

Supporting Information

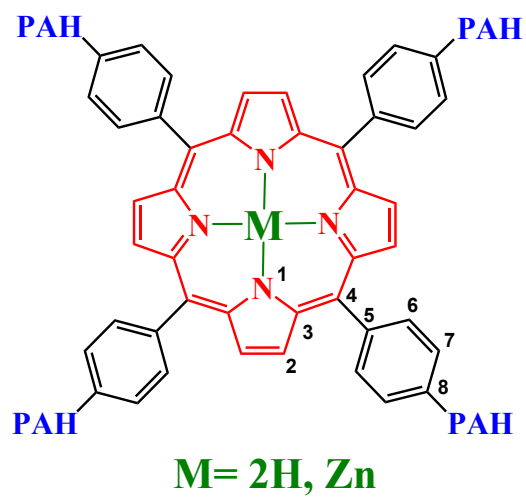
Decoupling the synergistic effects in aromatic-porphyrin-fullerene systems

Sergio Ferrero, Héctor Barbero, Daniel Miguel, Raúl García-Rodríguez* and Celedonio M. Álvarez*

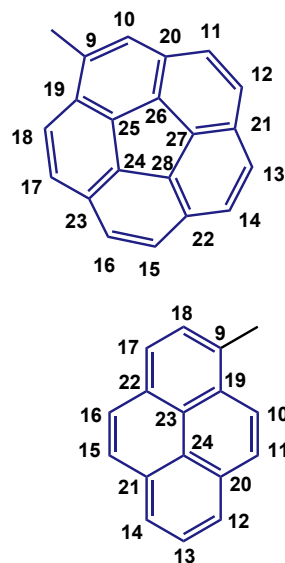
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General atom numbering



PAH:



NMR spectra

2H-PTetraBr

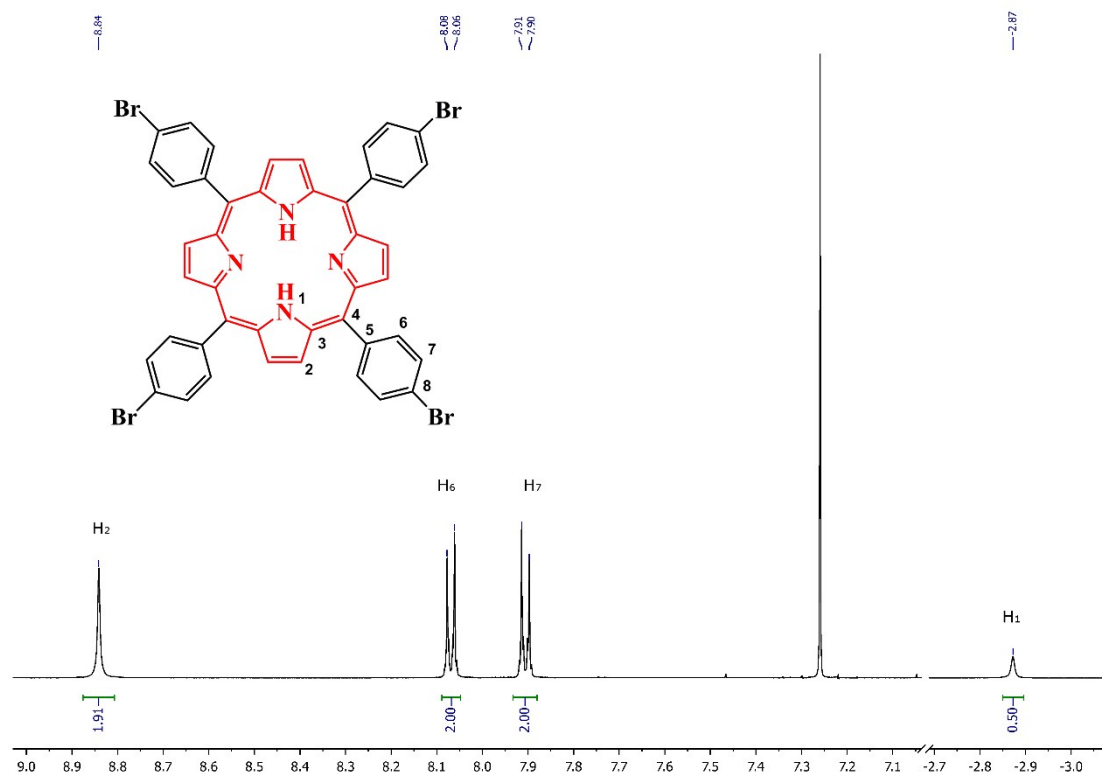


Figure S1: ^1H NMR (500 MHz, CDCl_3) spectrum of 2H-PTetraBr .

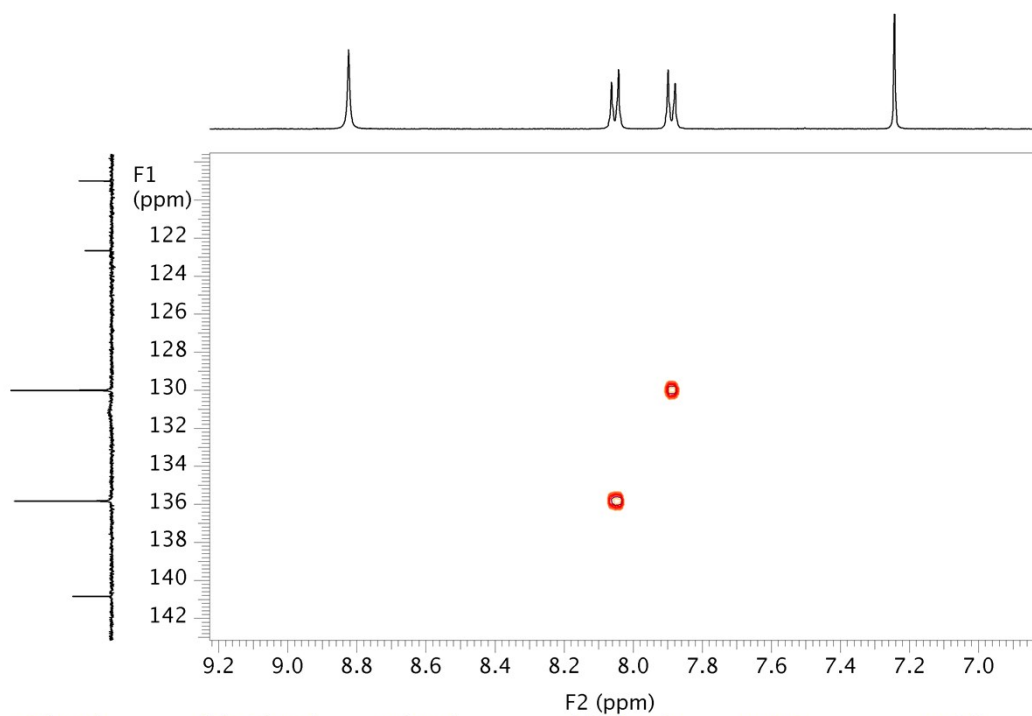


Figure S2: ^1H - ^{13}C band selective HSQC (CDCl_3) of 2H-PTetraBr .

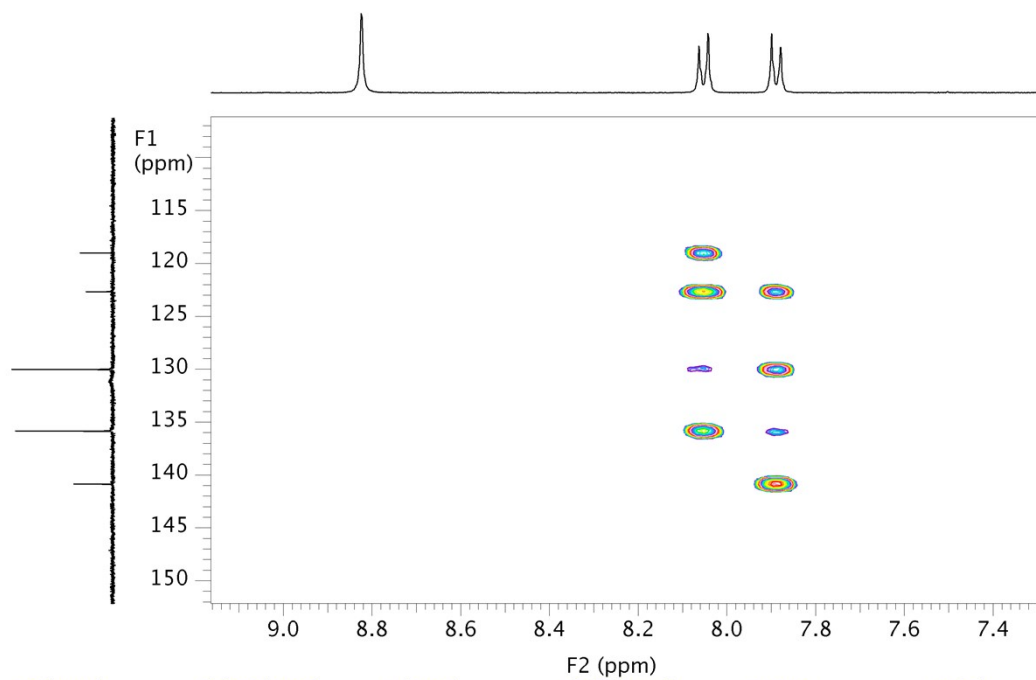


Figure S3: ^1H - ^{13}C band selective HMBC (CDCl_3) of **2H-PTetraBr**.

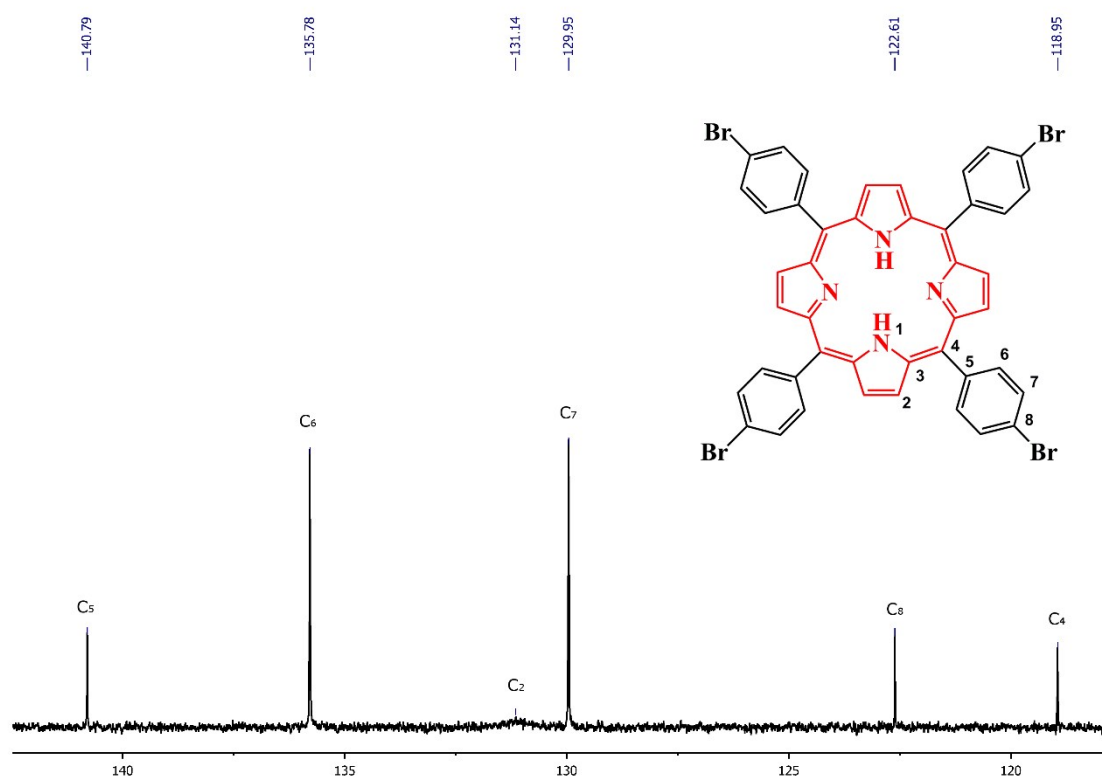


Figure S4: ^{13}C $\{^1\text{H}\}$ NMR (101 MHz, CDCl_3) spectrum of **2H-PTetraBr**.

Zn-PTetraBr

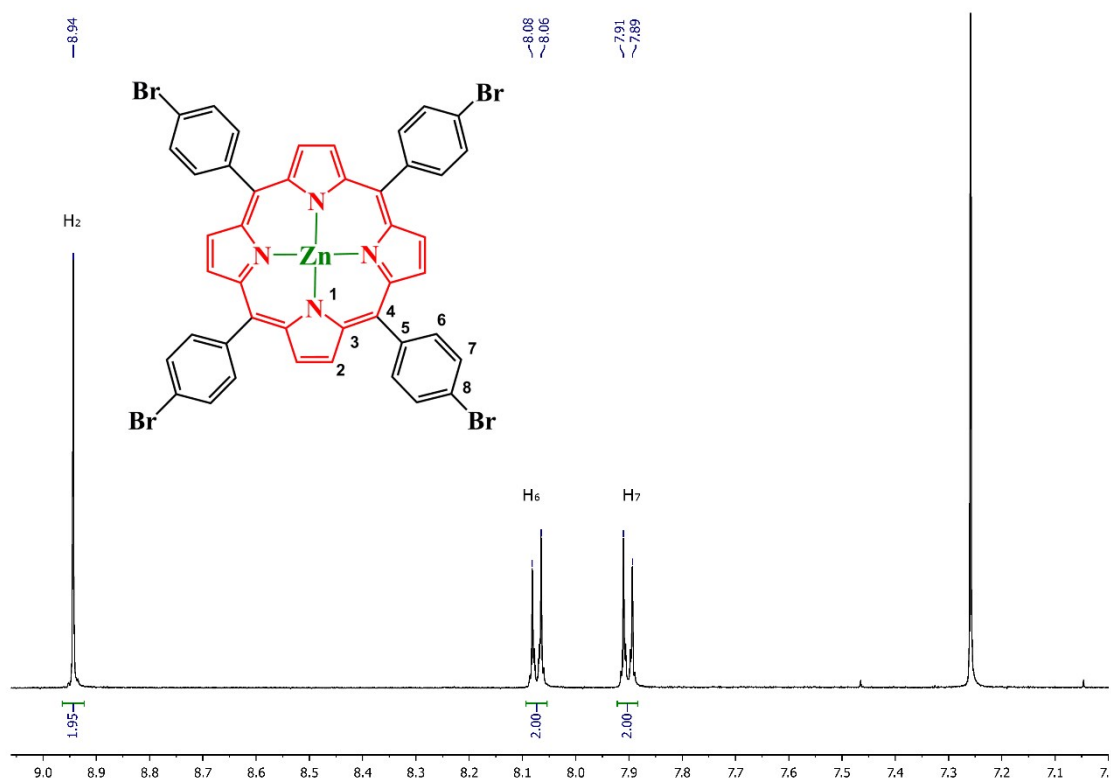


Figure S5: ^1H NMR (500 MHz, CDCl_3) spectrum of Zn-PTetraBr.

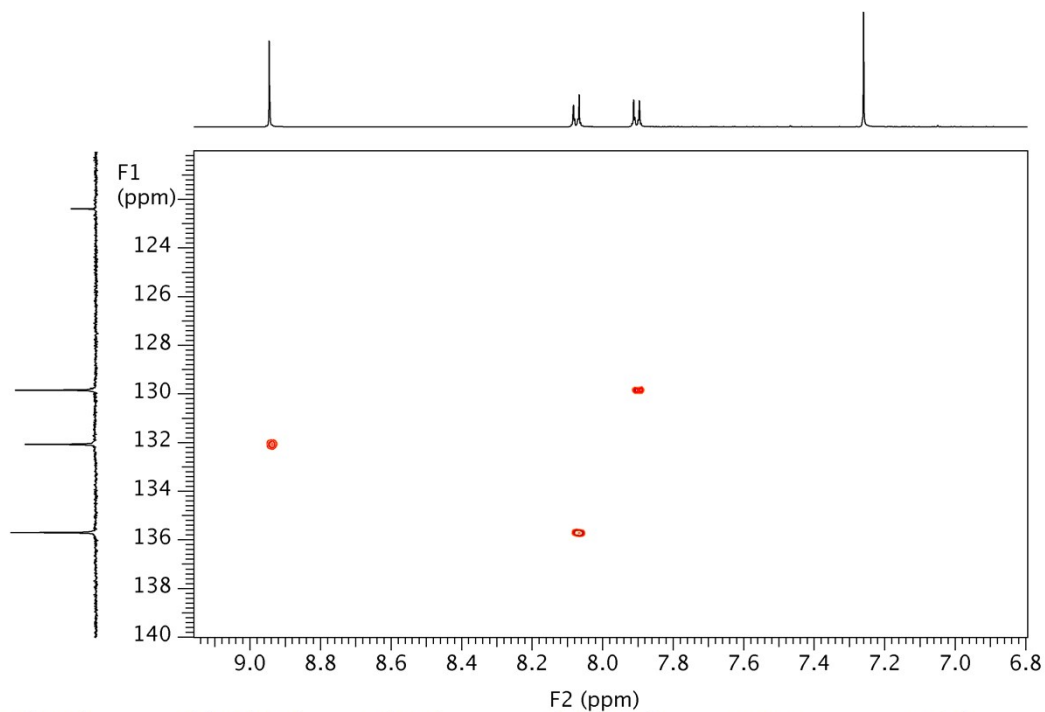


Figure S6: ^1H - ^{13}C band selective HSQC (CDCl_3) of Zn-PTetraBr.

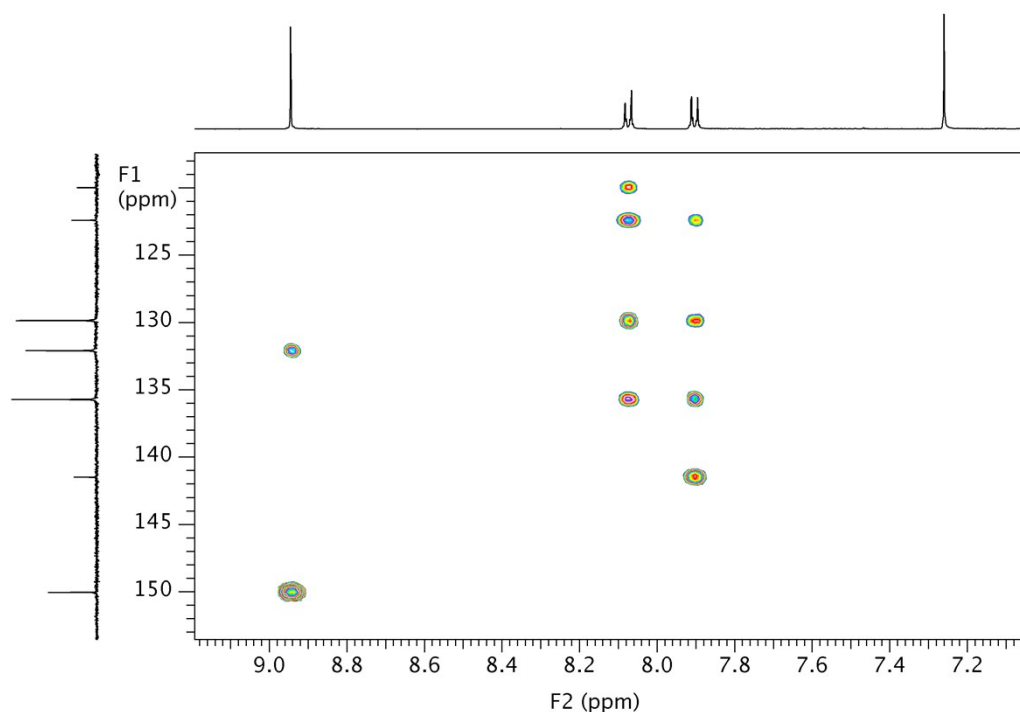


Figure S7: ^1H - ^{13}C band selective HMBC (CDCl_3) of **Zn-PTetraBr**.

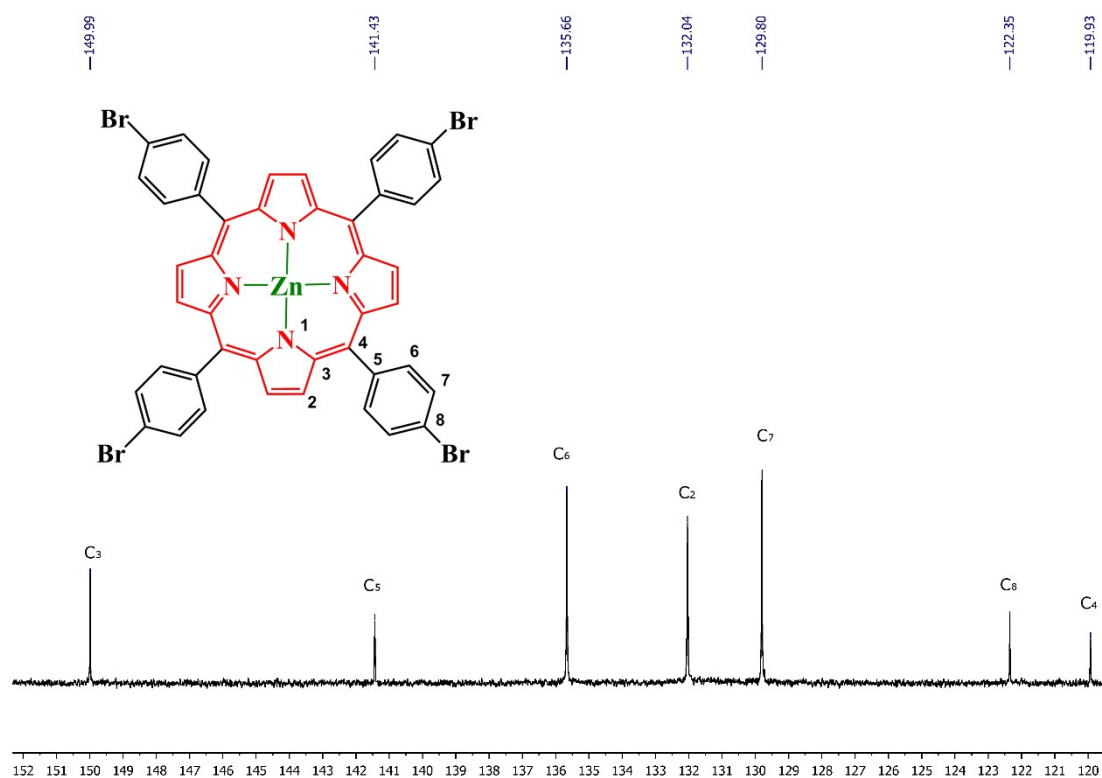


Figure S8: ^{13}C $\{^1\text{H}\}$ NMR (101 MHz, CDCl_3) spectrum of **Zn-PTetraBr**.

2H-PTetraBpin

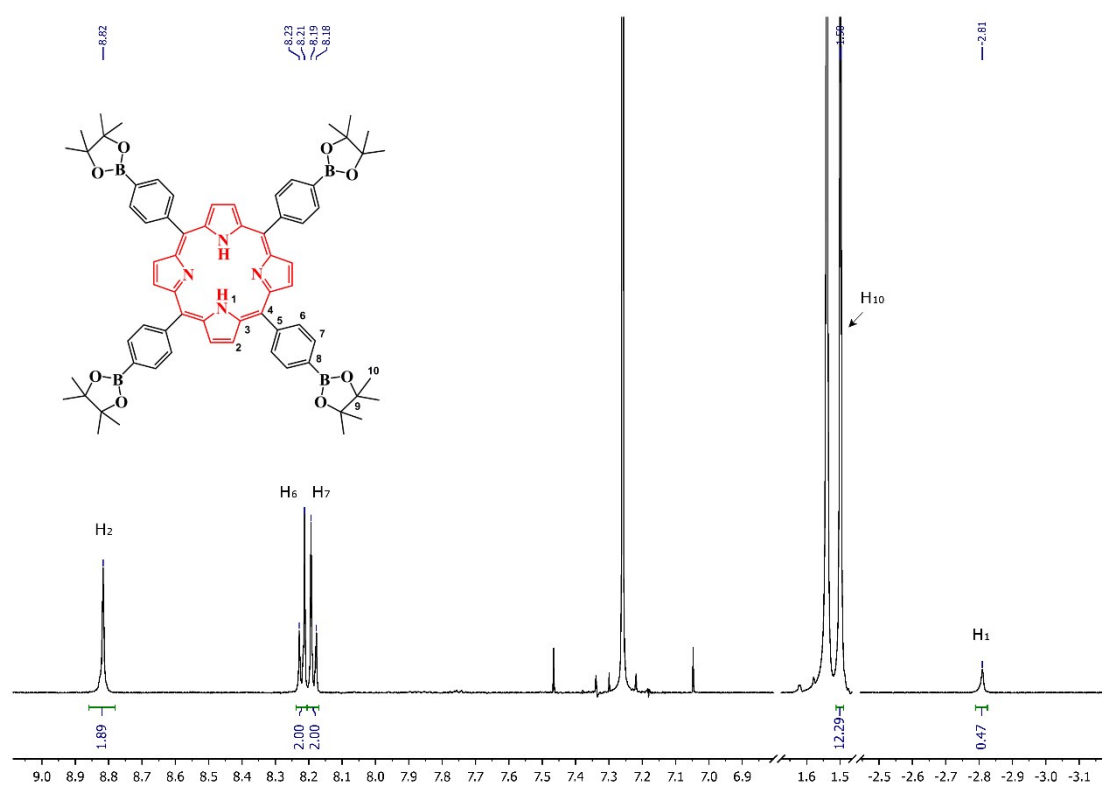


Figure S9: ¹H NMR (500 MHz, CDCl₃) spectrum of 2H-PTetraBpin.

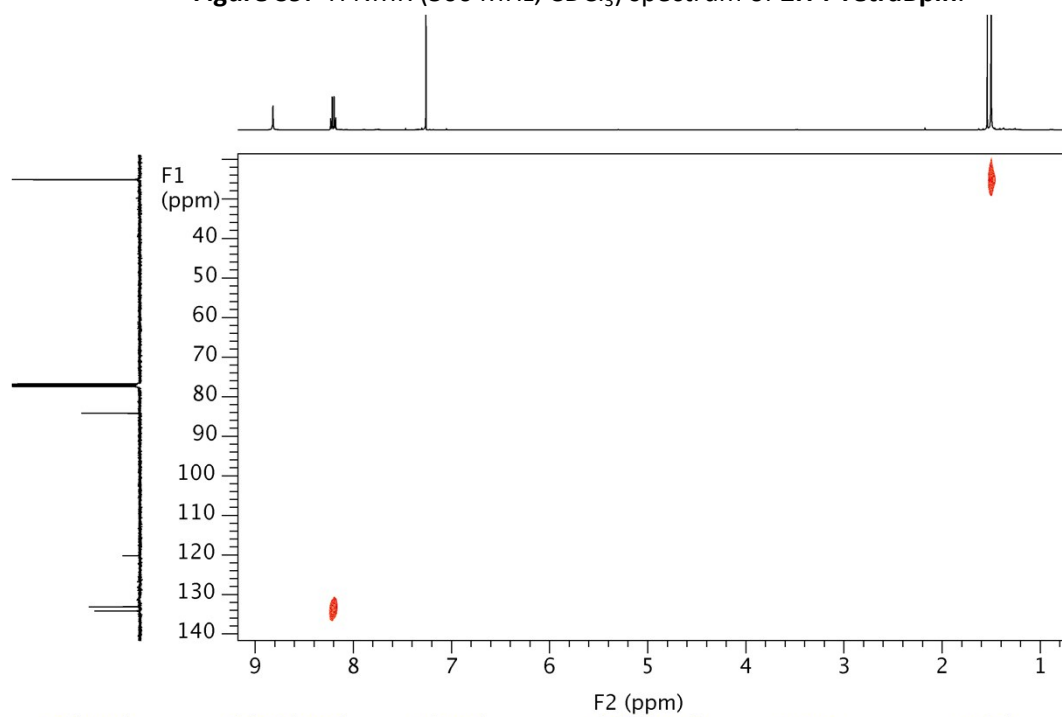


Figure S10: Full ¹H-¹³C HSQC (CDCl₃) of 2H-PTetraBpin.

