scaffolds in HT29 colon cancer cell line at 0, 0.25, 0.5, 1, 3, 6, 12, 24, 48 and 72 h. TBHP: t-butyl hydroperoxide, NAC: N-acetylcysteine. Data points denote mean \pm SEM. n = 3 from three independent experiments where samples were run in triplicates.