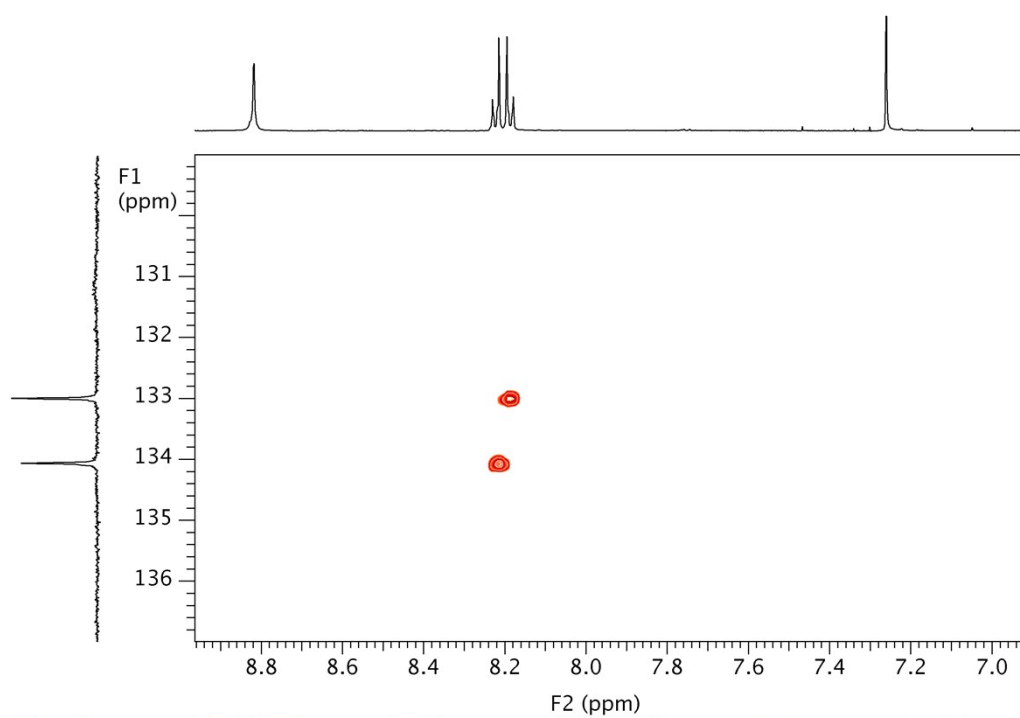


Figure S11: ^1H - ^{13}C band selective HSQC (expansion of the aromatic region) (CDCl_3) of **2H-PTetraBpin**.

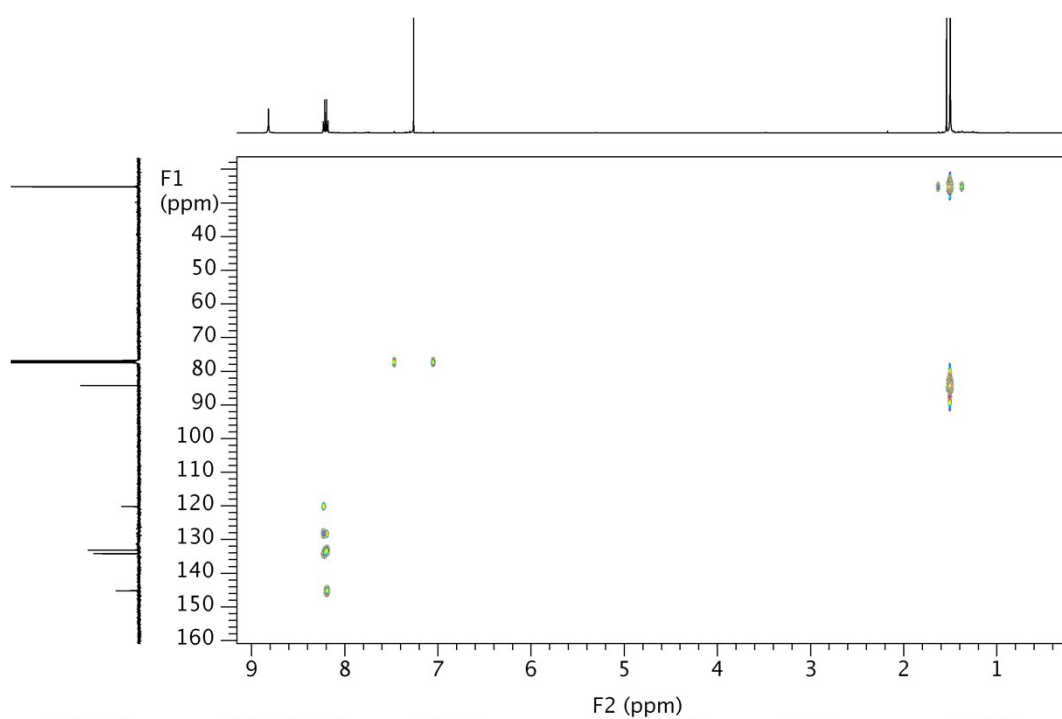


Figure S12: Full ^1H - ^{13}C HMBC (CDCl_3) of **2H-PTetraBpin**.

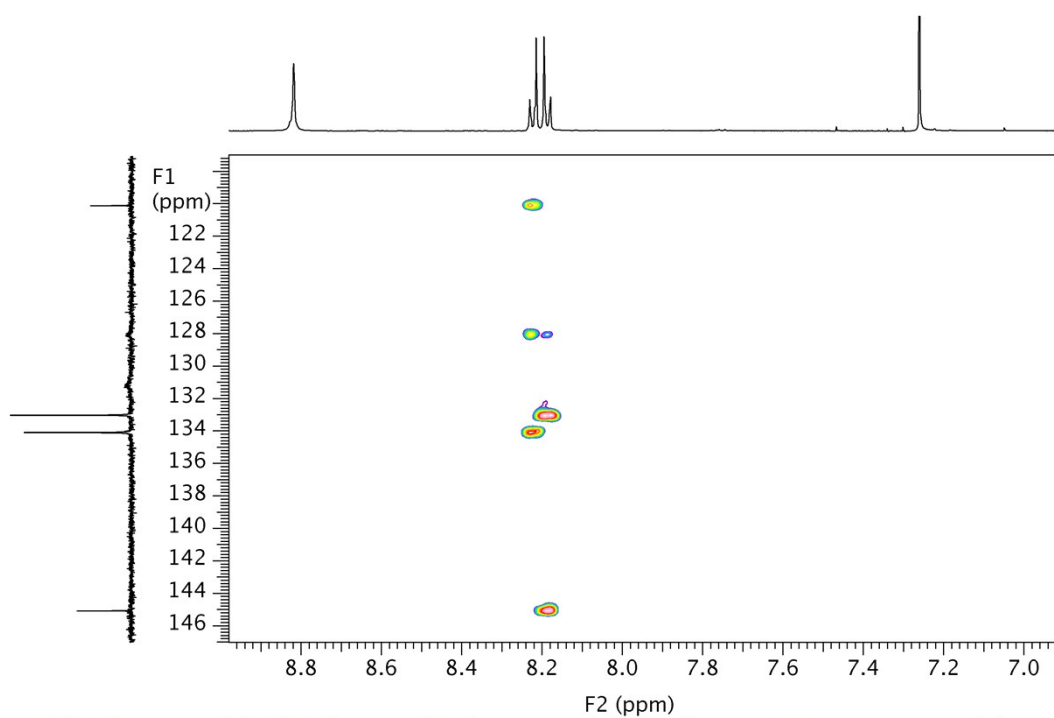


Figure S13: ^1H - ^{13}C band selective HMBC (expansion of the aromatic region) (CDCl_3) of **2H-PTetraBpin**.

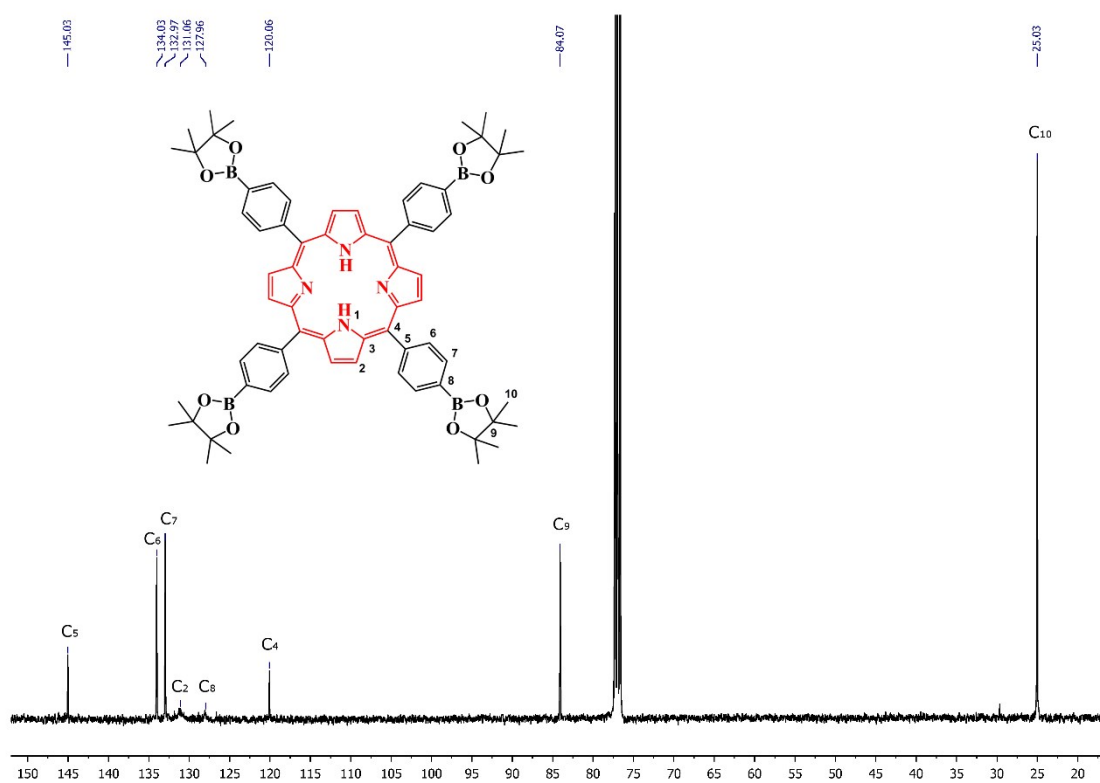


Figure S14: ^{13}C $\{^1\text{H}\}$ NMR (101 MHz, CDCl_3) spectrum of **2H-PTetraBpin**.

Zn-PTetraBpin

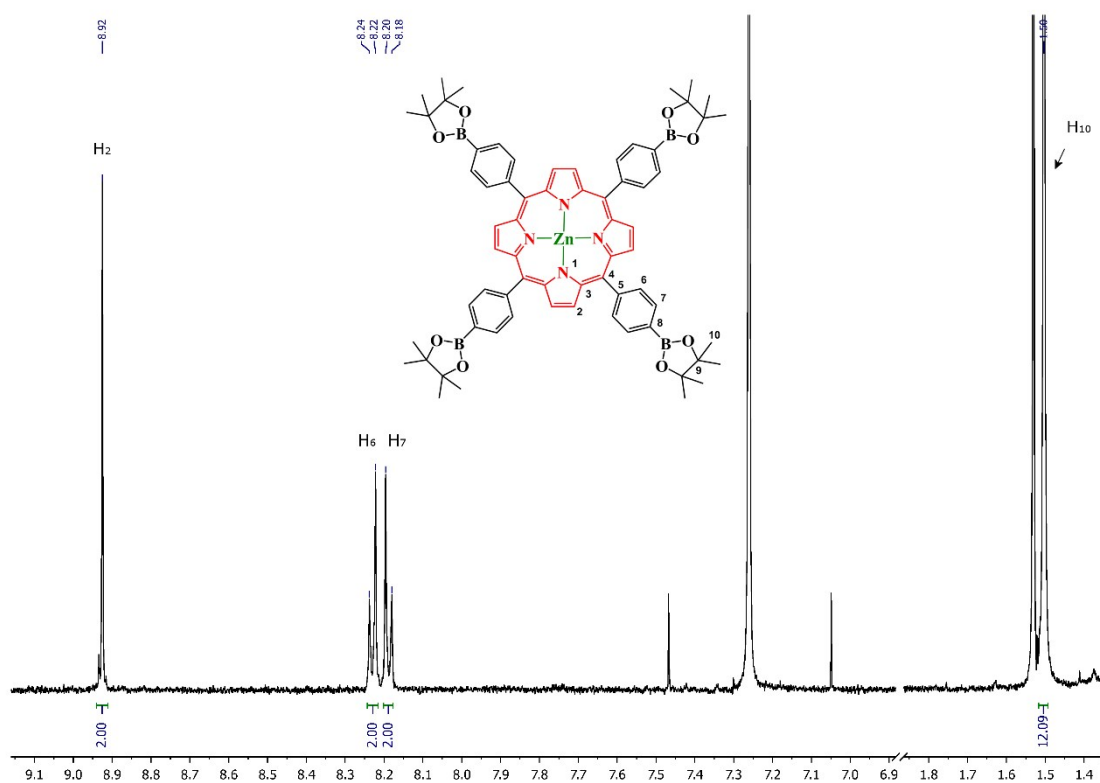


Figure S15: ¹H NMR (500 MHz, CDCl₃) spectrum of Zn-PTetraBpin.

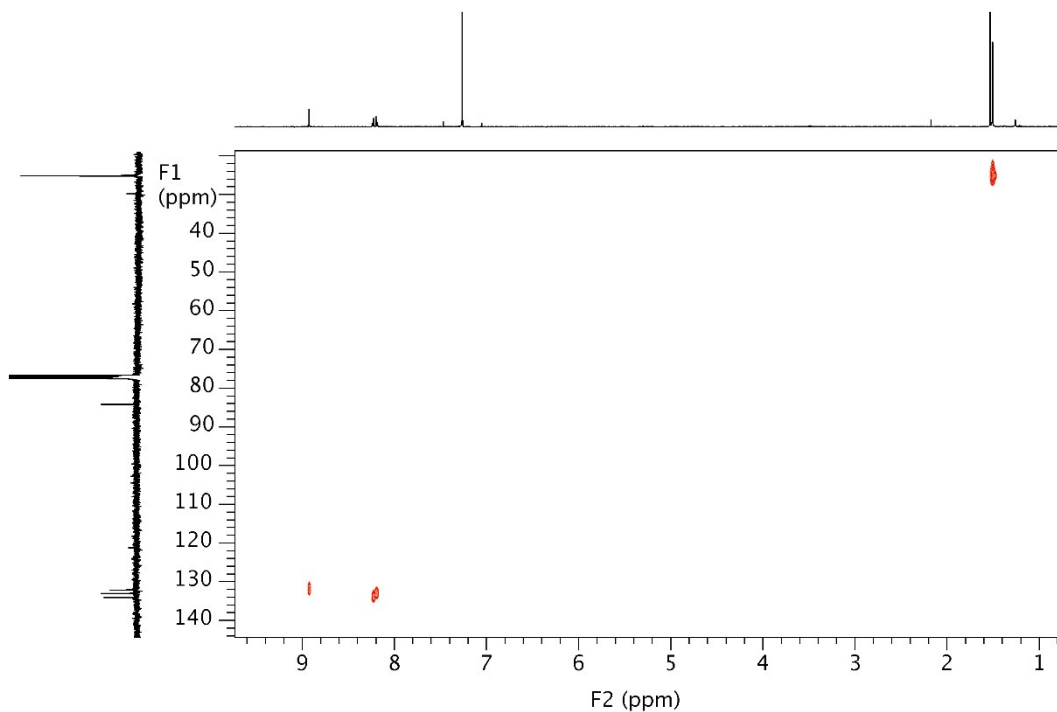


Figure S16: Full ¹H-¹³C HSQC (CDCl₃) of Zn-PTetraBpin.

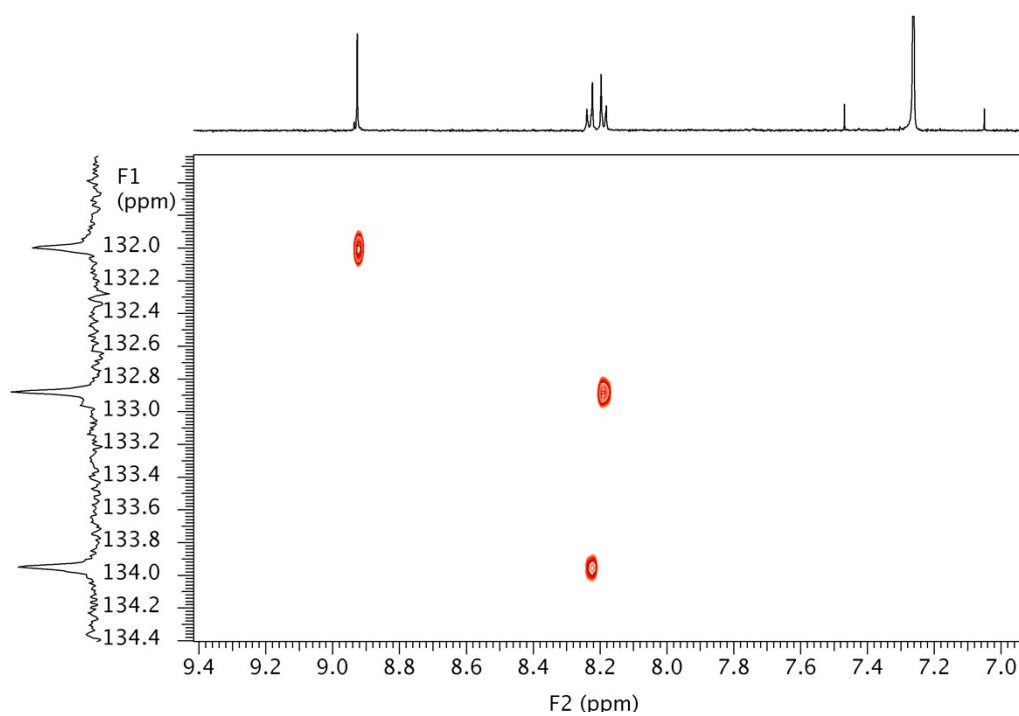


Figure S17: ^1H - ^{13}C band selective HSQC (expansion of the aromatic region) (CDCl_3) of **Zn-PTetraBpin**.

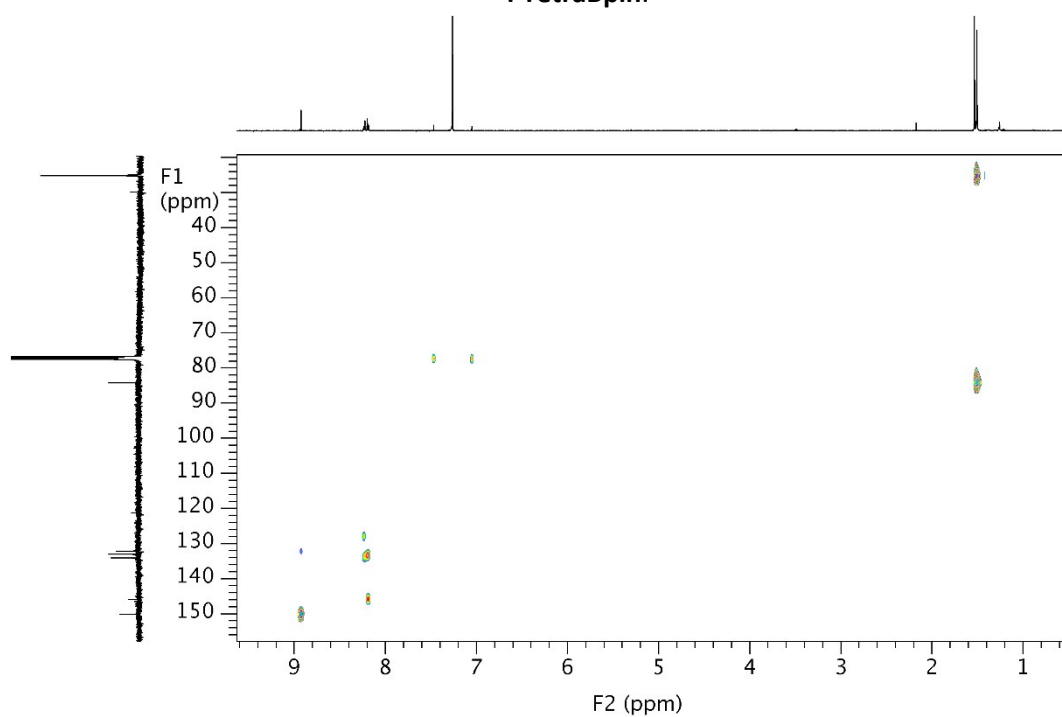


Figure S18: ^1H - ^{13}C full HMBC (CDCl_3) of **Zn-PTetraBpin**.

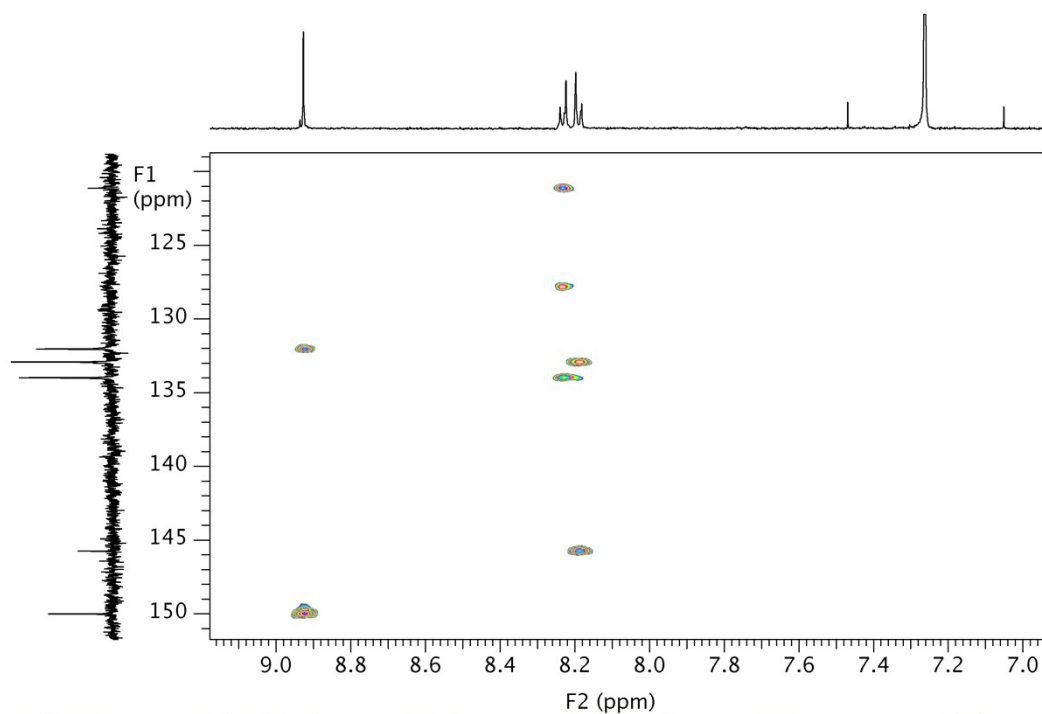


Figure S19: ^1H - ^{13}C band selective HMBC (expansion of the aromatic region) (CDCl_3) of **Zn-PTetraBpin**.

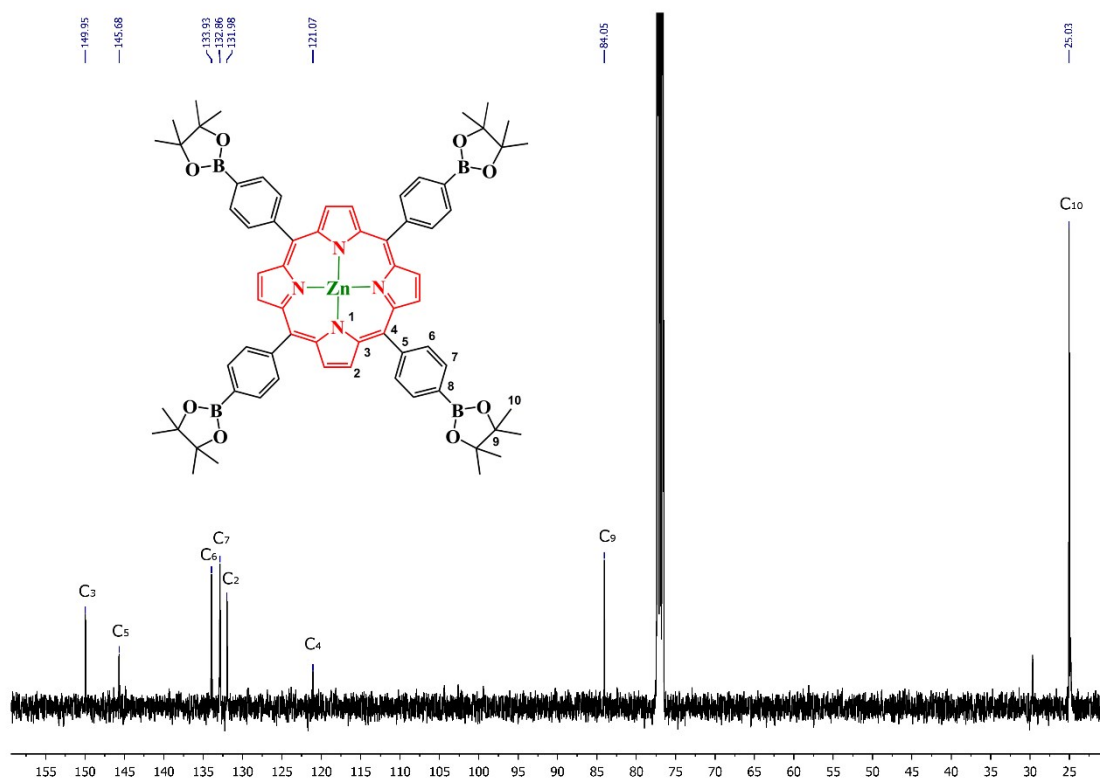


Figure S20: ^{13}C $\{^1\text{H}\}$ NMR (101 MHz, CDCl_3) spectrum of **Zn-PTetraBpin**.

Zn-PTetraPyr

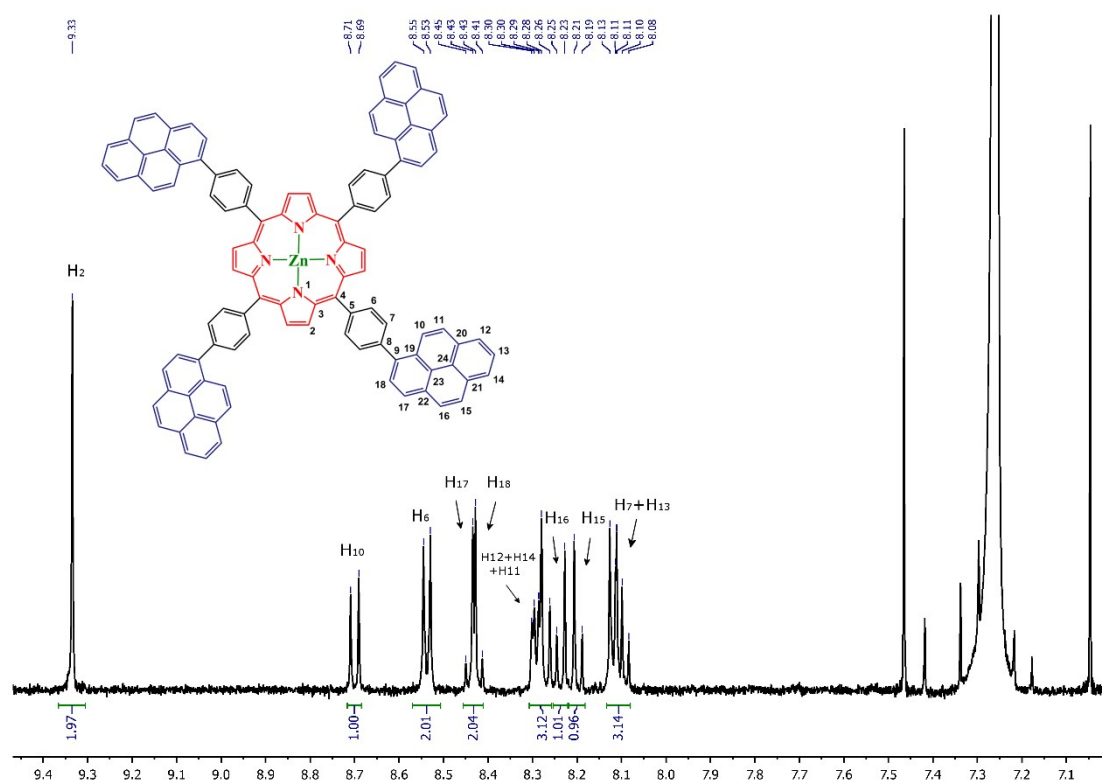


Figure S21: ^1H NMR (500 MHz, CDCl_3) spectrum of Zn-PTetraPyr.

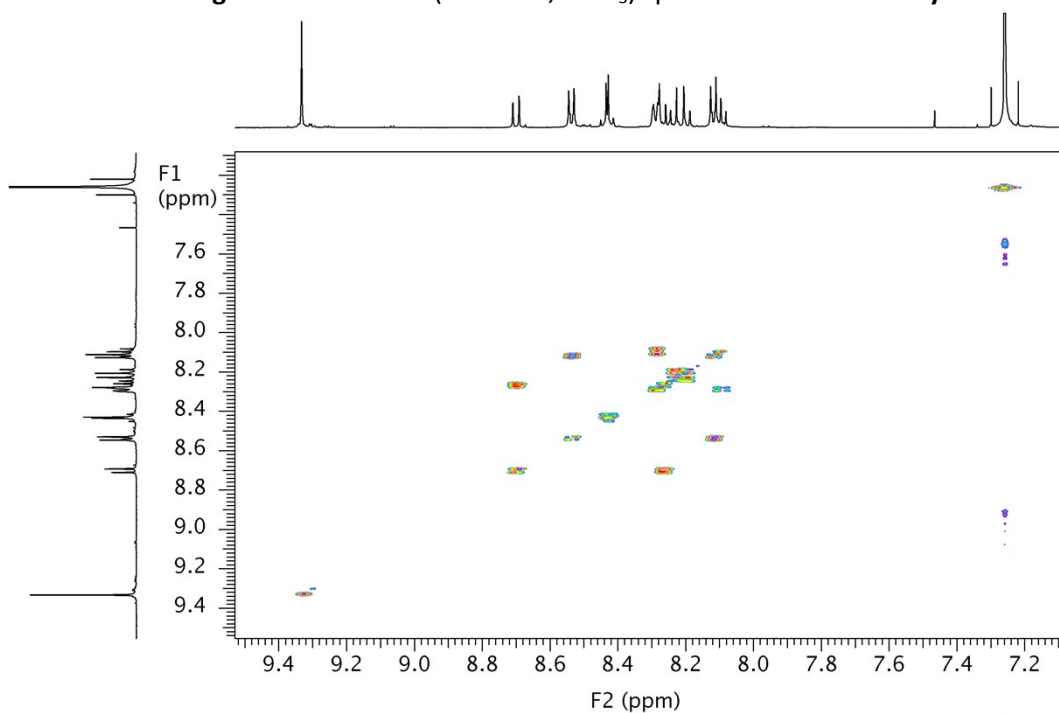


Figure S22: ^1H - ^1H COSY (CDCl_3) of Zn-PTetraPyr.

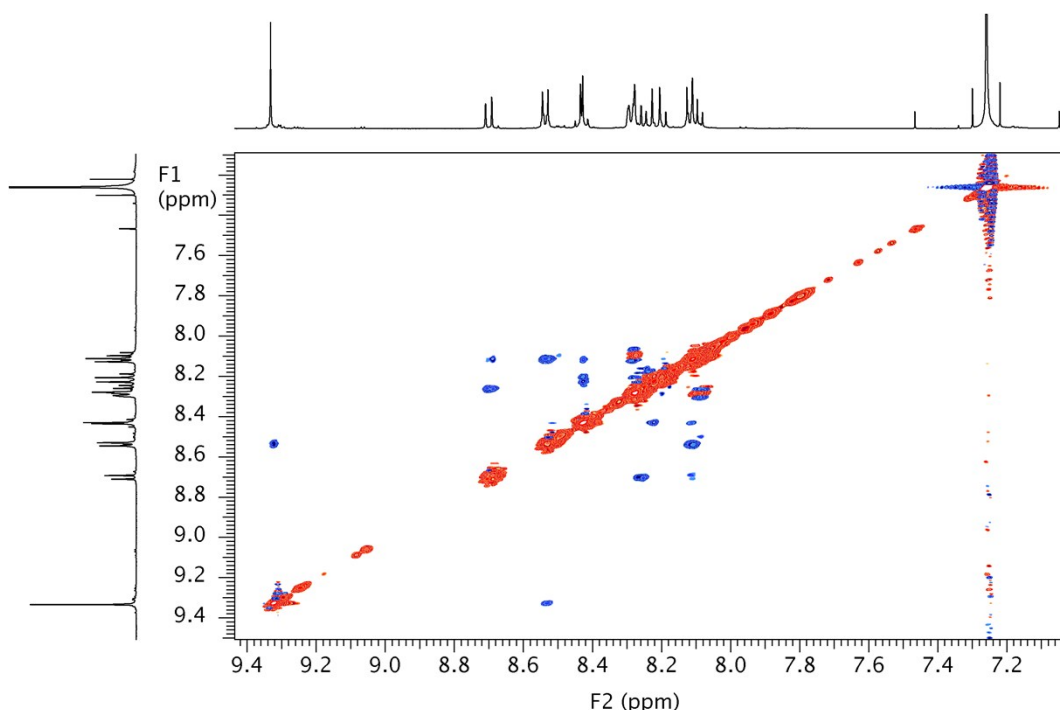


Figure S23: ^1H - ^1H band selective ROESY (CDCl_3) of Zn-PTetraPyr.

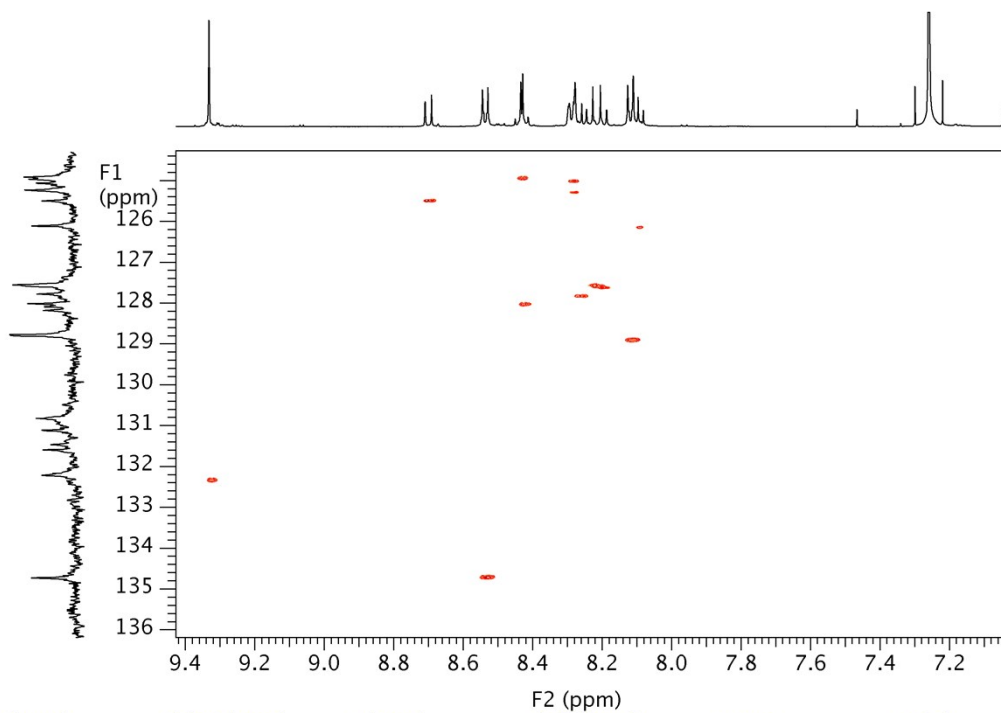


Figure S24: ^1H - ^{13}C band selective HSQC (CDCl_3) of Zn-PTetraPyr.

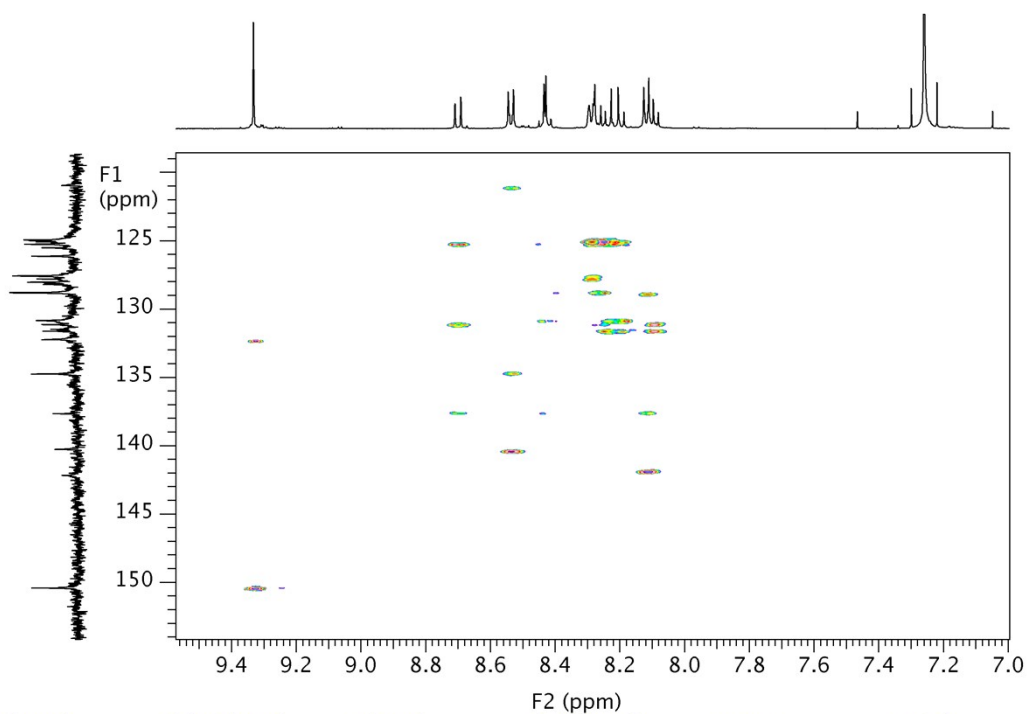


Figure S25: ^1H - ^{13}C band selective HMBC (CDCl_3) of Zn-PTetraPyr.

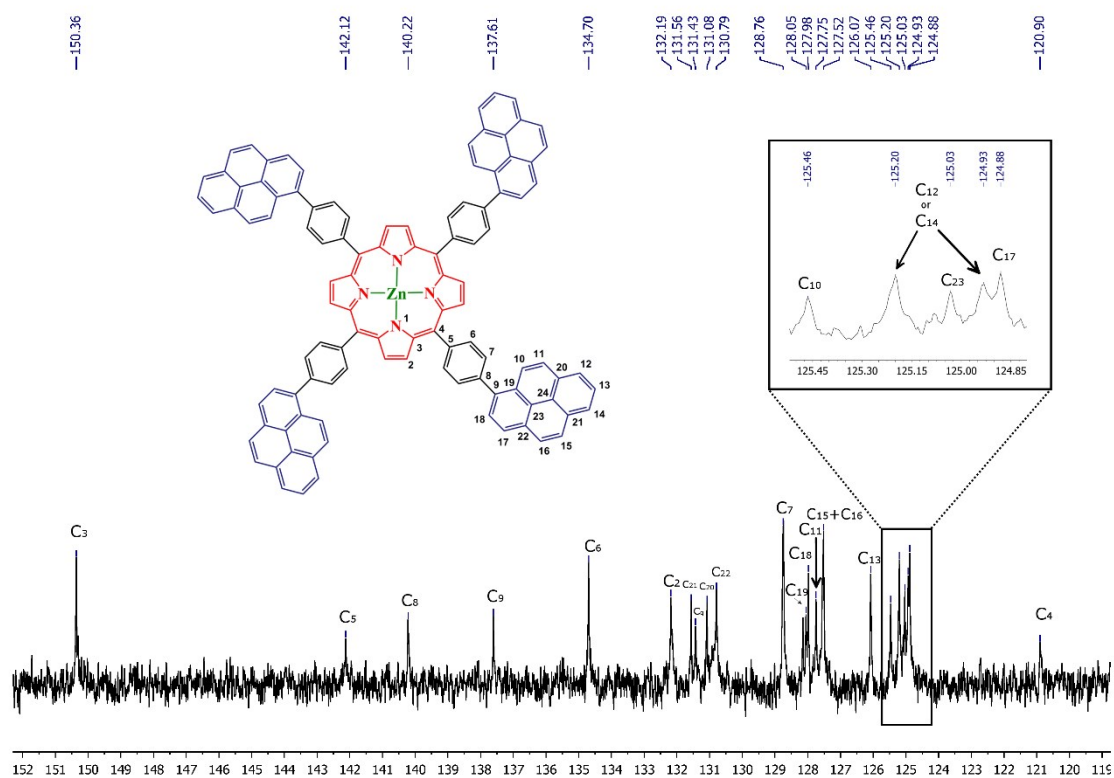


Figure S26: ^{13}C $\{^1\text{H}\}$ NMR (126 MHz, CDCl_3) spectrum of Zn-PTetraPyr.

2H-PTetraPyr

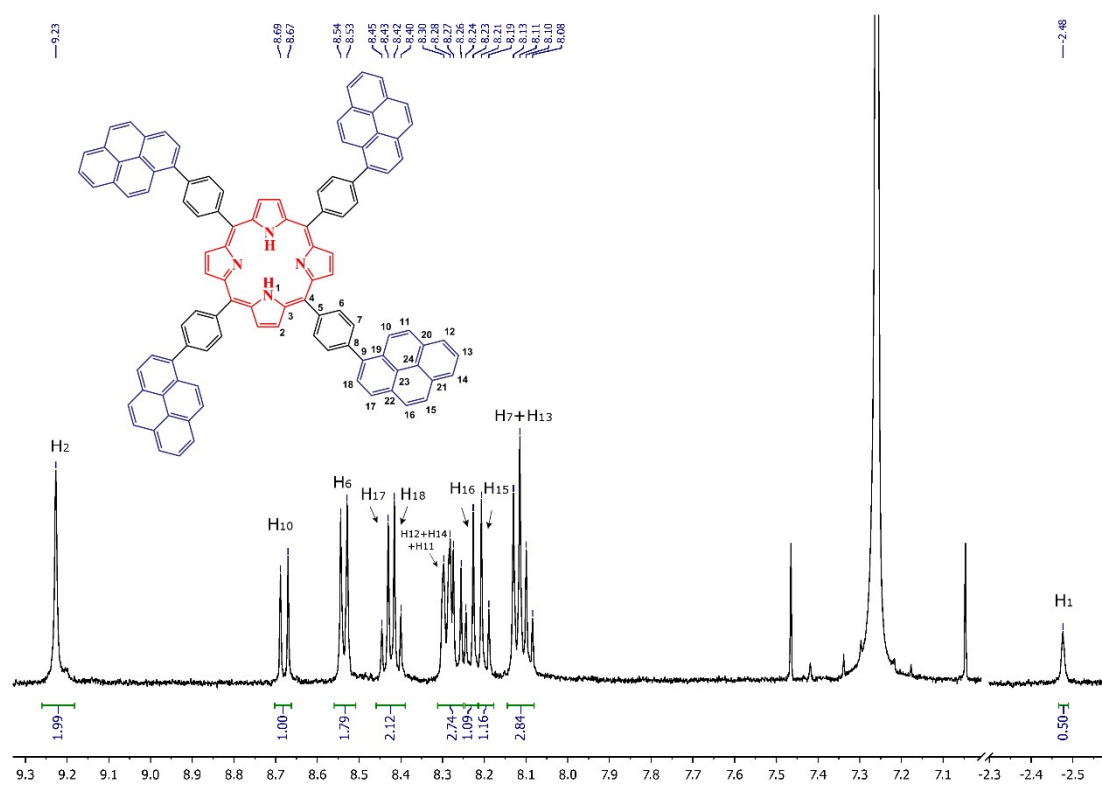


Figure S27: ¹H NMR (500 MHz, CDCl₃) spectrum of 2H-PTetraPyr.

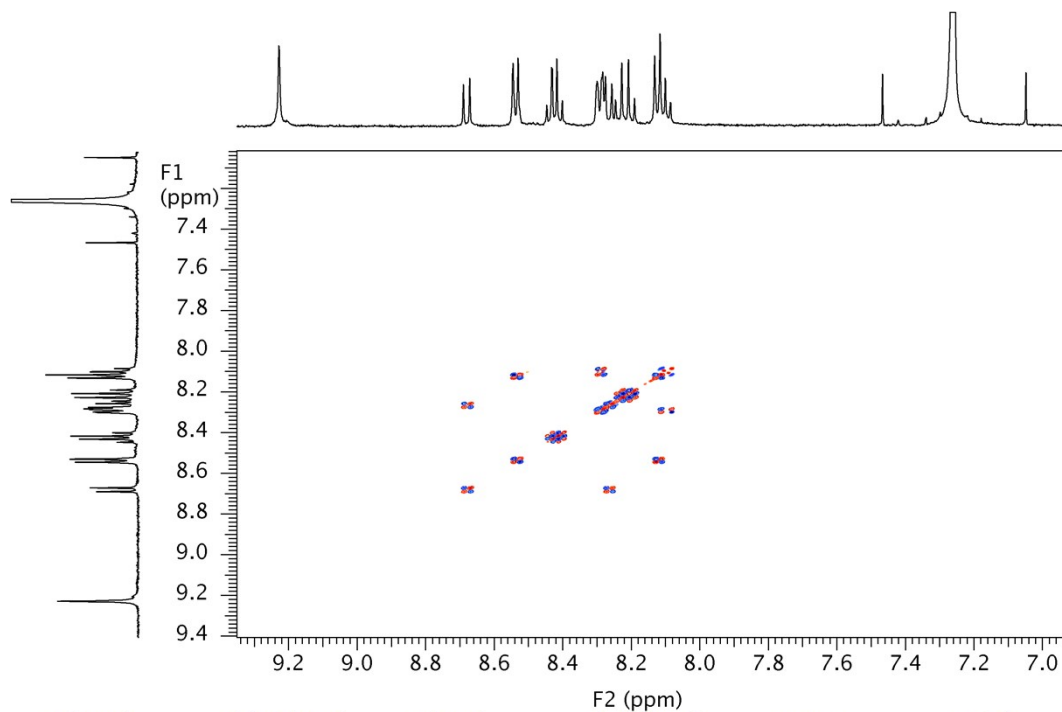


Figure S28: ¹H-¹H DQFCOSY (CDCl₃) of 2H-PTetraPyr.

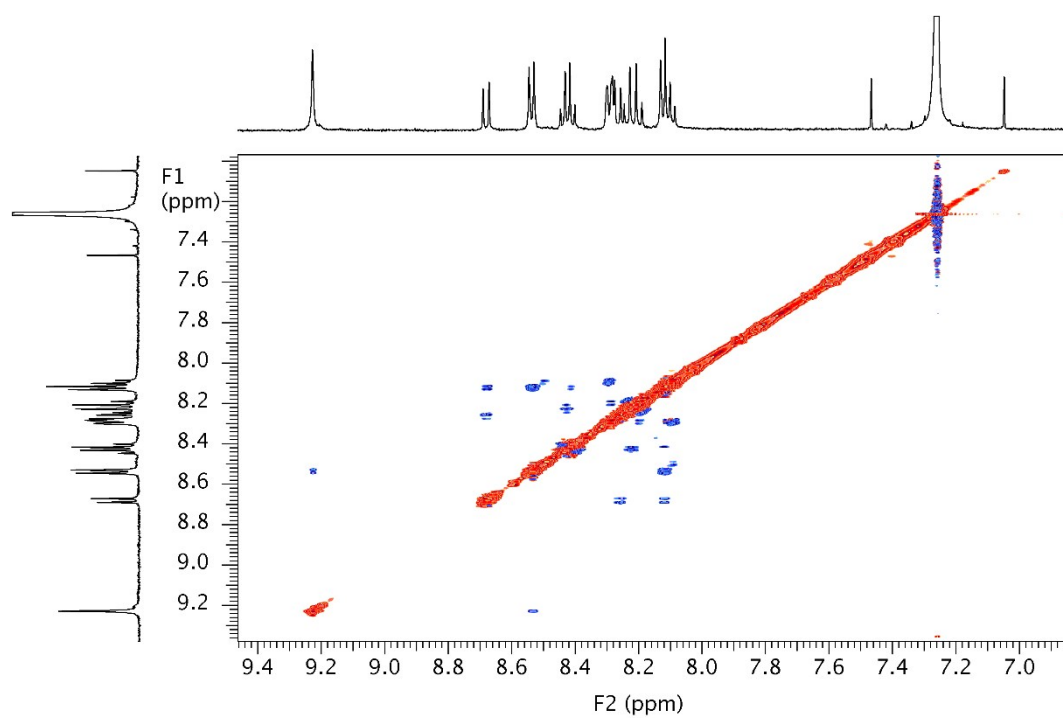


Figure S29: ^1H - ^1H band selective ROESY (CDCl_3) of **2H-PTetraPyr**.

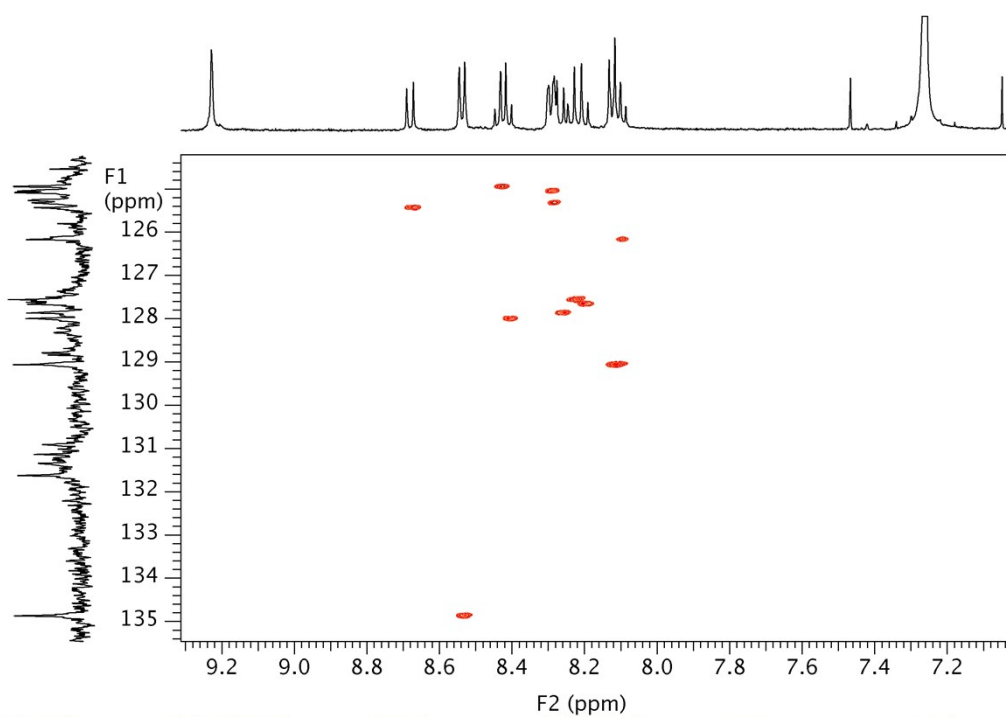


Figure S30: ^1H - ^{13}C band selective HSQC (CDCl_3) of **2H-PTetraPyr**.

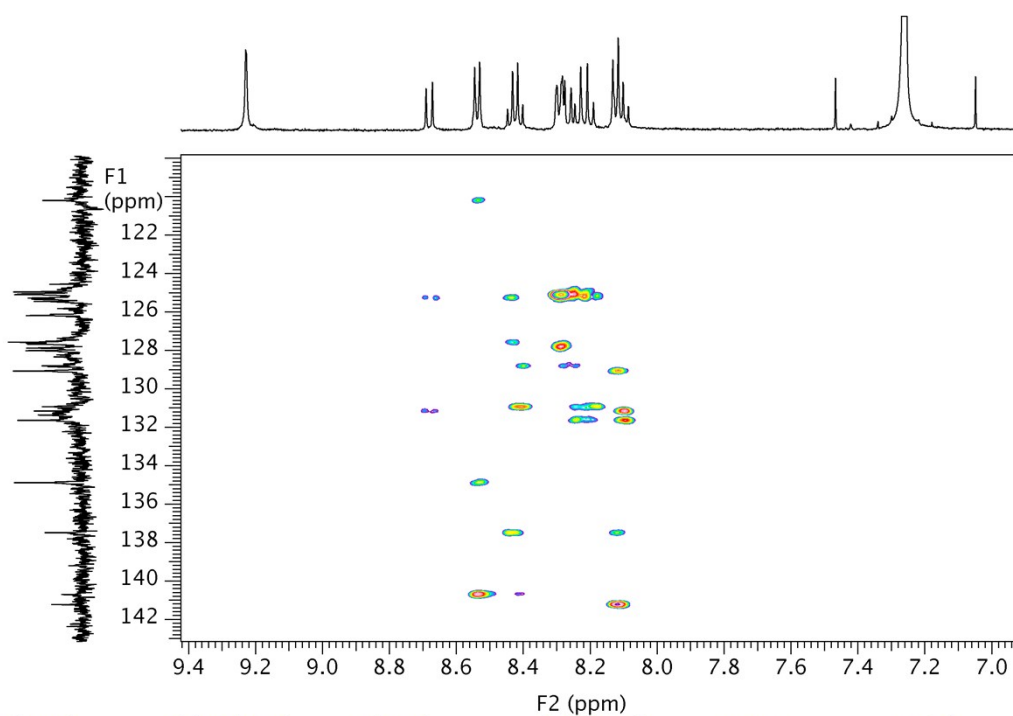


Figure S31: ^1H - ^{13}C band selective HMBC (CDCl_3) of **2H-PTetraPyr**.

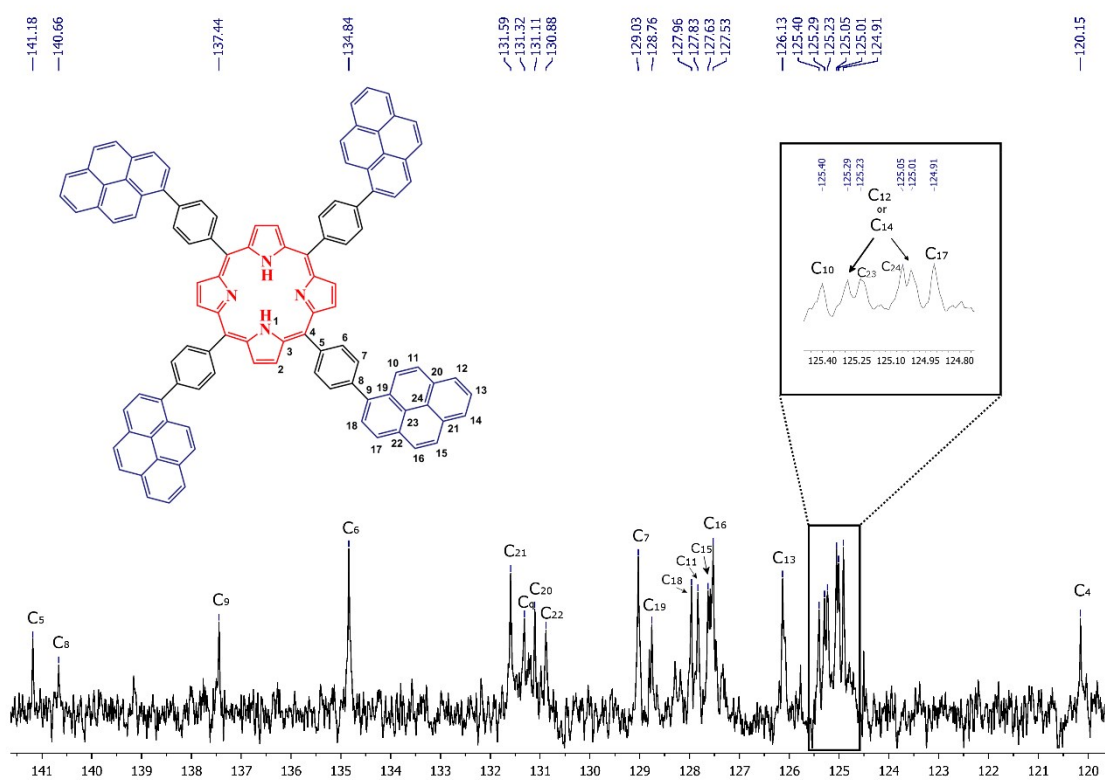
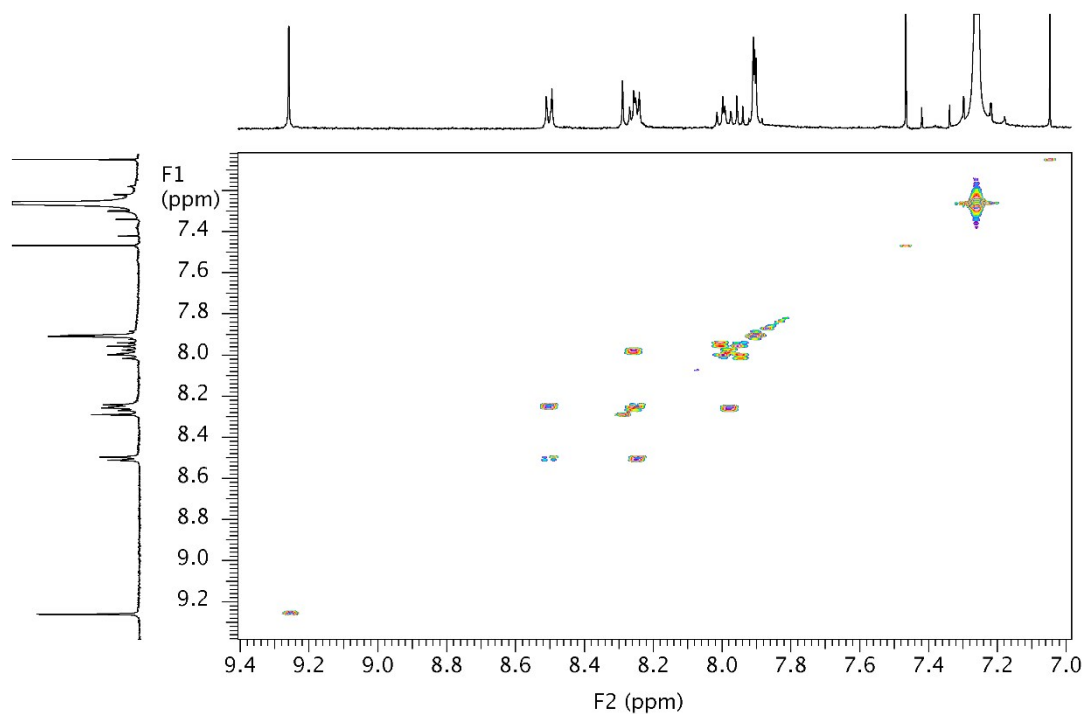
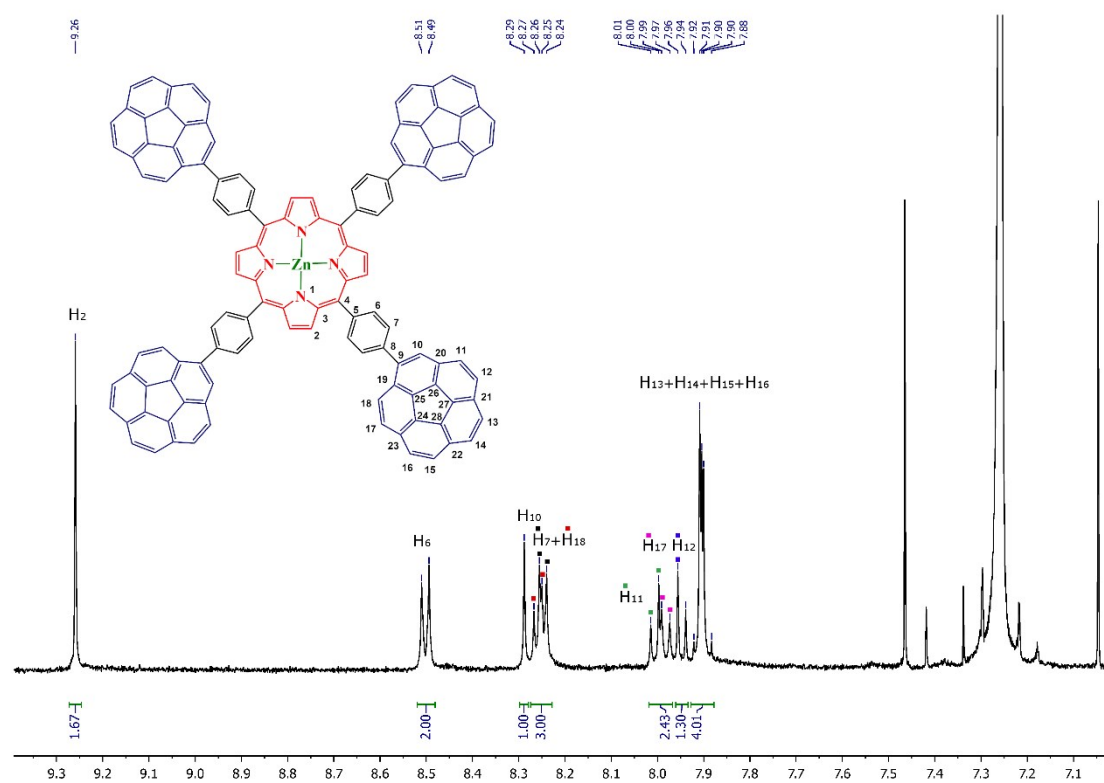


Figure S32: ^{13}C $\{^1\text{H}\}$ NMR (126 MHz, CDCl_3) spectrum of **2H-PTetraPyr**.

Zn-PTetraCor



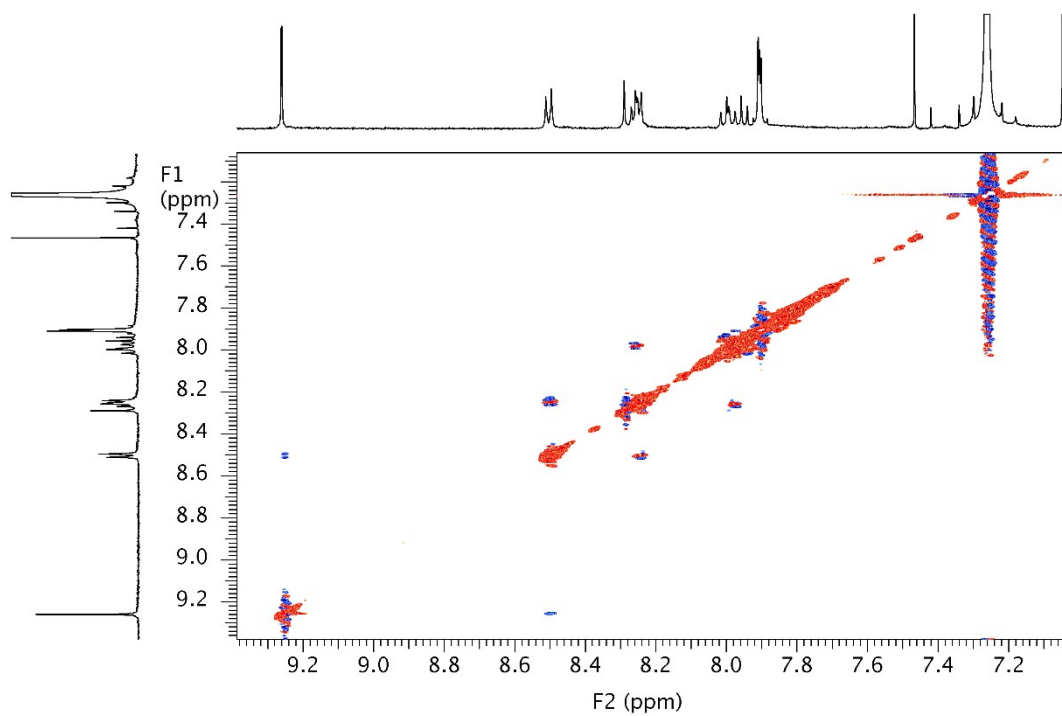


Figure S35: ^1H - ^1H band selective ROESY (CDCl_3) of **Zn-PTetraCor**.

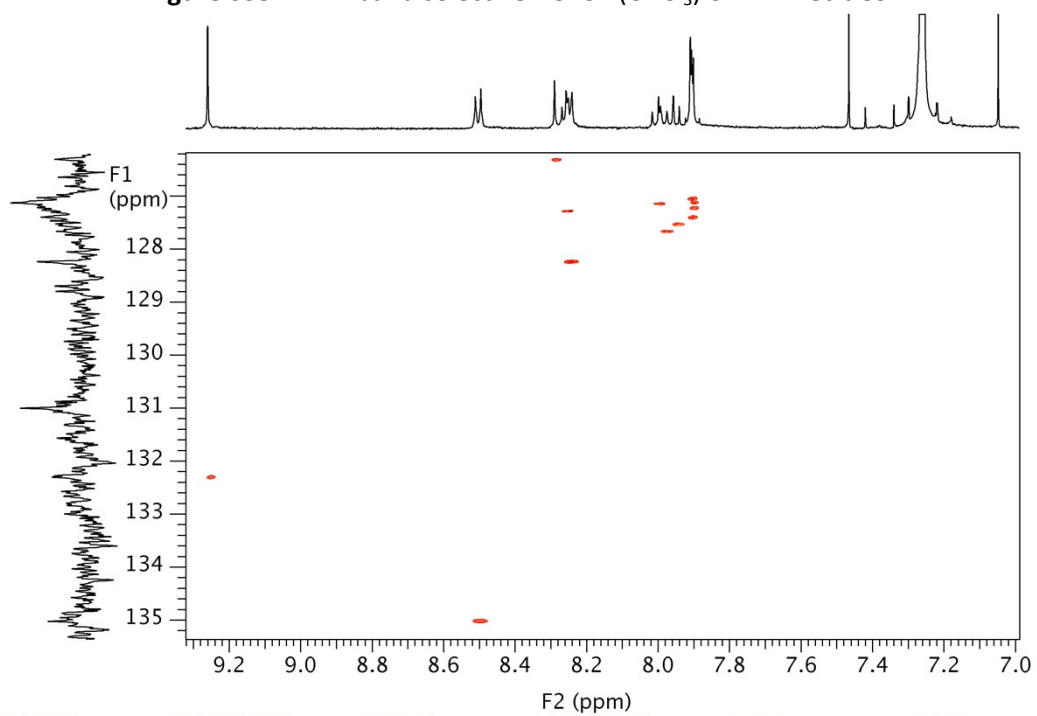


Figure S36: ^1H - ^{13}C band selective HSQC (CDCl_3) of **Zn-PTetraCor**.

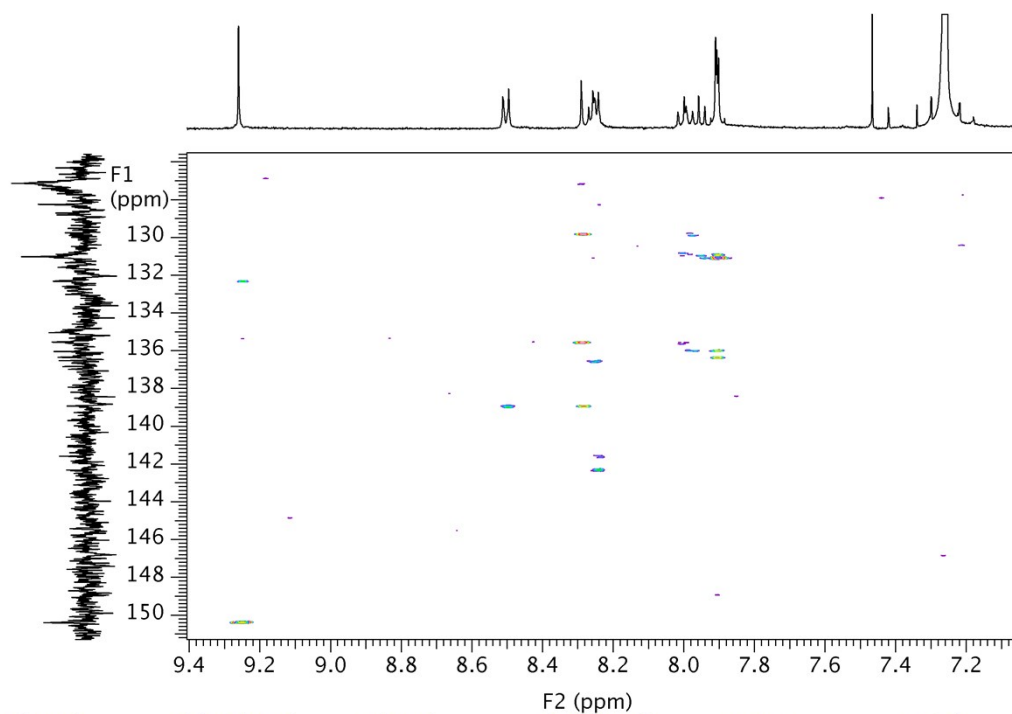


Figure S37: ^1H - ^{13}C band selective HMBC (CDCl_3) of Zn-PTetraCor.

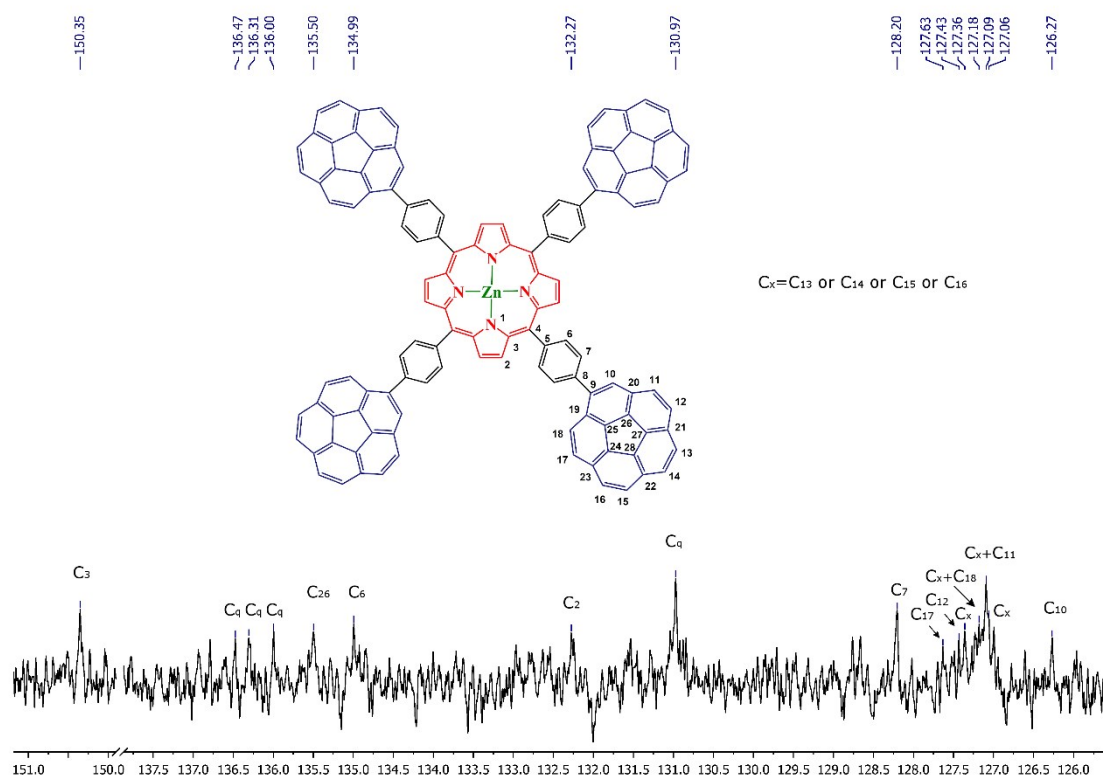


Figure S38: ^{13}C $\{^1\text{H}\}$ NMR (126 MHz, CDCl_3) spectrum of Zn-PTetraCor.

2H-PTetraCor

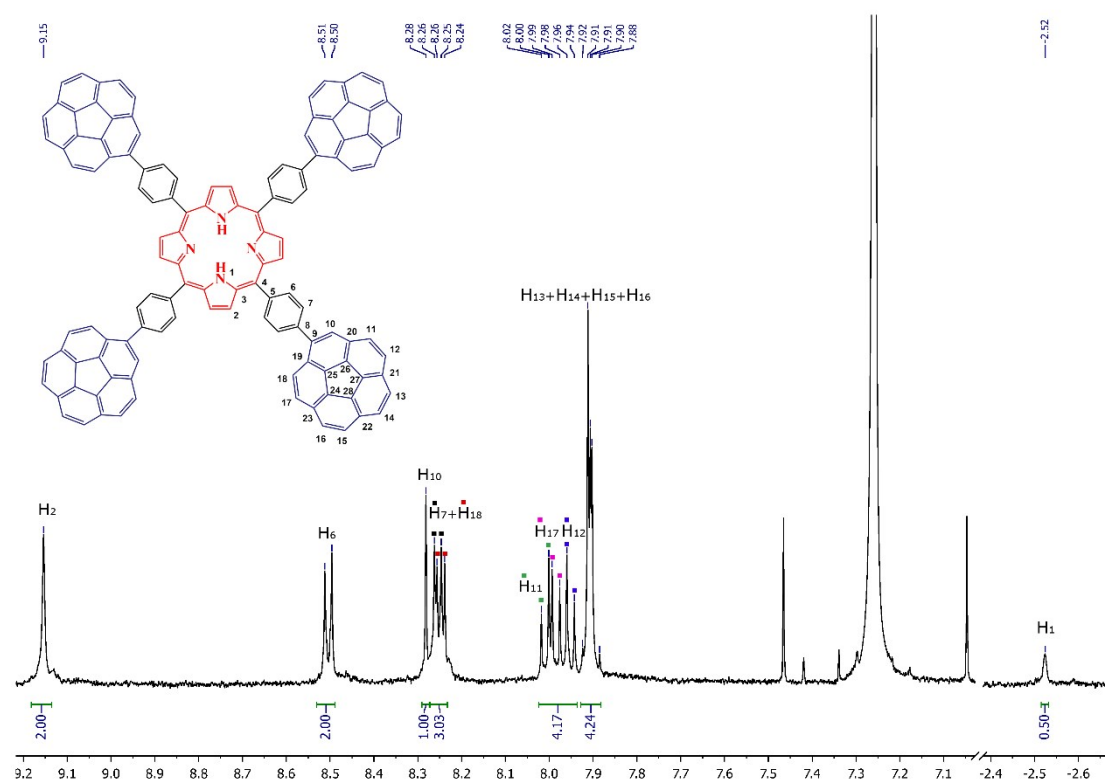


Figure S39: ¹H NMR (500 MHz, CDCl₃) spectrum of 2H-PTetraCor.

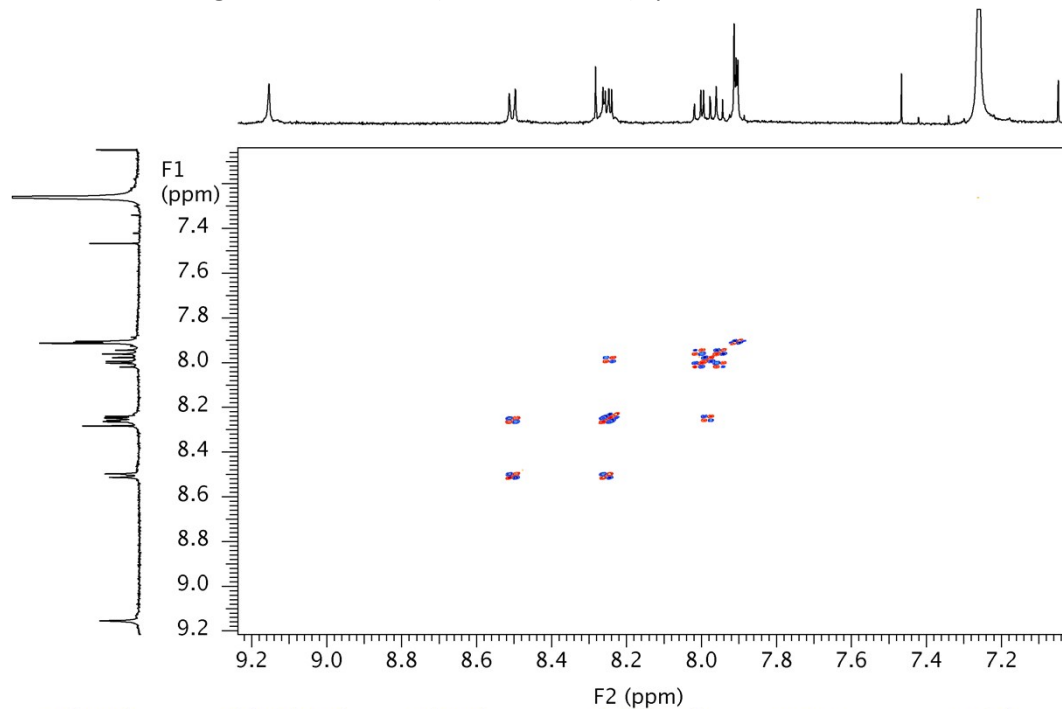


Figure S40: ¹H-¹H DQF-COSY (CDCl₃) of 2H-PTetraCor.

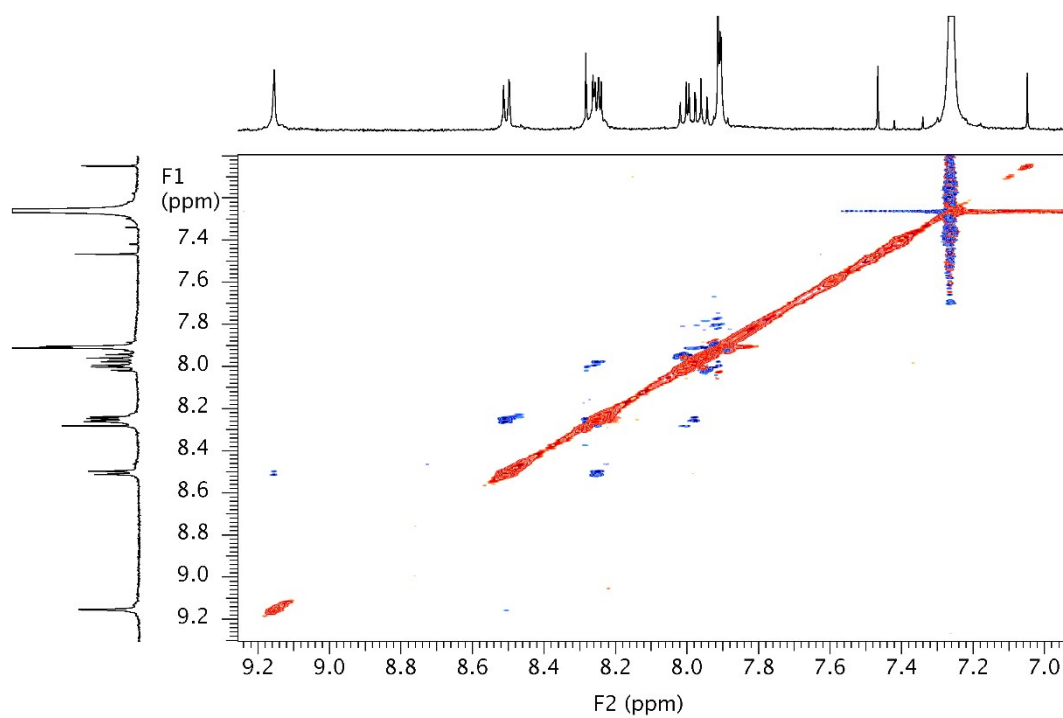


Figure S41: ^1H - ^1H band selective ROESY (CDCl_3) of **2H-PTetraCor**.

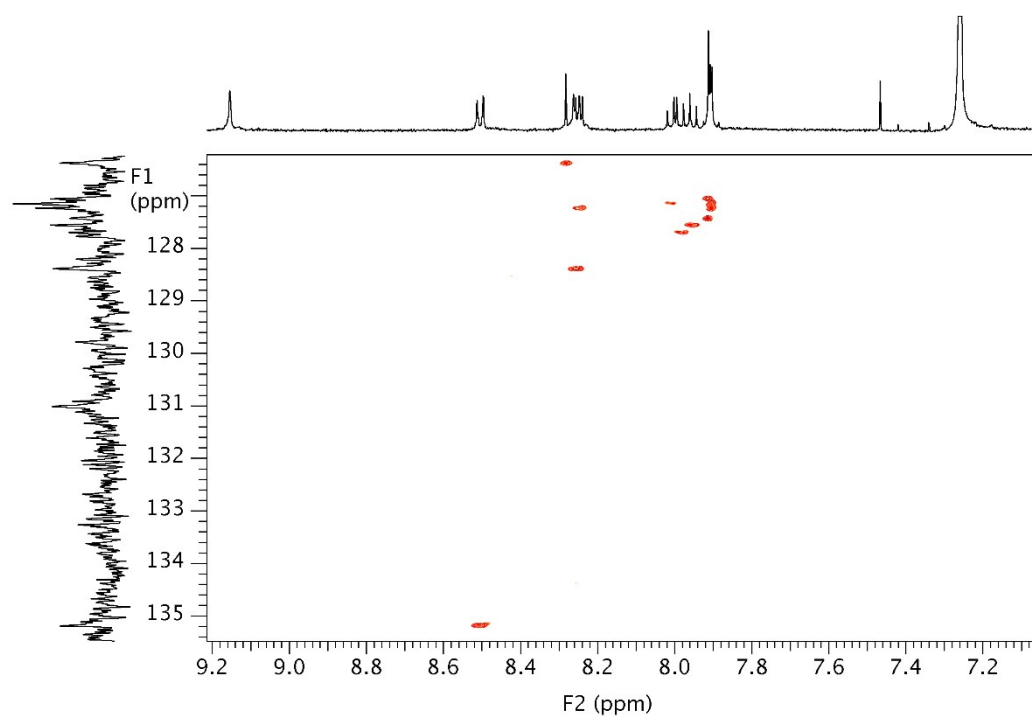


Figure S42: ^1H - ^{13}C band selective HSQC (CDCl_3) of **2H-PTetraCor**.

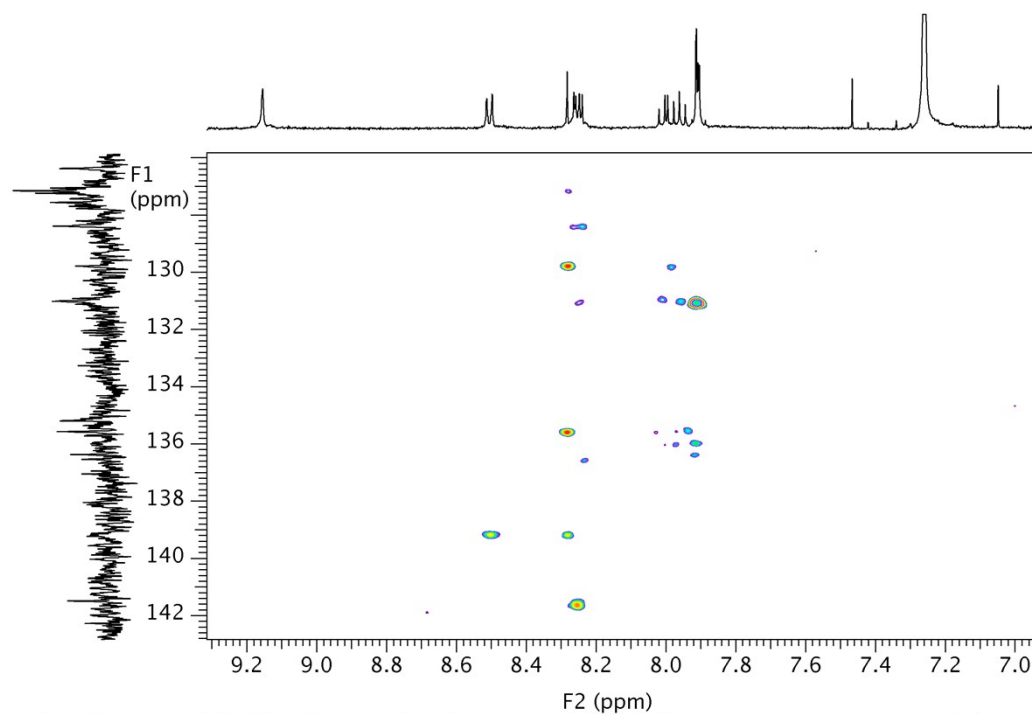


Figure S43: ^1H - ^{13}C band selective HMBC (CDCl_3) of **2H-PTetraCor**.

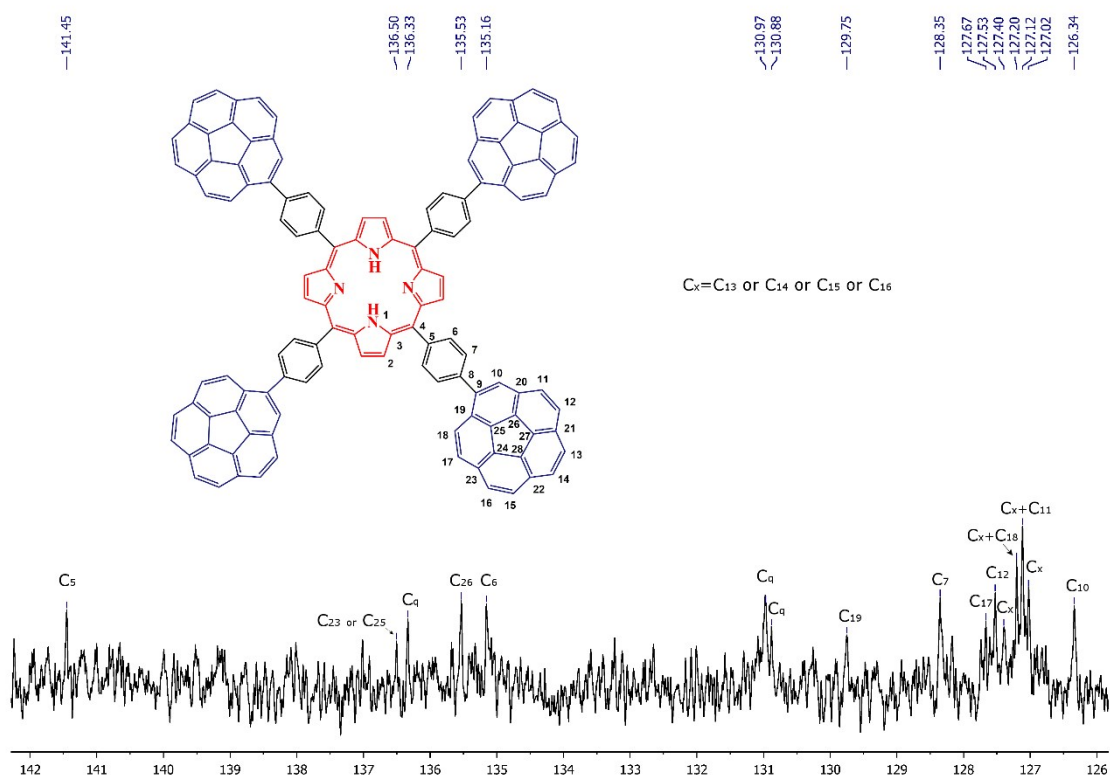


Figure S44: ^{13}C $\{^1\text{H}\}$ NMR (126 MHz, CDCl_3) spectrum of **2H-PTetraCor**.

UV/Vis spectra

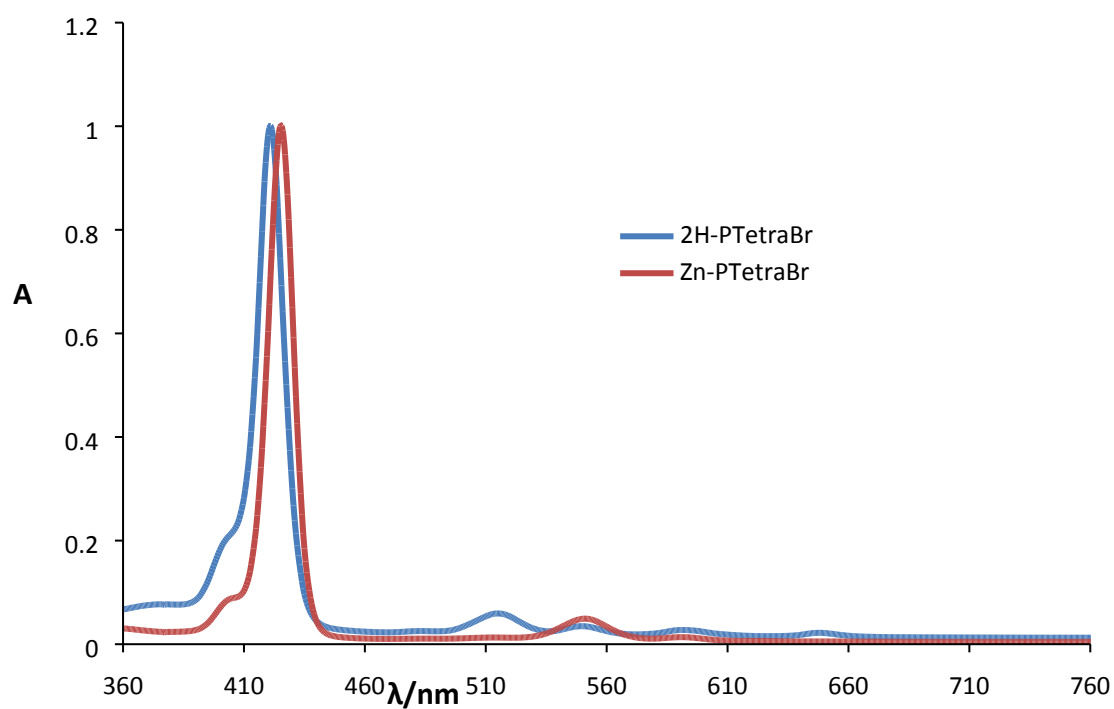


Figure S45: Normalized UV-Vis absorption spectra of **2H-PTetraBr** and **Zn-PTetraBr** in toluene.

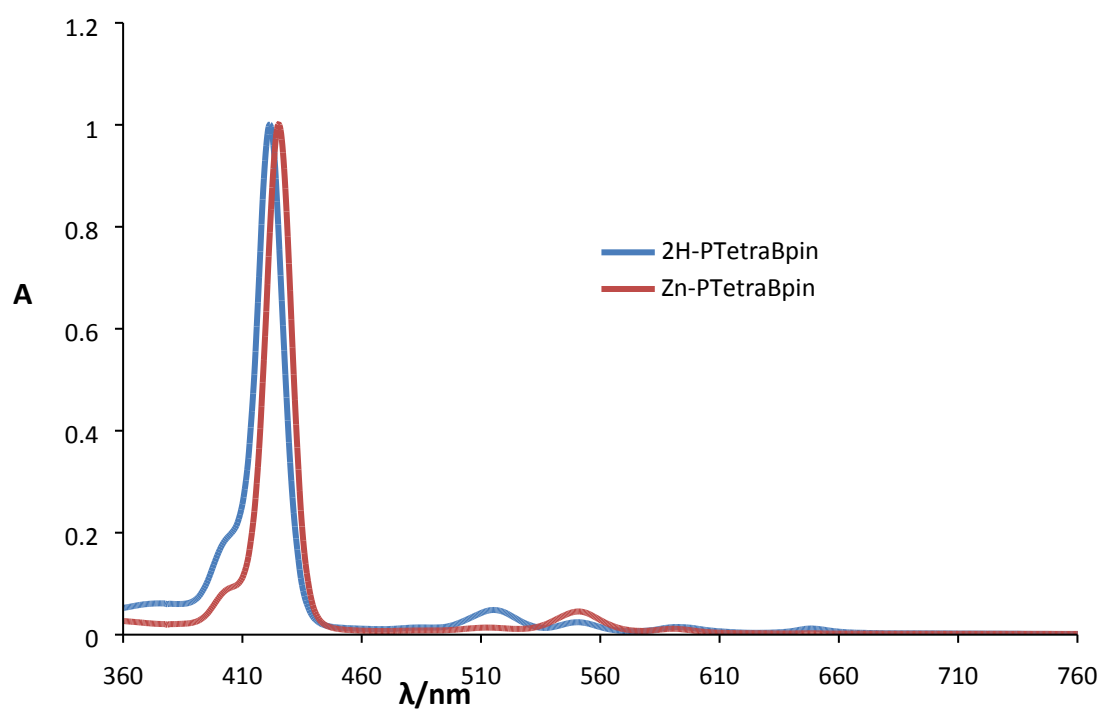


Figure S46: Normalized UV-Vis absorption spectra of **2H-PTetraBpin** and **Zn-PTetraBpin** in toluene.

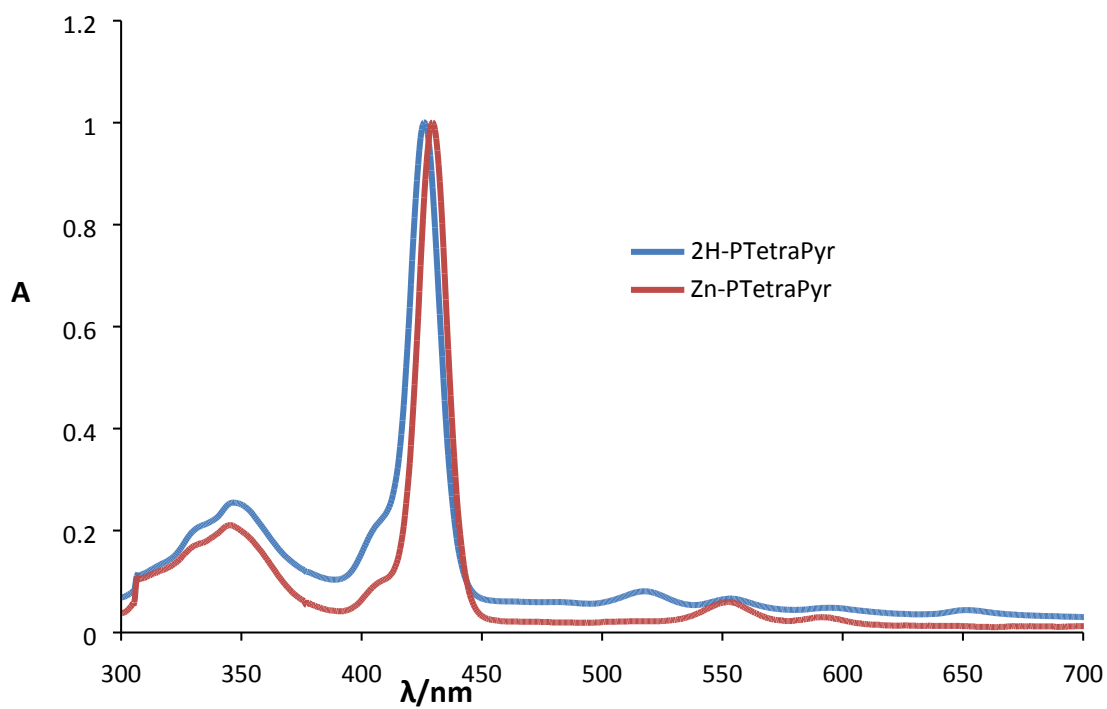


Figure S47: Normalized UV-Vis absorption spectra of **2H-PTetraPyr** and **Zn-PTetraPyr** in toluene.

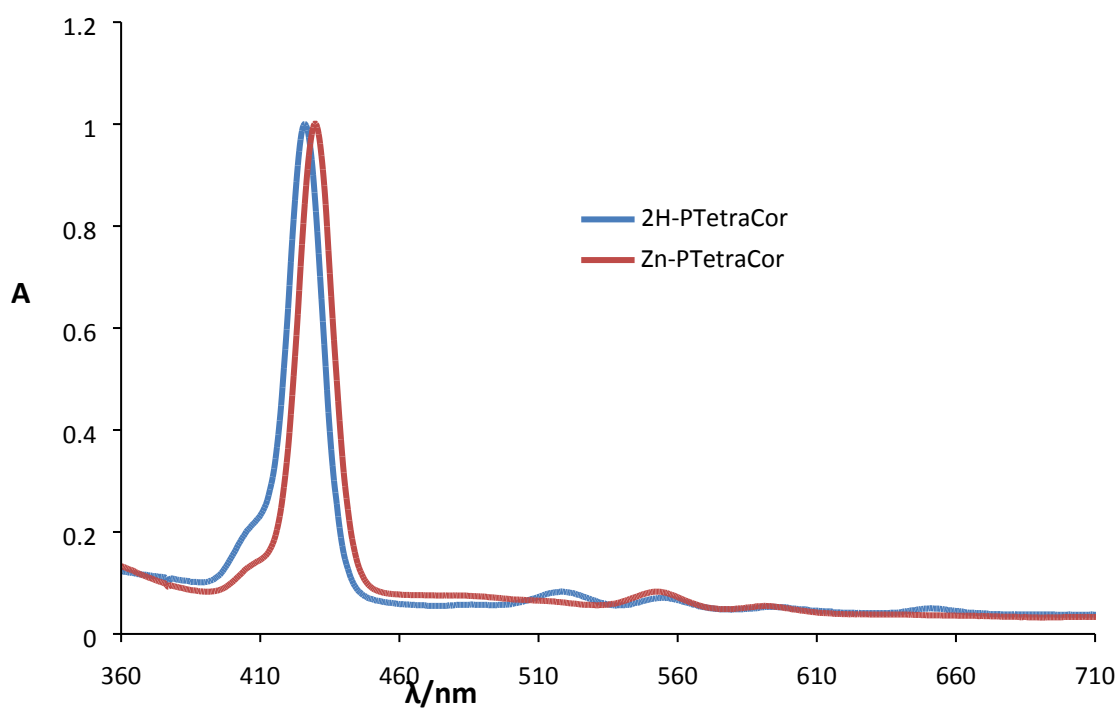


Figure S48: Normalized UV-Vis absorption spectra of **Zn-PTetraCor** and **2H-PTetraCor** in toluene.

MS spectra

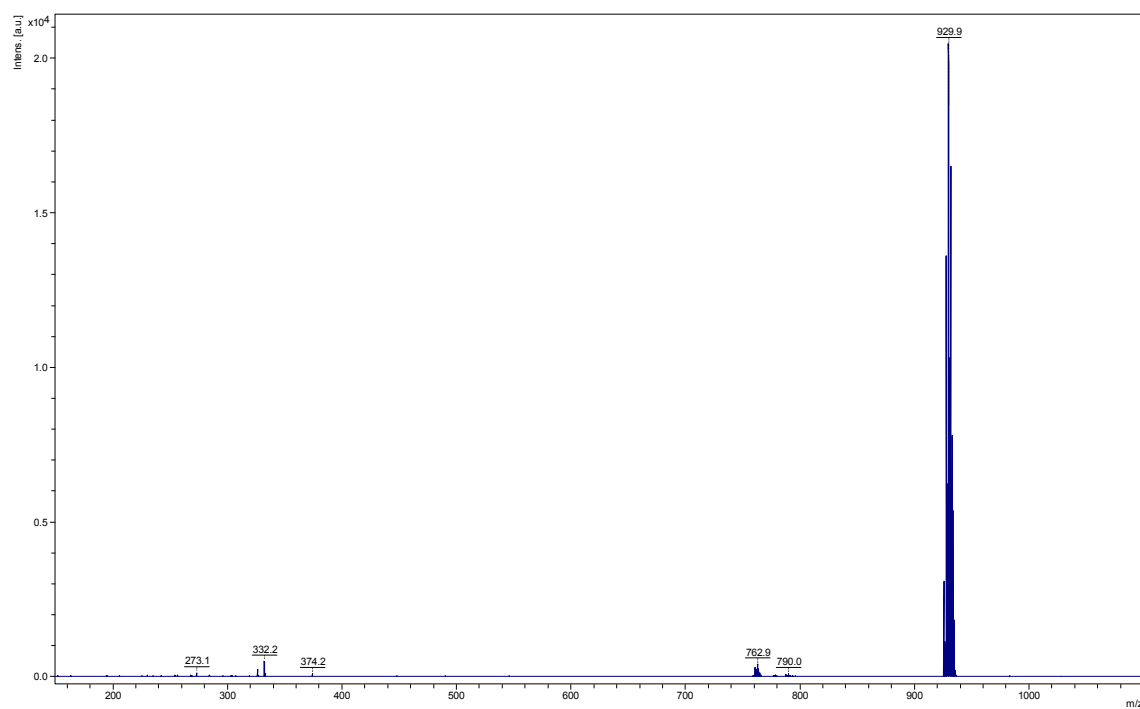


Figure S49: Full MS (MALDI-TOF) in DCTB of 2H-PTetraBr [M]⁺.

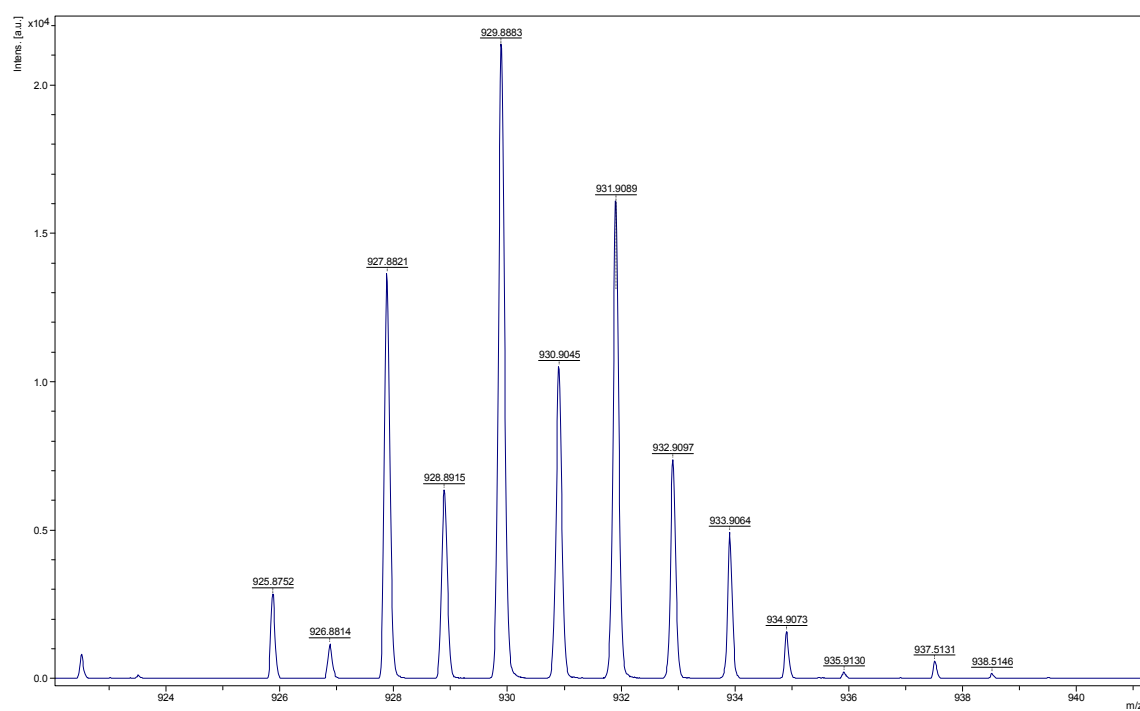


Figure S50: Expanded region of MS (MALDI-TOF) in DCTB of 2H-PTetraBr [M]⁺.

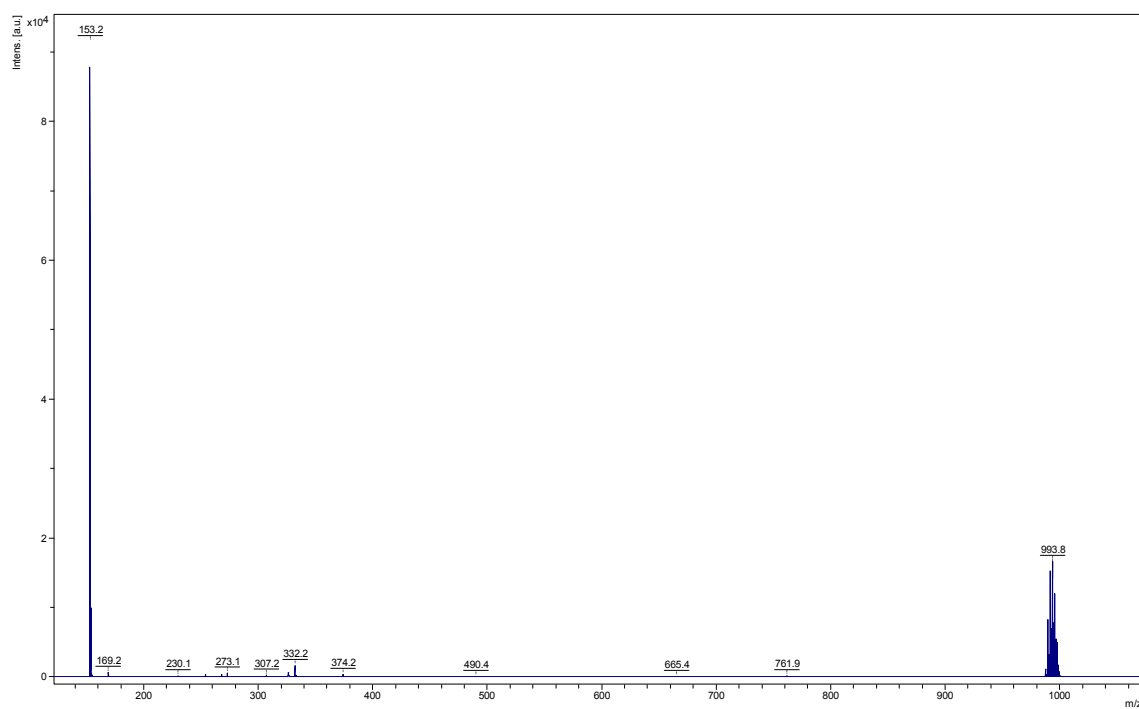


Figure S51: Full MS (MALDI-TOF) in DCTB of Zn-PTetraBr [M]⁺.

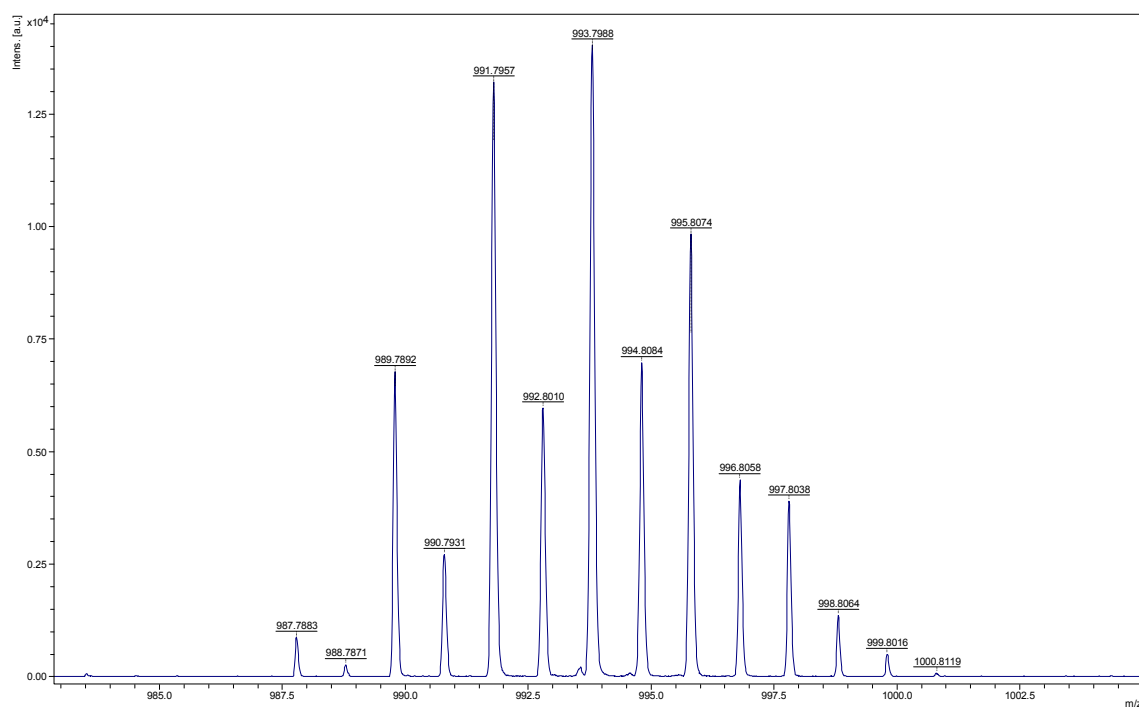


Figure S52: Expanded region of MS (MALDI-TOF) in DCTB of Zn-PTetraBr [M]⁺.

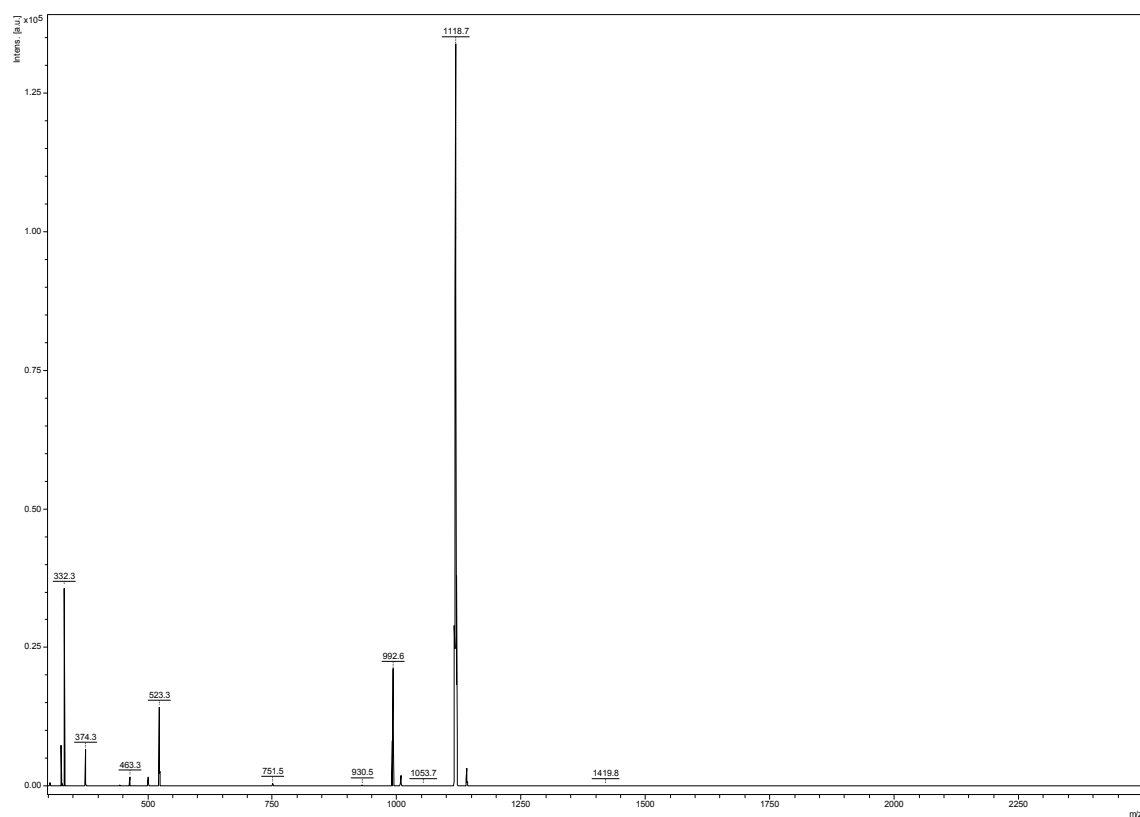


Figure S53: Full MS (MALDI-TOF) in DCTB of 2H-PTetraBpin [M]⁺.

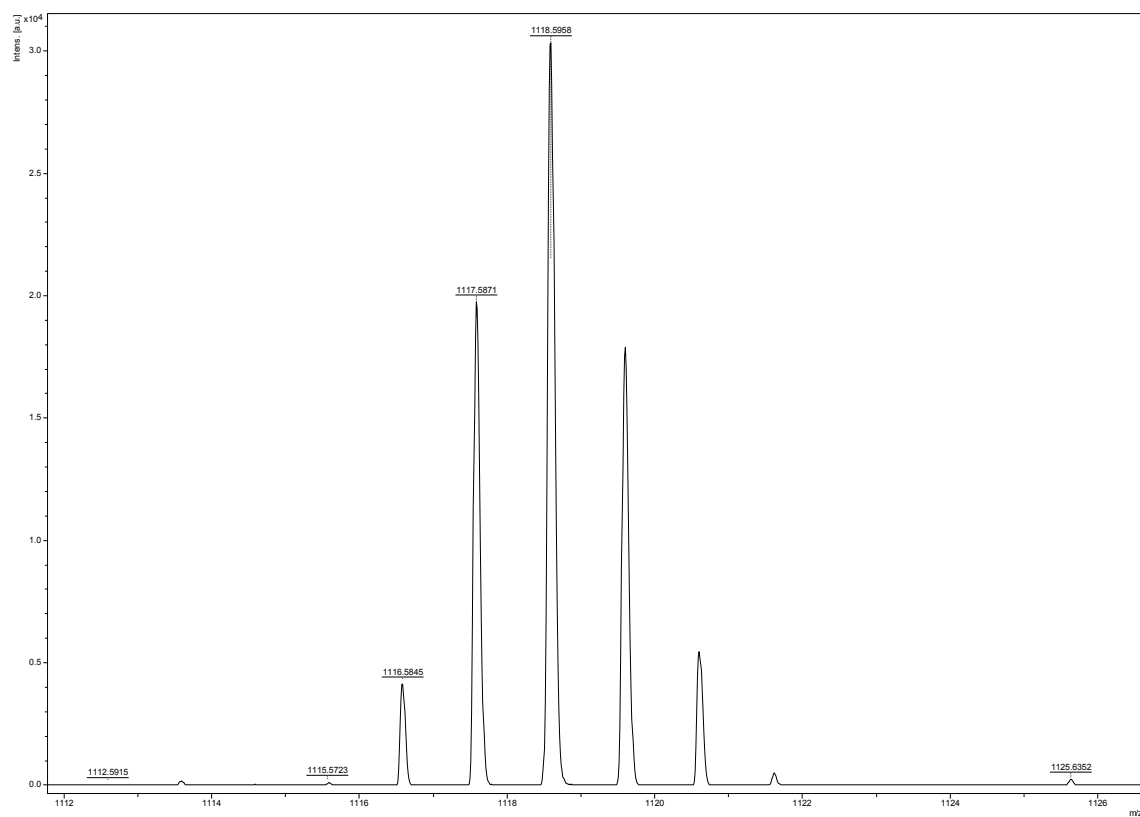


Figure S54: Expanded region of MS (MALDI-TOF) in DCTB of 2H-PTetraBpin [M]⁺.

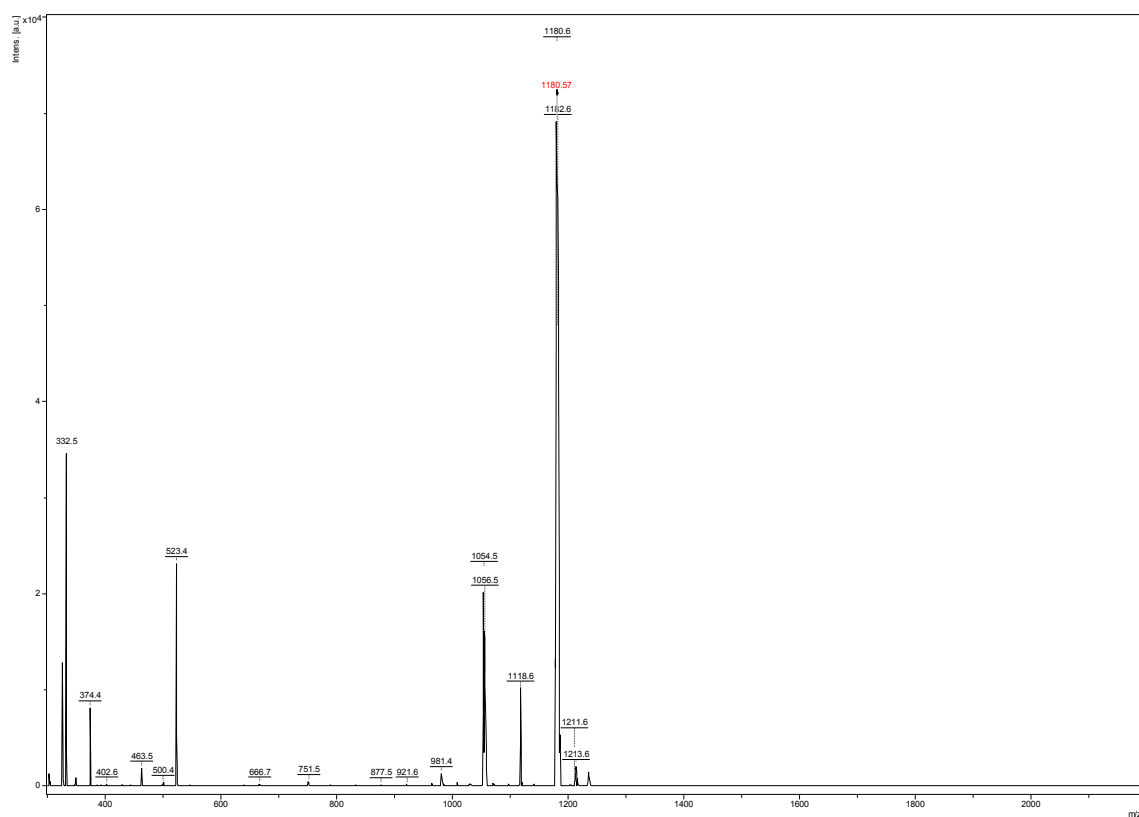


Figure S55: Full MS (MALDI-TOF) in DCTB of Zn-PTetraBpin [M]⁺.

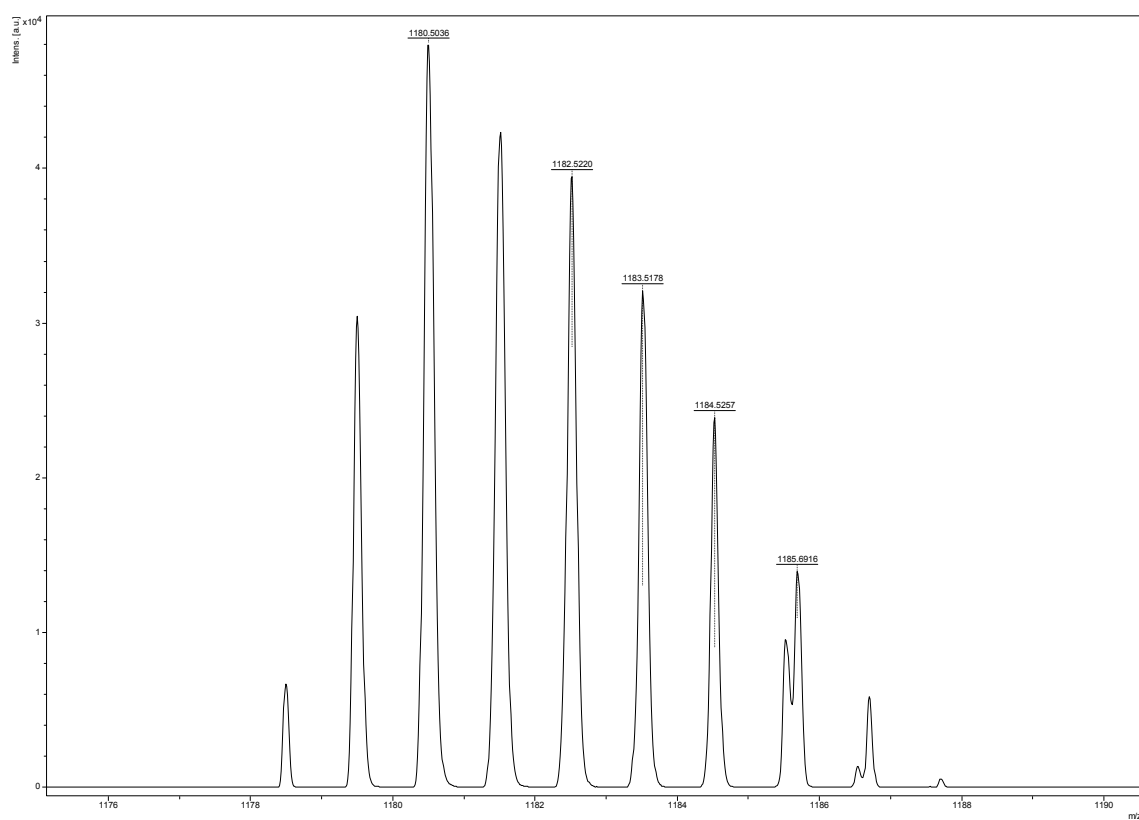


Figure S56: Expanded region of MS (MALDI-TOF) in DCTB of Zn-PTetraBpin [M]⁺.

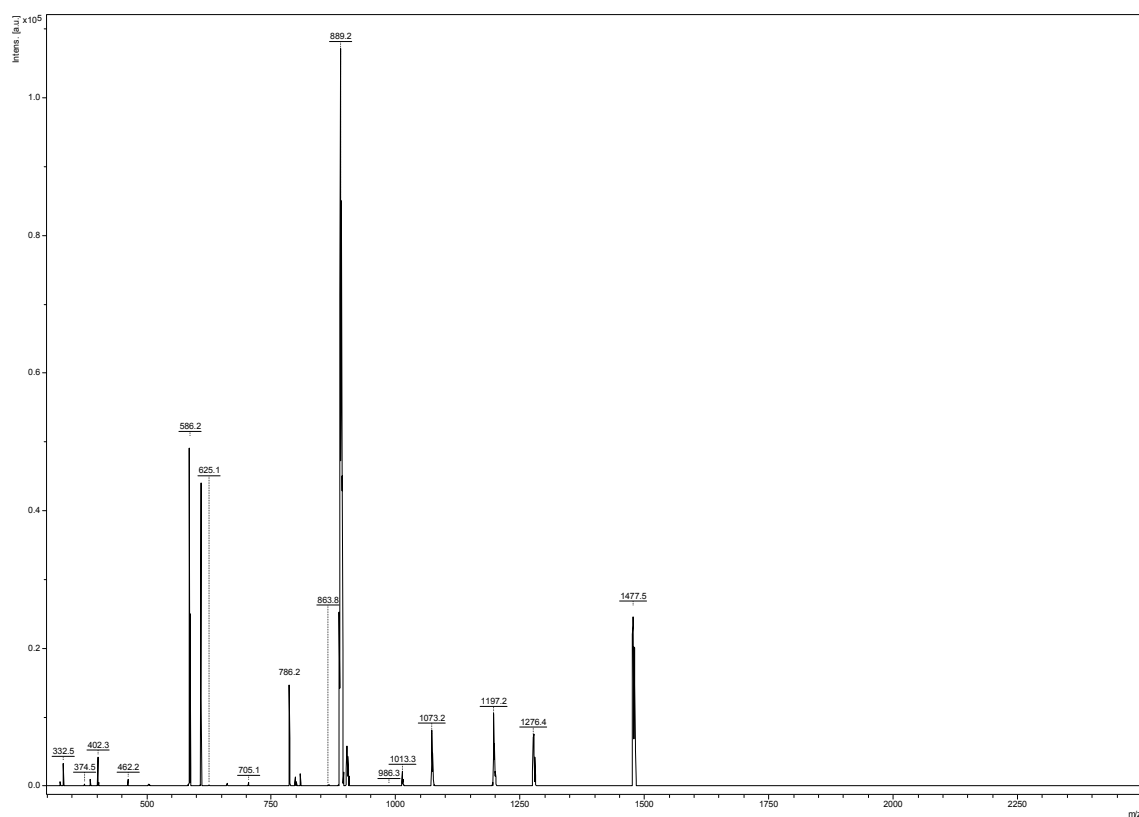


Figure S57: Full MS (MALDI-TOF) in DCTB of Zn-PTetraPyr [M]⁺.

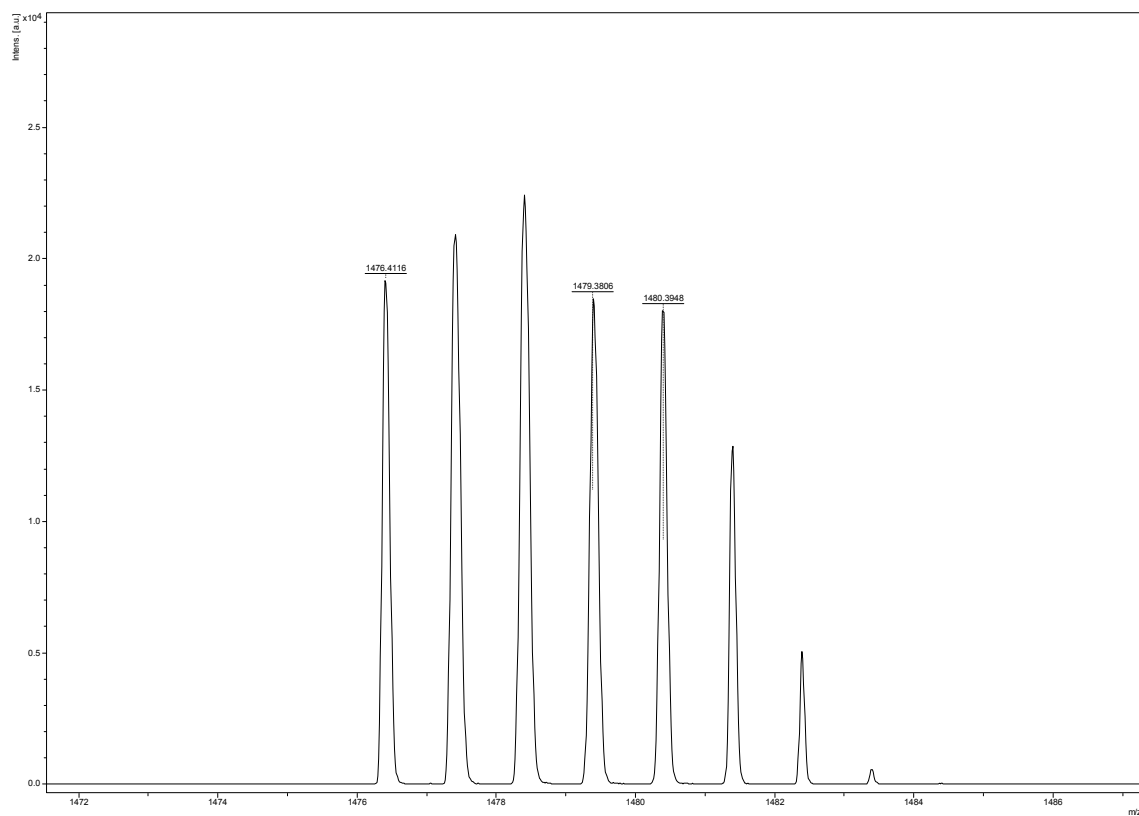


Figure S58: Expanded region of MS (MALDI-TOF) in DCTB of Zn-PTetraPyr [M]⁺.

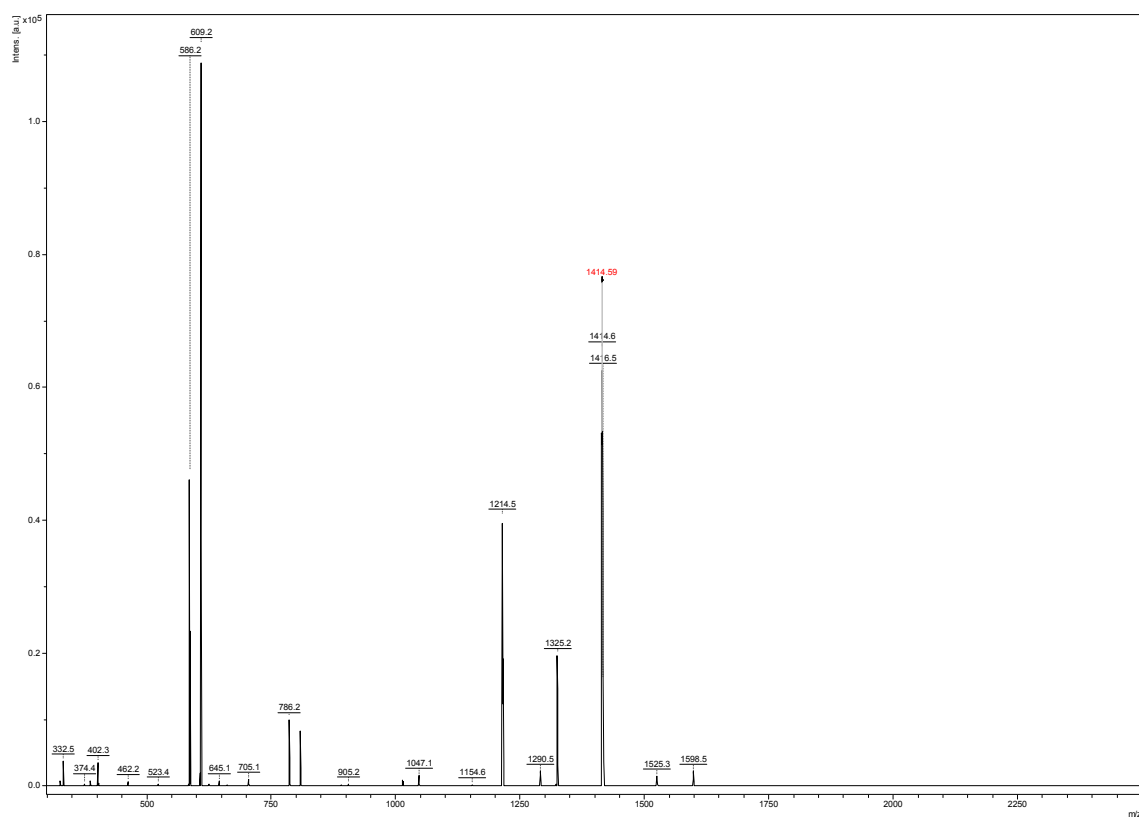


Figure S59: Full MS (MALDI-TOF) in DCTB of 2H-PTetraPyr [M]⁺.

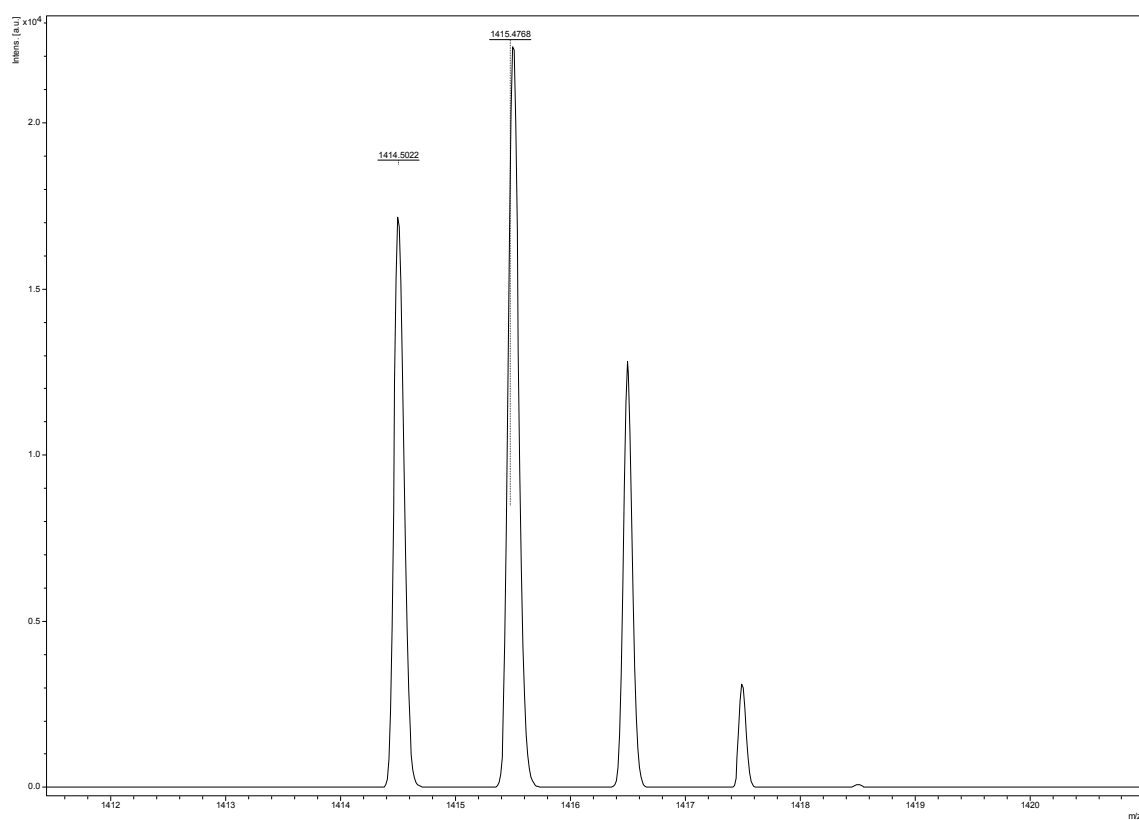


Figure S60: Expanded region of MS (MALDI-TOF) in DCTB of 2H-PTetraPyr [M]⁺.

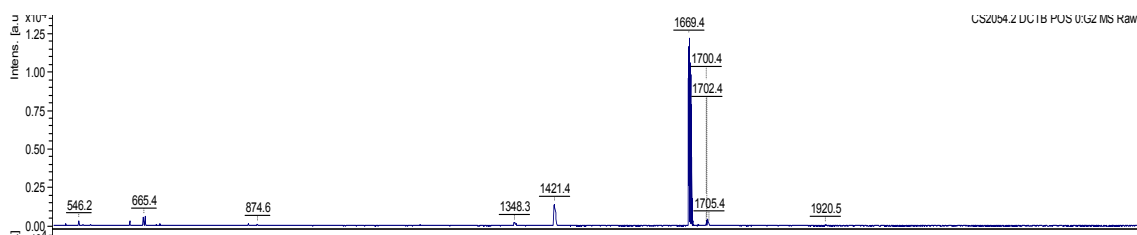


Figure S61: Full MS (MALDI-TOF) in DCTB of Zn-PTetraCor [M]⁺.

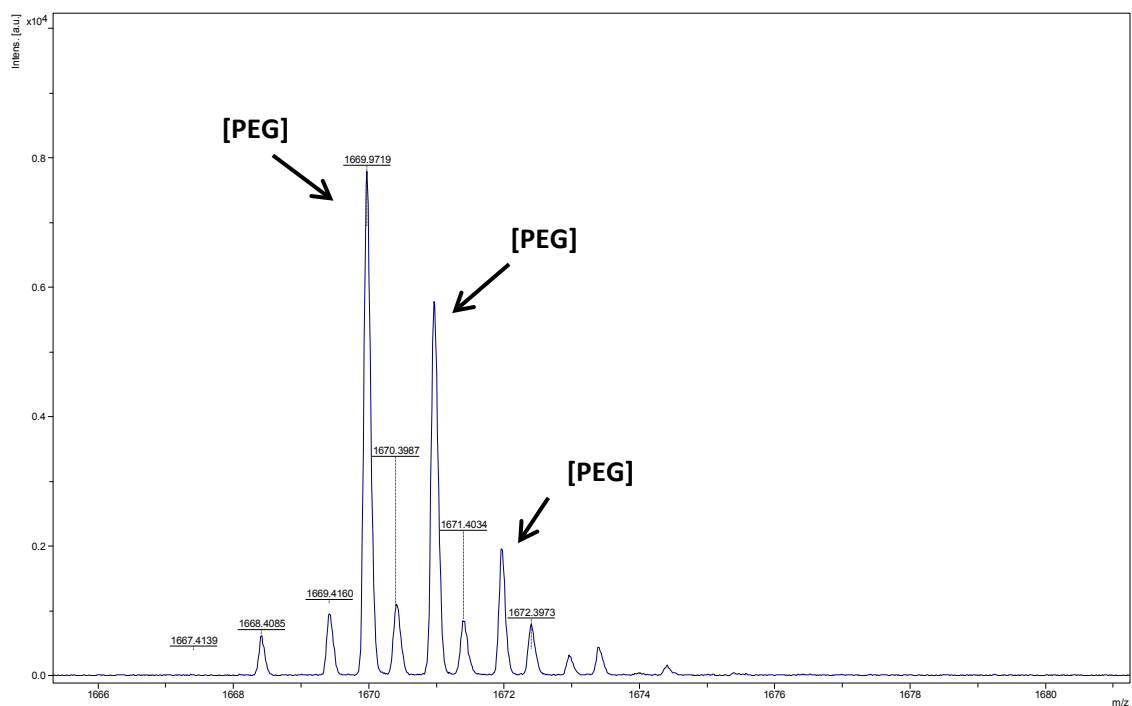


Figure S62: Expanded region of MS (MALDI-TOF) in DCTB of Zn-PTetraCor [M]⁺.

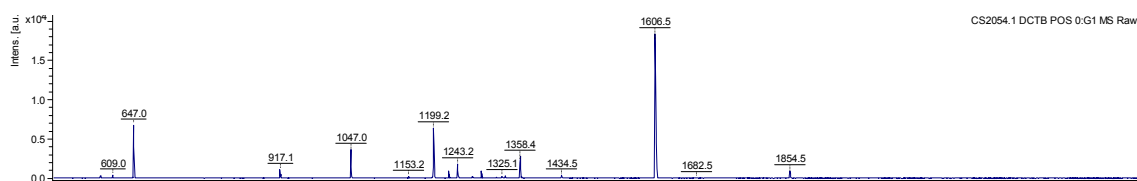


Figure S63: Full MS (MALDI-TOF) in : 1-Dithranol-negative, 2-Dithranol-positive, 3-DCTB-positive, 4-DCTB-negative of 2H-PTetraCor [M]⁺.

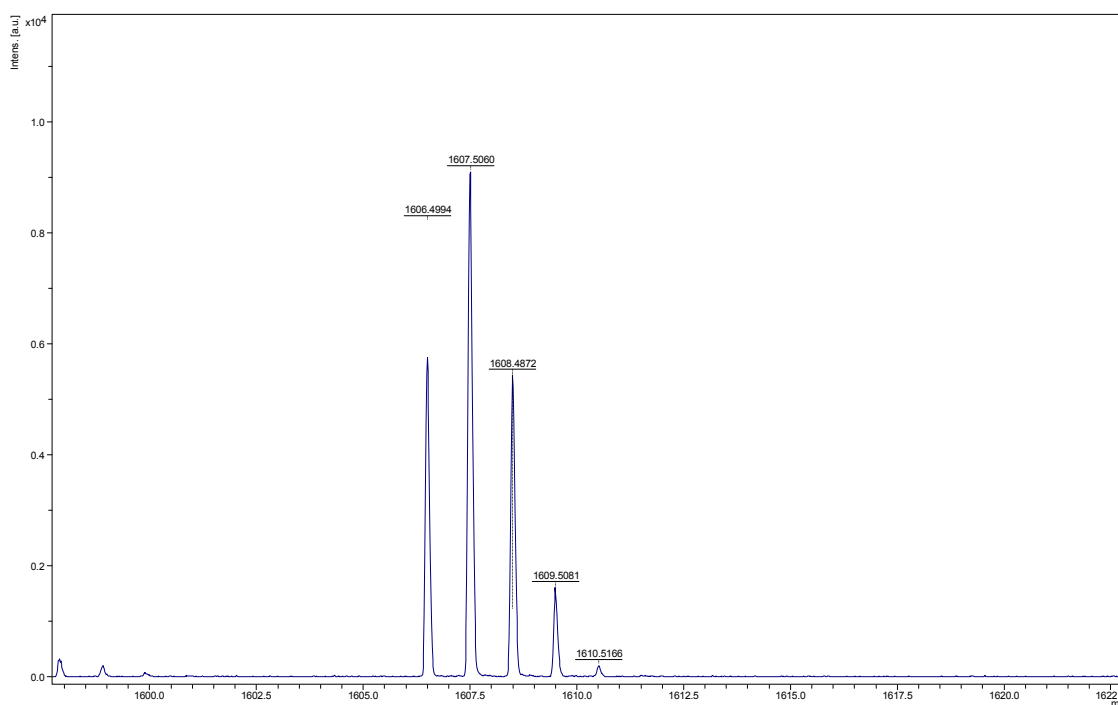


Figure S64: Expanded region of MS (MALDI-TOF) in DCTB of **2H-PTetraCor** [M]⁺.

Complexation measurements

In order to estimate the association constants (K_a) of the compounds **Zn-PTetraCor** and **2H-PTetraCor** with fullerenes, the dilution method was applied. A 10^{-4} M deuterated toluene solution of each compound was prepared, and a known volume was transferred to an NMR tube (500 μ L). The titration was carried out by adding known portions of a stock solution of C₆₀ or C₇₀ (10^{-3} M) in deuterated toluene to cover a wide range of equivalents. A ^1H NMR spectrum was recorded at room temperature after each addition. Once all data had been obtained, the changes in the chemical shifts ($\Delta\delta$) of selected protons were plotted as a function of the molar fraction of the guest, and the resulting curve was fitted by a nonlinear method using the global analysis approach according to the following equations, depending on the type of equilibrium:¹

1:1 Equilibria

General expression for the equilibrium constant:

$$K_a = \frac{[HG]}{[H][G]} \quad \text{eq.1}$$

Changes upon NMR titration:

$$\Delta\delta = \Delta\delta_{\Delta HG} \left(\frac{[HG]}{[H]_0} \right) \quad \text{eq.2}$$

Where:

[HG] is the concentration of the guest of the complex, and is calculated using the following equation:

$$[HG] = \frac{1}{2} \left([G_0] + [H_0] + \frac{1}{K_a} \right) - \sqrt{\left([G_0] + [H_0] + \frac{1}{K_a} \right)^2 - 4[G_0][H_0]} \quad \text{eq.3}$$

Where:

[G₀] is the total concentration of the guest

[H₀] is the total concentration of the host

Δδ_{ΔHG} is Δδ at maximum complexation (100% supramolecular complex formation)

K_a is the estimated association constant for 1:1 equilibrium

Δδ_{ΔHG} and K_a for a 1:1 equilibrium were extracted using the non-linear curve fitting tool at the open access web portal <http://supramolecular.org> (accessed since 2016). Links to all the fittings of the data are provided below for every case.

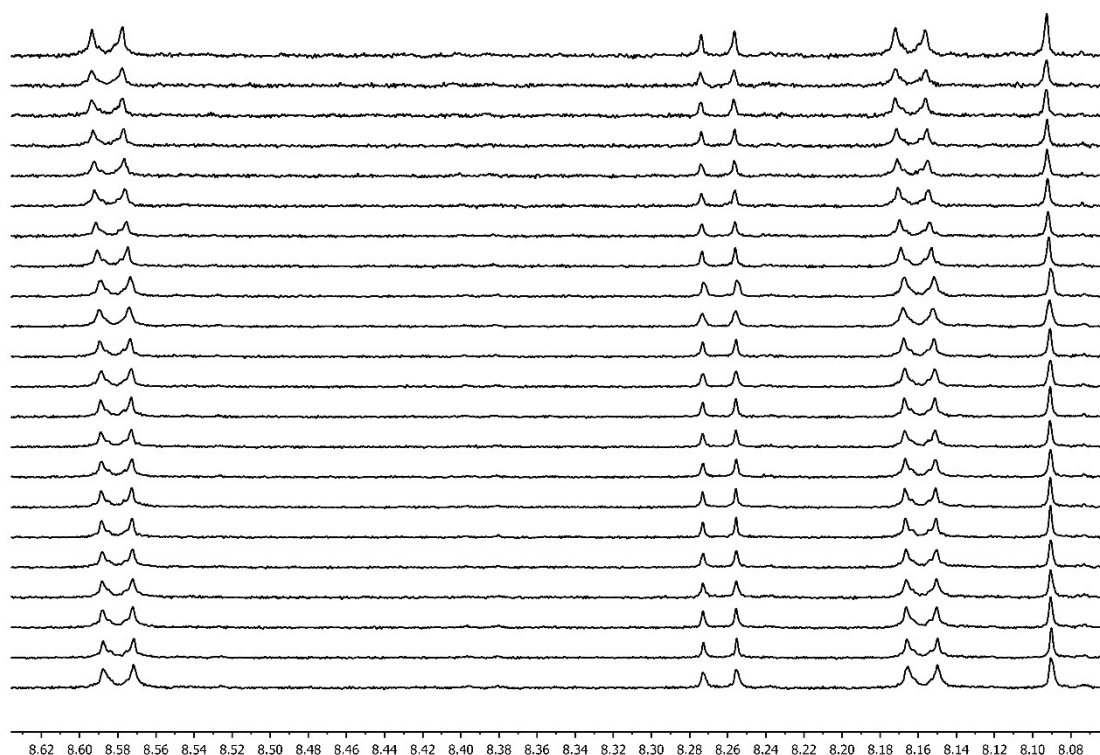


Figure S65: ^1H -NMR spectra of the titration of **Zn-PTetraCor** (10^{-4} M) with variable concentrations of C_{60} (10^{-3} M) in toluene- d_8 .

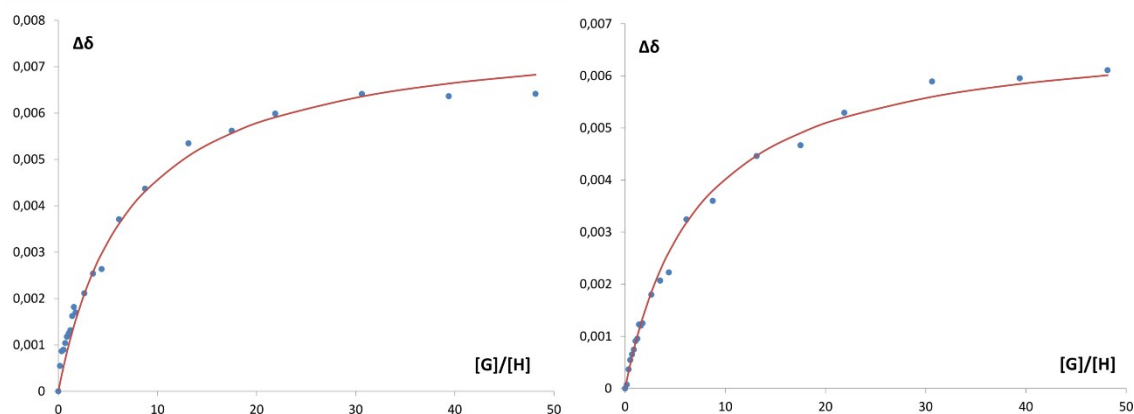


Figure S66: Nonlinear regressions for the results of the titration of **Zn-PTetraCor** (10^{-4} M) with C_{60} (10^{-3} M) for selected protons (right plot: H_6 proton, left plot: H_7 , 1:1 binding model).

For additional information see:

<http://app.supramolecular.org/bindfit/view/c40ba3c1-1658-43b7-965e-17b2e510871b>

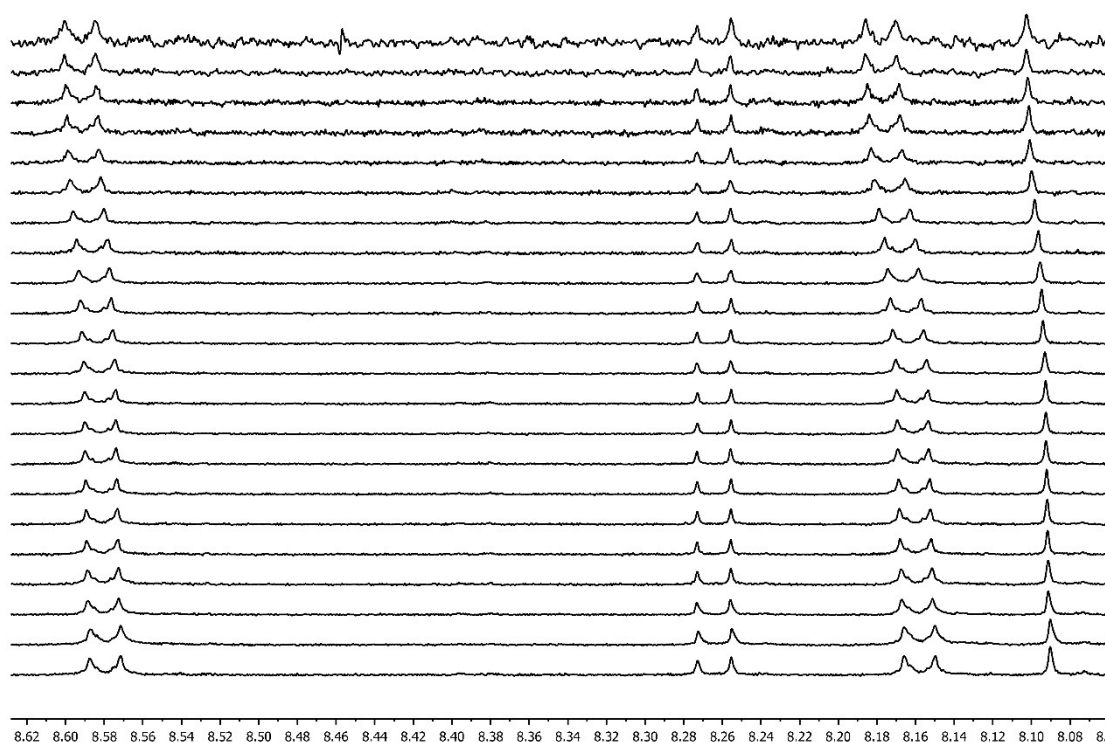


Figure S67: ^1H -NMR spectra of the titration of **Zn-PTetraCor** (10^{-4} M) with variable concentrations of **C₇₀** (10^{-3} M) in toluene- d_8 .

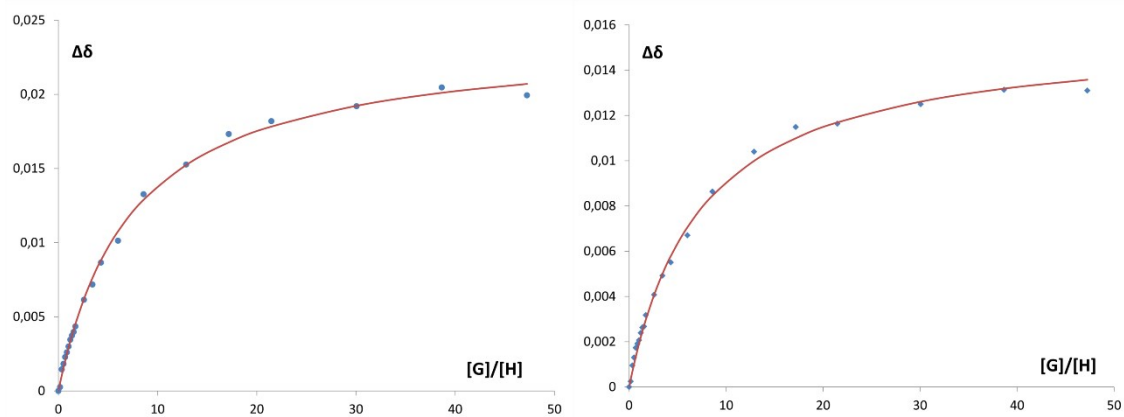


Figure S68: Nonlinear regressions for the results of the titration of **Zn-PTetraCor** (10^{-4} M) with **C₇₀** (10^{-3} M) for selected protons (right plot: H_6 proton, left plot: H_7 , 1:1 binding model).

For additional information see:

<http://app.supramolecular.org/bindfit/view/1d6856af-192c-4e5a-9d53-f609627ac179>

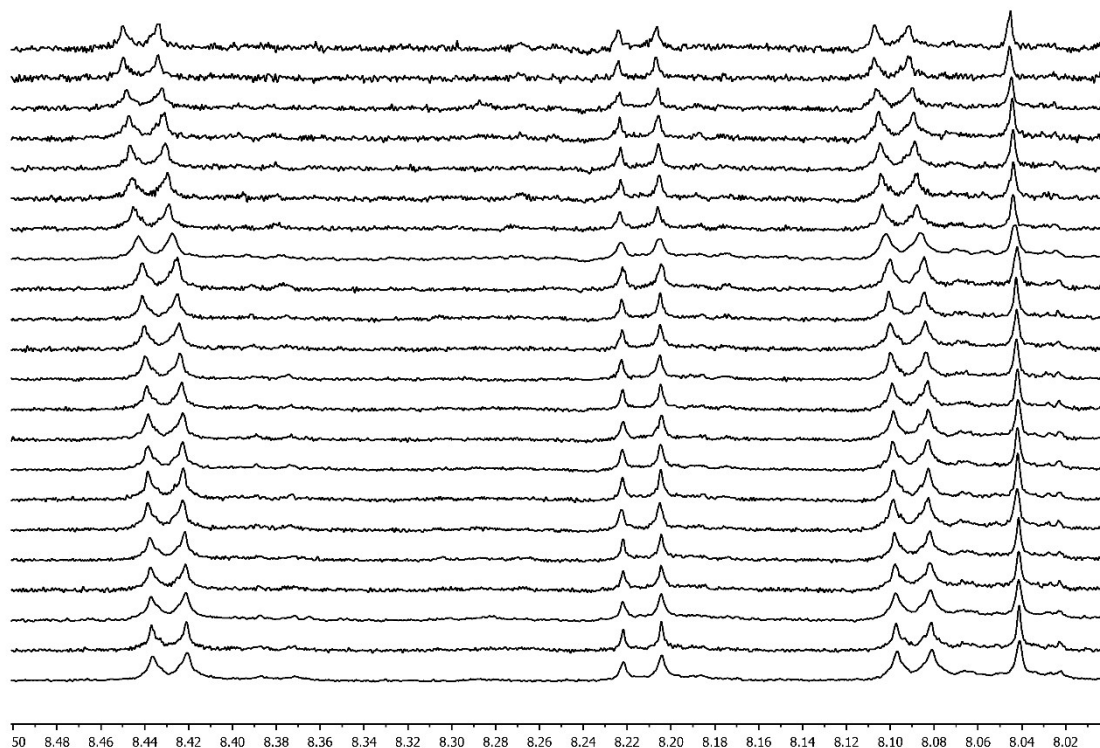


Figure S69: ^1H -NMR spectra of the titration of **2H-PTetraCor** (10^{-4} M) with variable concentrations of C_{60} (10^{-3} M) in toluene- d_8 .

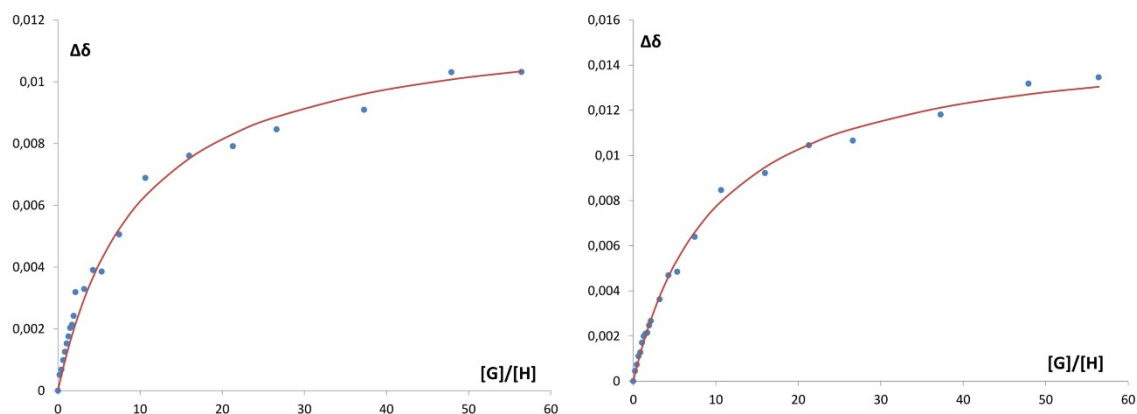


Figure S70: Nonlinear regressions for the results of the titration of **2H-PTetraCor** (10^{-4} M) with C_{60} (10^{-3} M) for selected protons (right plot: H_6 proton, left plot: H_7 , 1:1 binding model).

For additional information see:

<http://app.supramolecular.org/bindfit/view/41c6686e-e06a-41fd-bc7c-ee6955080966>

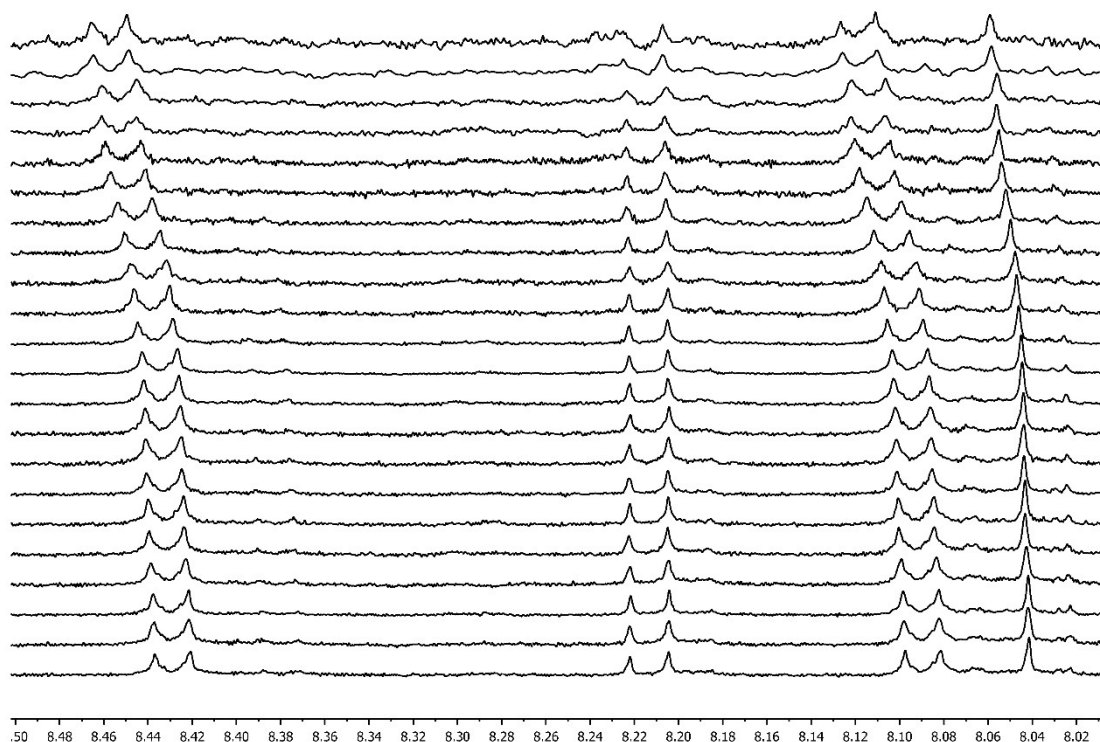


Figure S71: ^1H -NMR spectra of the titration of **2H-PTetraCor** (10^{-4} M) with variable concentrations of **C₇₀** (10^{-3} M) in toluene- d_8 .

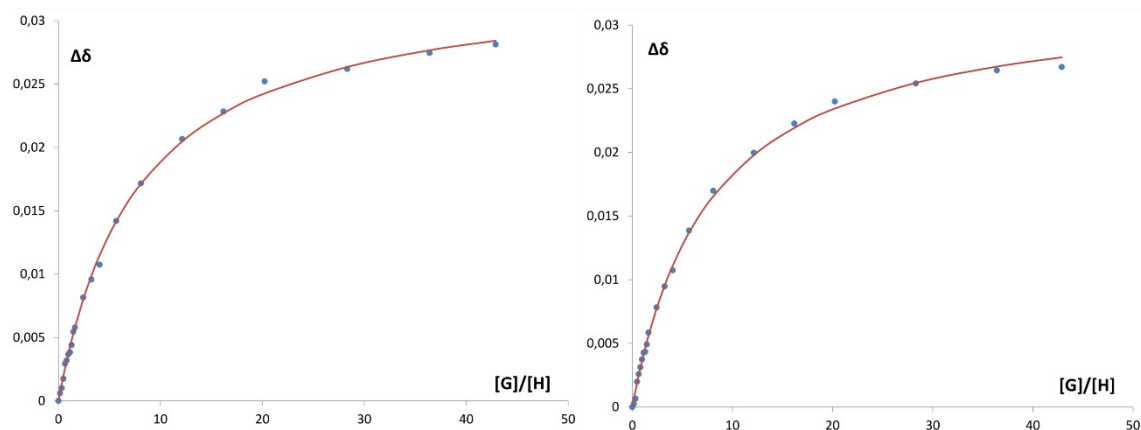


Figure S72: Nonlinear regressions for the results of the titration of **2H-PTetraCor** (10^{-4} M) with **C₇₀** (10^{-3} M) for selected protons (right plot: H₆ proton, left plot: H₇, 1:1 binding model).

For additional information see:

<http://app.supramolecular.org/bindfit/view/f3cbc3e0-aacf-4516-852f-a773e3b87783>

Computational methods

Optimized geometries of porphyrin **2H-PTetraCor** and the supramolecular assemblies $C_{60}@2H-PTetraCor$, $C_{70}@2H-PTetraCor$, $(C_{60})_2@2H-PTetraCor$ and $(C_{70})_2@2H-PTetraCor$ were obtained by DFT methods with the B97D3 functional, which contains the Becke-Johnson damping empirical dispersion correction and was provided by Grimme and collaborators.² Pople's split valence set 6-31G(d,p) was chosen as the basis set.³ Solvent corrections applied using the polarizable continuum model (PCM) using toluene ($\epsilon=2.3741$).⁴

The strategy to obtain the inclusion complexes $C_{60}@2H-PTetraCor$ and $C_{70}@2H-PTetraCor$ consisted of using the optimized structure of compound **2H-PTetraCor** and manually placing the corresponding fullerene molecule halfway between the two corannulenes, rotating the single C-C bonds at the same time so that both PAH fragments matched with the fullerene surface. For the assembly $C_{70}@2H-PTetraCor$, several attempts were carried out by imposing different orientations on the fullerene C_{70} ; only the most stable one was considered and is reported here. Once both 1:1 adducts were optimized, their structures were used as starting geometries for a second round of optimizations to obtain the complexes $(C_{60})_2@2H-PTetraCor$ and $(C_{70})_2@2H-PTetraCor$ by placing a second fullerene molecule following the same protocol described above.

All minima were confirmed by vibrational analysis to show no imaginary frequencies. The electronic energies of the optimized geometries were further evaluated using a more extended 6-31+G(d,p) basis set that includes diffuse functions.⁵

Deformation energies were estimated by subtracting the electronic energy of the optimized porphyrin **2H-PTetraCor** (H) from the electronic energy of the porphyrin in the optimized structure of the adduct (HG, i.e., $C_{60}@2H-PTetraCor$ and $C_{70}@2H-PTetraCor$) according to eq. 4. For the supramolecular assemblies $(C_{60})_2@2H-PTetraCor$ and $(C_{70})_2@2H-PTetraCor$ (HG₂), the subtraction was carried out from the porphyrin structure shown in the parent 1:1 inclusion complexes (HG) according to eq. 5:

$$E_{def} = E_{HG}(H) - E_H(H) \quad \text{eq.4}$$

$$E_{def} = E_{HG_2}(H) - E_{HG}(H) \quad \text{eq.5}$$

Where the subscripts denote the geometry used and the letter between parentheses corresponds to the molecular entity studied (porphyrin in all cases).

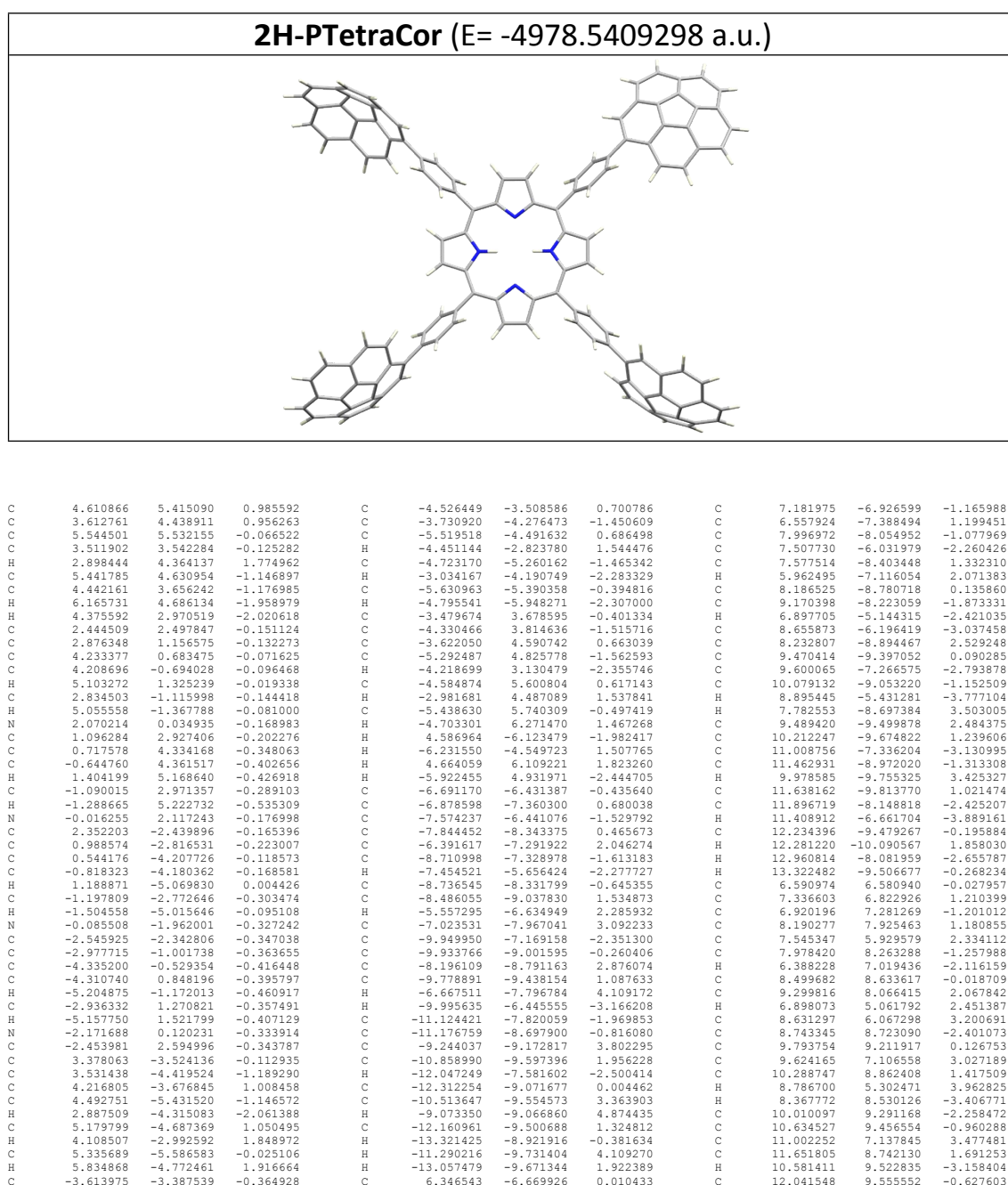
Interaction energies were calculated taking into account basis set superposition error (BSSE) with the Boys–Bernardi functional counterpoise scheme⁶ as follows (eq. 6):

$$E_{int}(HG) = E_{HG}^{HG}(HG) - E_{HG}^{HG}(H) - E_{HG}^{HG}(G) \quad \text{eq. 6}$$

Where the subscripts denote the geometry used (inclusion complex in all cases) and the superscripts refer to the basis set (from the supramolecular assembly in all cases); H and G correspond to the host and guest molecular entities, respectively, and HG to the supramolecular adduct.

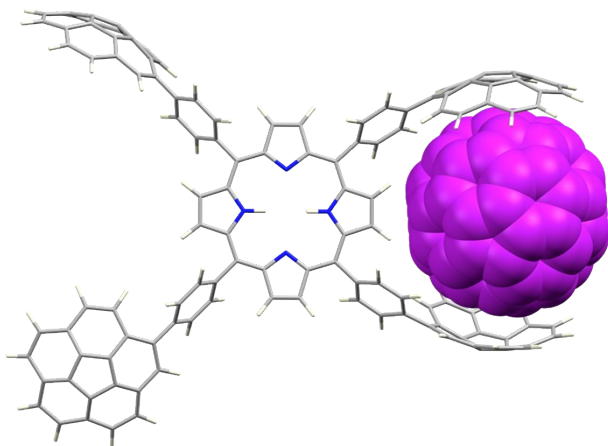
All the above-described computational methods were performed using the Gaussian 16 package.⁷

Non-covalent interactions were obtained from the location critical points at which the reduced density gradient decreases to low electronic density values according to the scheme of Yang et al. with the help of the NCIPLOT package.⁸ Calculations were performed with promolecular densities, and gradient isosurfaces were plotted with an isovalue of 0.3 a.u. and coloured on a blue-green-red scale according to the values of the sign of λ_2 (the second eigenvalue of the electron-density Hessian). Red indicates repulsion, green indicates weak attraction, and blue represents strong attraction. Graphics were visualized in Chimera⁹ with the help of Tangram NCIPLOT GUI built by Insilichem Group.¹⁰



C	11.968716	7.917243	2.840717	H	-7.911861	4.311527	-0.977097	H	-12.716319	5.521093	-0.468468
H	11.318979	6.458980	4.270367	C	-9.886907	5.128786	-1.068925	C	-11.991047	10.025626	-0.732909
C	12.526477	9.215628	0.636782	C	-7.126582	10.368996	0.764382	H	-11.076108	11.686509	1.732241
H	12.758659	9.805289	-1.410740	C	-9.510744	9.619813	-0.455179	H	-13.627856	7.642921	0.315526
H	13.008003	7.821344	3.157935	C	-10.738740	6.298862	-1.035418	H	-12.945771	10.204000	1.229465
H	13.605406	9.211672	0.797390	C	-10.603163	8.751266	-0.747488	H	1.053761	0.052817	-0.209021
C	-6.457856	6.813974	-0.565029	H	-10.336995	4.138552	-0.986338	H	-1.155077	0.102556	-0.298362
C	-7.811011	6.506730	-1.036990	H	-6.212137	10.678754	1.271913				
C	-6.146522	8.102408	-0.098445	C	-8.308746	11.058706	1.037279				
C	-8.645142	7.612272	-1.201917	C	-9.587849	10.627311	0.507853				
C	-8.494272	5.227416	-1.065432	C	-12.106429	6.419876	-0.568856				
C	-7.109624	9.178805	-0.064022	C	-11.832773	8.846028	-0.094496				
H	-5.149287	8.268453	0.310157	H	-8.277622	11.884527	1.749320				
C	-8.303503	8.916767	-0.735320	C	-10.920546	10.874471	1.020682				
C	-10.069034	7.510922	-1.206597	C	-12.627864	7.633874	-0.120225				

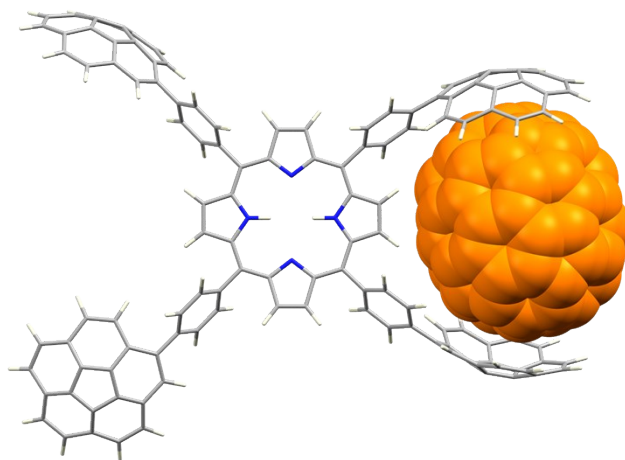
C₆₀@2H-PTetraCor (E= -7263.3537726 a.u.)



C	1.268664	5.224657	0.962646	C	-7.199843	4.557200	0.365641	H	8.899534	-6.381882	2.319063
C	0.113317	4.462660	0.772810	C	-8.855837	4.648548	-1.881170	C	9.897321	-6.265487	-0.418790
C	2.369997	5.111863	0.085754	H	-7.738600	2.941183	-2.583692	C	8.726318	-5.317151	-3.944232
C	0.007235	3.574044	-0.315271	C	-8.183911	5.542452	0.264035	H	7.373387	-4.427337	-5.346795
H	-0.695257	4.509711	1.501021	H	-6.565249	4.509434	1.249458	C	10.035405	-5.968445	-1.775612
C	2.214618	4.302879	-1.060206	C	-9.030473	5.609907	-0.862582	H	10.750030	-6.098732	0.240068
C	1.050319	3.562407	-1.264121	H	-8.324466	6.250035	1.080203	H	9.591824	-4.882536	-4.445187
H	3.041922	4.194671	-1.758844	H	1.476771	-5.884798	-1.844870	H	10.990251	-5.577023	-2.127685
H	0.973020	2.907052	-2.130172	H	-9.651360	-4.99637	1.785596	C	3.670071	5.746381	0.403410
C	-1.084696	2.556667	-0.361515	H	1.338454	5.888496	1.822906	C	4.162316	5.725270	1.787305
C	-0.644298	1.217387	-0.295391	H	-9.480887	4.697991	-2.771779	C	4.497268	6.215425	-0.633157
C	0.718838	0.762927	-0.252001	C	-10.072588	-6.537231	-0.004090	C	5.372218	6.389861	1.993755
C	0.710377	-0.610818	-0.233686	C	-10.261176	-7.372099	1.183631	C	3.753142	4.874091	2.890092
H	1.585944	1.405563	-0.240659	C	-10.942600	-6.651148	-1.103000	C	5.844965	6.682511	-0.414706
C	-0.655584	-1.053991	-0.203504	C	-11.211692	-8.383016	1.042100	H	4.125276	6.140901	-1.654766
H	1.570117	-1.261863	-0.236174	C	-9.791491	-7.183464	2.544705	C	6.193334	6.857075	0.923652
N	-1.437913	0.084579	-0.262848	C	-12.066140	-7.559088	-1.124127	C	6.225381	6.113945	3.105399
C	-2.436839	2.965763	-0.414365	H	-10.824916	-5.929942	-1.912573	H	2.817905	4.323072	2.819855
C	-2.843668	4.356946	-0.605161	C	-12.090216	-8.477446	-0.075781	C	4.583525	4.613007	3.980636
C	-4.207217	4.352841	-0.663159	C	-11.857321	-8.994286	2.158692	C	6.957222	6.733862	-1.344364
H	-2.171454	5.200987	-0.710742	H	-8.967923	-6.498406	2.738188	C	7.546603	6.865648	1.369686
C	-4.623113	2.957254	-0.499912	C	-10.427589	-7.777052	3.636575	C	5.919973	5.159963	4.075158
H	-4.869361	5.194981	-0.825638	C	-13.297885	-7.479518	-1.886946	C	7.567317	6.408948	2.719911
N	-3.531118	2.132871	-0.357429	C	-13.283198	-9.129407	0.350761	H	4.254911	3.877679	4.715562
C	-1.095077	-2.391735	-0.102546	C	-11.587041	-8.632129	3.477584	H	6.760965	6.656854	-2.414182
C	-2.446716	-2.805610	-0.102435	C	-13.139223	-9.448853	1.732396	C	8.282367	6.736857	-0.907314
C	-2.862697	-4.190750	0.112300	H	-10.085633	-7.517739	4.639426	C	8.625287	6.691841	0.500668
C	-4.227130	-4.190651	0.077292	H	-13.343045	-6.827698	-2.760365	C	7.059576	4.659118	4.818304
H	-2.198123	-5.028459	0.290856	C	-14.468203	-8.112791	-1.465286	C	8.670658	5.763267	3.279276
C	-4.633547	-2.802731	-0.158507	C	-14.523149	-8.891207	-0.242359	H	9.075760	6.682616	-1.651392
H	-4.897141	-5.029974	0.221449	C	-12.641214	-9.549617	4.420940	C	9.840654	6.621698	1.131467
N	-3.536342	-1.979129	-0.255885	C	-14.227736	-9.949809	2.599232	C	8.371023	4.947200	4.438919
C	-5.987639	-2.392961	-0.224291	H	-15.387510	-9.323899	-2.024244	H	6.896078	3.947666	5.628207
C	-6.437237	-1.060940	-0.334625	C	-15.663661	-9.210621	0.593988	C	9.862786	5.775373	2.455364
C	-7.801560	-0.613558	-0.418677	C	-13.900366	-9.383873	4.002059	H	10.740673	6.095707	0.528040
C	-7.798946	0.762595	-0.488340	H	-12.485083	-8.751139	5.482201	H	9.186049	4.450115	4.965624
H	-8.661543	-1.270611	-0.421087	C	-15.522992	-9.524602	1.947459	H	10.779633	5.326763	2.838797
C	-6.431875	1.210138	-0.476615	H	-16.669873	-9.108208	0.185300	C	-10.071153	6.657371	-0.989197
H	-8.656952	1.419932	-0.543220	H	-14.683792	-9.508168	4.750870	C	-11.414630	6.299273	-1.452576
N	-5.649251	0.074461	-0.380217	H	-16.424442	-9.656954	2.547307	C	-9.789252	7.973220	-0.584556
C	-5.977238	2.543958	-0.537112	C	3.696162	-5.635306	-0.260757	C	-12.270785	7.377681	-1.674663
C	0.003804	-3.398989	-0.013574	C	4.300955	-5.768223	-1.591544	C	-12.070668	5.005831	-1.422544
C	0.186405	-4.374522	-1.013196	C	4.428046	-5.996466	0.884931	C	-10.774684	9.029519	-0.606987
C	0.983568	-3.297094	0.996689	C	5.509267	-6.466715	-1.626249	H	-8.798004	8.175552	-0.179740
C	1.350795	-5.146832	-1.054192	C	4.006615	-5.036453	-2.808922	C	-11.959122	8.710087	-1.270161
H	-0.568698	-4.488321	-1.789845	C	5.780603	-6.496245	0.830190	C	-13.692207	7.246023	-1.680768
C	2.158345	-4.044861	0.938324	H	3.975617	-5.810013	1.858522	H	-11.469501	4.107874	-1.287796
H	0.848411	-2.568496	1.794278	C	6.231585	-6.822655	-0.448073	C	-13.460873	4.877654	-1.427505
C	2.387096	-4.958470	-0.113183	C	6.455442	-6.325144	-2.687228	C	-10.821131	10.257703	0.162750
H	2.936748	-3.868710	1.678234	H	3.073094	-4.472603	-2.877400	C	-13.182337	9.400131	-1.029776
C	-7.041035	-3.450392	-0.162611	C	4.922068	-4.905695	-3.848355	C	-14.337213	6.029701	-1.454024
C	-7.957534	-3.500534	0.905553	C	6.811200	-6.460796	1.848781	C	-14.254742	8.495483	-1.284500
C	-7.138658	-4.425510	-1.173897	C	7.615447	-6.899429	-0.777192	H	-13.890568	3.883187	-1.299638
C	-8.935575	-4.496753	0.965588	C	6.250181	-5.472267	-3.772317	H	-9.916254	10.618083	0.659027
H	-7.897181	-2.749818	1.692425	C	7.754783	-6.592973	-2.162690	C	-12.018933	10.934634	0.396579
C	-8.115422	-4.522741	-1.113926	H	4.669904	-4.249186	-4.681555	C	-13.285795	10.451172	-0.117068
H	-6.439044	-4.395664	-2.008068	H	6.531576	-6.260994	2.883570	C	-15.709502	6.143999	-0.999461
C	-9.027522	-5.480948	-0.040612	H	8.167172	-6.531006	1.525051	C	-15.489513	8.595352	-0.642099
H	-8.172335	-6.177090	-1.898053	C	8.623660	-6.649522	0.155067	H	-12.009003	11.794269	1.068049
C	-7.027762	3.599331	-0.653070	C	7.456950	-5.057033	-4.460542	C	-14.626169	10.694593	0.377219
C	-7.872078	3.662605	-1.778721	C	8.913235	-6.020797	-2.690408	C	-16.258691	7.368689	-0.612258

H	-16.300825	5.238345	-0.858029	C	4.260709	0.722897	1.418491	C	5.755768	3.106993	-0.821572
C	-15.676992	9.810220	0.126361	C	4.135592	-0.686859	1.059936	C	5.125276	2.178884	-1.755839
H	-14.802694	11.536770	1.047684	C	4.947464	-1.466264	1.990255	C	4.234779	1.308570	-0.992575
H	-17.260880	7.375584	-0.181556	C	5.577814	-0.537229	2.923581	C	4.311489	1.701985	0.412388
H	-16.637932	9.992244	0.609420	C	5.153869	0.815013	2.570334	C	5.253878	2.810996	0.516211
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C	9.754430	-2.070050	1.626092	C	8.768520	-2.791508	-1.750671	C	7.989120	0.876023	-3.492975
C	9.124703	-2.999735	0.693216	C	7.374787	-3.179469	-1.562947	C	7.047789	-0.234762	-3.599458
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C	7.426319	1.639591	3.126763	C	7.987957	3.569471	0.150599				
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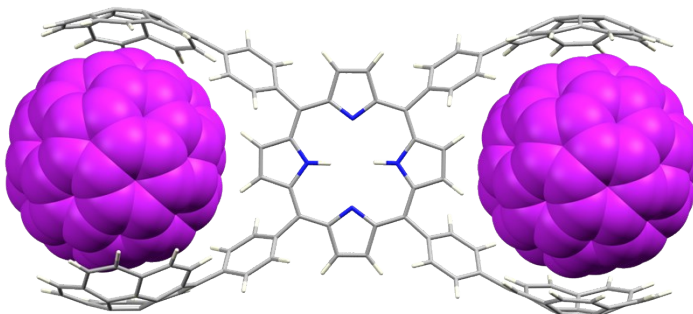
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C	-0.738834	-3.352361	-1.370165	C	8.191775	-3.714299	-1.741381	C	-5.745871	5.684191	-3.965256
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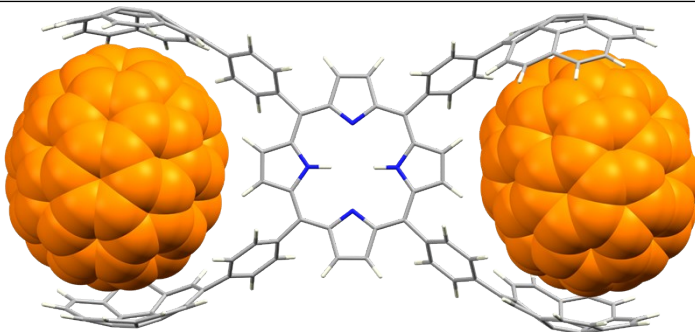
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C	-3.547943	-3.496384	-0.489632	C	4.745367	5.264321	-0.541895	H	-7.646590	5.570580	2.600879
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C	0.683286	-4.230691	-0.794268	C	6.555258	-3.567354	-2.111549	C	-13.477442	6.291863	0.161178
H	-1.346196	-5.063363	-1.038923	H	7.151185	5.723749	0.017293	C	-12.166648	5.762282	-3.402820
C	1.083252	-2.874381	-0.417649	C	8.073486	6.186279	1.060965	H	-10.757119	5.042536	-4.846698
H	1.357796	-5.041172	-1.046000	C	7.558832	5.723595	-1.334076	C	-13.561460	6.156201	-1.225530
N	-0.016834	-2.076114	-0.208037	C	9.252109	6.762820	0.584125	H	-14.354718	6.049954	0.761939
C	-2.479747	2.409040	0.471293	C	8.082523	5.947947	2.496481	H	-13.010568	5.390300	-3.984552
C	-1.133287	2.835987	0.542407	C	8.862702	6.149553	-1.774210	H	-14.500582	5.810056	-1.658056
C	-0.733771	4.203550	0.876047	H	6.903054	5.231703	-2.051651	C	-7.225725	-5.739923	-0.198155
C	0.632029	4.218213	0.862609	C	9.637072	6.755664	-0.787779	C	-7.774273	-5.883358	1.156816
H	-1.408620	5.018971	1.110336	C	10.443692	6.846806	1.366996	C	-8.007916	-6.079766	-1.316725
C	1.054087	2.857155	0.531660	H	7.168133	5.642099	2.999064	C	-8.989391	-6.565821	1.232444
C	1.294852	5.047257	1.083734	C	9.250834	6.015153	3.256751	C	-7.413233	-5.170312	2.369403
N	-0.032586	2.034705	0.346696	C	9.591382	5.781235	-2.975139	C	-9.361709	-6.568575	-1.211406
C	2.408697	2.452935	0.478849	C	11.058695	6.820179	-0.855434	H	-7.594174	-5.883029	-2.305634
C	2.851161	1.123215	0.296811	C	10.530048	6.357561	2.668888	C	-9.764131	-6.900671	0.081237
C	4.214040	0.668027	0.276335	C	11.557584	6.874429	0.478280	C	-9.888409	-6.423169	2.333010
C	4.206656	-0.696387	0.120971	H	9.202018	5.723038	4.306635	H	-6.477211	-4.617192	2.403776
C	5.081966	1.301260	0.373962	H	9.049171	5.344400	-3.814246	C	-8.288895	-5.039198	3.444742
C	2.846527	-1.129648	-0.032644	C	10.983700	5.841776	-3.040070	C	-10.434971	-6.508073	-2.185334
H	5.065508	-1.347652	0.109637	C	11.787581	6.273849	-1.911586	C	-11.134548	-6.960832	0.467030
N	2.060886	0.000728	0.110473	C	11.875984	6.054485	3.114077	C	-9.626468	-5.590431	3.420871
C	2.428762	-2.449944	-0.317854	C	12.815673	6.388008	0.833002	C	-11.212273	-6.668160	1.859998
C	-3.590040	3.395855	0.625633	H	11.480076	5.447760	-3.927370	H	-7.993142	-4.397352	4.277712
C	-3.746269	4.469159	-0.273396	C	13.170010	5.969446	-1.595691	H	-10.195920	-6.304888	-3.229522
C	-4.602618	3.187549	1.586173	C	12.964072	6.066692	2.239134	C	-11.776918	-6.561896	-1.806587
C	-4.912033	5.240157	-0.271688	H	12.033829	5.706679	4.135347	C	-12.177552	-6.684236	-0.418620
H	-2.967292	4.664696	-1.009041	C	13.659597	6.024471	-0.289641	C	-10.797524	-5.180290	4.170181
C	-5.780420	3.932478	1.564239	H	13.821742	5.580450	-2.378385	C	-12.340118	-6.092551	2.446305
H	-4.489124	2.381539	2.309151	H	13.931255	5.724363	2.608070	H	-12.539409	-6.394614	-2.568102
C	-5.978891	4.951741	0.607764	H	14.675920	5.677836	-0.101081	C	-13.419671	-6.286766	0.213688
H	-6.582671	3.676019	2.253528	C	-7.284767	5.639836	0.487726	C	-12.091236	-5.419692	3.075553
C	3.496702	3.473193	0.547948	C	-7.838523	5.926474	-0.841419	H	-10.669991	-4.569847	5.064550
C	4.542214	3.369566	1.486833	C	-8.061923	5.866939	1.638124	C	-13.497652	-6.005908	1.578688

H	-14.294822	-6.094108	-0.407519	C	-8.608411	1.212632	2.202631	C	10.344008	-0.701452	3.686954
H	-12.929154	-4.987407	4.253221	C	-9.287577	0.184153	2.985193	C	7.685791	1.205835	1.163772
H	-14.431130	-5.605504	1.975092	C	-8.850411	-1.119887	2.493474	C	14.131800	-1.207831	-0.696202
C	7.247462	-5.677392	-0.470422	C	-9.139209	2.947349	0.507943	C	10.742984	-1.960371	3.210097
C	8.104156	-5.949182	-1.631081	C	-10.430563	3.490790	0.097005	C	12.587727	-1.297757	-2.635832
C	7.728676	-5.921788	0.834463	C	-11.379881	3.264017	1.182846	C	14.311806	1.059174	0.294377
C	9.298058	-6.608202	-1.334204	C	-10.674322	2.587247	2.266709	C	10.578582	3.294611	1.561063
C	8.043554	-5.450589	-2.997280	C	-9.288043	2.390900	1.849251	C	13.051061	-1.123460	2.855011
C	9.045961	-6.433899	1.115400	C	-9.995538	2.691622	-2.209100	C	12.570848	-2.976366	-0.810798
H	7.119213	-5.566547	1.664828	C	-10.848925	1.931701	-3.115398	C	14.167350	0.525284	1.584874
C	9.755094	-6.853670	-0.007965	C	-12.234619	2.128324	-2.698029	C	7.648811	-0.525623	-1.117486
C	10.446046	-6.555858	-2.183636	C	-12.233459	3.010891	-1.536096	C	7.512120	-1.059420	0.173336
H	7.109206	-5.056918	-3.389116	C	-10.849821	3.360992	-1.234401	C	10.289138	-3.507950	-0.002515
C	9.171491	-5.377346	-3.816242	C	-13.146782	1.063923	-2.772099	C	14.294420	0.175648	-0.868124
C	9.843924	-6.316401	2.322692	C	-12.710078	-0.239306	-3.264311	C	12.756253	0.141426	-2.813511
C	11.176422	-6.944951	-0.032444	C	-13.389100	-1.267262	-2.481352	C	11.469874	0.700025	-3.218280
C	10.473853	-5.830872	-3.372856	C	-14.245010	-0.599625	-1.505676	C	13.015023	-2.854870	0.574086
C	11.603962	-6.758042	-1.378680	C	-14.095314	0.841002	-1.685194	C	11.070348	1.959520	-2.742045
H	9.072097	-4.894658	-4.789081	C	-12.710753	-2.445418	-2.129871	C	8.634928	-2.177033	1.922121
H	9.356781	-6.604207	3.258787	C	-11.324493	-2.641943	-2.547363	C	12.810236	2.942520	0.875094
C	11.236634	-6.406797	2.298293	C	-10.619288	-2.318940	-1.463767	C	7.834353	1.763076	-0.178435
C	11.971790	-6.622078	1.066116	C	-11.567542	-3.538815	-0.377069	C	13.979863	-1.762944	0.645823
C	11.799273	-5.455011	-3.824148	C	-12.860593	-3.001628	-0.788721	C	11.308390	0.392317	3.758244
C	12.851941	-6.227504	-1.704744	C	-9.288140	-2.983284	-1.174860	C	9.057746	-0.143141	3.282451
H	11.785817	-6.199642	3.217139	C	-8.608923	-1.955508	-1.958416	C	12.105201	-3.056542	1.622749
C	13.341886	-6.282940	0.736339	C	-7.754928	-1.191104	-1.053052	C	9.226073	1.296388	3.104077
C	12.931599	-5.642355	-3.028959	C	-7.903149	-1.749477	0.288841	C	10.505472	-0.393735	-3.290708
H	11.908728	-4.919171	-4.767492	C	-8.852863	-2.854635	0.211991	C	11.940329	2.707150	-1.839817
C	13.760737	-6.095278	-0.582753	C	-7.612805	0.193941	-1.225528	C	11.523534	3.500646	0.470895
H	14.044221	-6.060255	1.540026	C	-8.316553	0.872492	-2.309193	C	8.797796	2.856034	-0.105925
H	13.884036	-5.242018	-3.378058	C	-8.753077	2.174590	-1.815782	C	11.279408	2.606993	2.641551
H	14.774257	-5.733224	-0.758372	C	-8.316619	2.304533	-0.429212	C	10.534740	-2.607511	-2.172423
C	-10.620418	0.373568	3.385231	C	-7.617190	1.078175	-0.063534	C	9.004795	-2.945742	-0.406420
C	-11.326441	1.597422	3.019216	C	-9.137492	0.138215	-3.180203	C	9.241498	2.978729	1.278674
C	-12.712286	1.246801	2.719187	C	-9.287007	-1.302864	-3.000827	C	13.997073	-0.913137	1.763614
C	-12.862659	-0.193238	2.898553	C	-10.672252	-1.653121	-3.301253	C	7.815334	0.913544	-1.296551
C	-11.569469	-0.733082	3.309473	C	-11.378244	-0.428376	-3.665932	C	11.196545	-1.629365	-2.930558
C	-13.146242	-2.798569	-0.492708	C	-10.429524	0.678629	-3.591170	C	13.325292	1.204009	2.565023
C	-14.094494	1.691673	-0.568138	C	-1.100057	-0.049028	0.083227	C	8.552323	1.958938	2.065208
C	-14.244940	1.135060	0.773609	H	1.046765	0.018585	0.054444	C	13.623152	2.294805	-0.066606
C	-13.389515	1.898769	1.676449	C	9.689495	2.175193	-2.317508	C	10.715469	-3.386028	1.327199
C	-12.711085	2.927922	0.894383	C	10.617367	1.627690	3.398448	C	9.872868	-2.711980	2.309896
C	-14.388883	-1.135389	-0.215891	C	12.123432	-2.175016	2.785243	C	9.177997	-0.185608	-2.884211
C	-13.685183	-2.360675	0.149240	C	11.097552	3.382861	-0.857801	C	8.193671	-2.295508	0.534704
C	-13.249837	-2.233259	1.537770	C	12.635437	0.184967	3.350727	C	11.235098	-3.294486	-1.092669
C	-13.684523	-0.928290	2.029150	C	13.261619	-1.959805	-1.596883	C	8.221468	-0.866514	2.417131
C	-14.389264	-0.250793	0.945746	C	8.488507	-1.205397	-2.098098				
C	-11.149222	-3.415596	0.954057	C	9.708875	3.054831	-1.153125				
C	-9.767106	-3.062876	1.253157	C	9.154000	-2.392100	-1.748569				
C	-9.765560	-2.180984	2.415218	C	13.593117	0.864027	-1.947621				
C	-11.150285	-2.834635	2.833542	C	8.761629	1.123939	-2.388746				
C	-12.005249	-2.746680	1.928654	C	13.178572	2.173368	-1.453299				
C	-7.900543	-0.890304	1.406470	C	7.525517	-0.176958	1.336044				
C	-7.753536	0.543088	1.226562	C	12.660782	2.390754	2.217491				

(C₇₀)₂@2H-PTetraCor (E= -10309.9502828 a.u.)



C	-4.870274	-5.216258	0.602796	C	4.229413	-0.613018	0.092918	H	-4.972563	-6.015275	1.335200
C	-3.756879	-4.373454	0.657644	H	5.103501	1.403281	0.149984	H	6.394606	-4.214312	-1.548820
C	-5.875825	-5.038670	-0.372879	C	2.859442	-1.050907	0.083696	C	7.141978	5.994896	0.659367
C	-3.588628	-3.334348	-0.279221	C	5.089379	-1.265303	0.042039	C	7.971556	6.273829	1.840470
H	-3.021813	-4.497421	1.451539	N	2.082178	0.090771	0.179028	C	7.627733	6.308314	-0.628888
C	-5.672295	-4.033703	-1.343553	C	2.397732	-2.382316	-0.002773	C	9.139704	6.992526	1.578629
C	-4.546301	-3.213660	-1.307972	C	-3.506160	3.642519	0.389770	C	7.913855	5.744401	3.196274
H	-6.429313	-3.859170	-2.104579	C	-3.590419	4.635860	-0.606165	C	8.920850	6.894335	-0.876094
H	-4.423441	-2.435111	-2.058945	C	-4.532622	3.582770	1.354124	H	7.053580	5.945609	-1.480546
C	-2.494742	-2.328958	-0.139543	C	-4.703491	5.476653	-0.686189	C	9.598839	7.306822	0.267811
C	-2.920413	-0.987670	-0.016564	H	-2.794831	4.714388	-1.345796	C	10.280590	6.954408	2.439046
C	-4.278676	-0.517497	-0.002212	C	-5.651834	4.409527	1.263184	H	6.987652	5.320641	3.574753
C	-4.258011	0.854459	0.086627	H	-4.474989	2.838270	2.146511	C	9.034215	5.687799	4.025233
H	-5.151379	-1.150837	-0.052619	C	-5.781131	5.347526	0.216905	C	9.737489	6.840604	-2.076758
C	-2.888465	1.278788	0.183678	H	-6.465833	4.280827	1.973222	C	11.017525	7.451706	0.311413
H	-5.109853	1.516979	0.090548	C	3.557218	3.548905	0.609764	C	10.324726	6.195929	3.606540
N	-2.117413	0.132289	0.107850	C	4.559516	3.344968	1.579291	C	11.437872	7.227097	1.654506
C	-1.145829	-2.758763	-0.128517	C	3.696130	4.659699	-0.247691	H	8.939660	5.183914	4.987763
C	-0.739046	-4.147152	-0.350376	C	5.698328	4.147367	1.625529	H	9.271603	6.571692	-3.025039
C	0.624896	-4.161152	-0.294858	H	4.466854	2.510455	2.271907	C	11.124059	6.984621	-2.033826
H	-1.406526	-4.977873	-0.547741	C	4.818351	5.488454	-0.173664	C	11.836771	7.195643	-0.786914
C	1.041066	-2.779483	-0.053309	H	2.931231	4.851004	-0.998947	C	11.658131	5.859134	4.063129
H	1.291323	-5.004381	-0.436286	C	5.871850	5.227977	0.733235	C	12.700816	6.730476	1.977227
N	-0.048877	-1.945283	0.041153	H	4.896422	6.339911	-0.848483	H	11.691205	6.823474	-2.951097
C	-2.432743	2.605378	0.344878	C	3.464971	-3.427078	0.012016	C	13.215694	6.894334	-0.452652
C	-1.078021	3.002030	0.433758	C	4.477711	-3.457580	-0.965446	C	12.785819	6.110775	3.285181
C	-0.662701	4.368794	0.749549	C	3.555649	-4.334292	1.086585	H	11.778559	5.302824	4.993265
C	0.701968	4.354584	0.790105	C	5.590193	-4.289329	-0.824351	C	13.626850	6.670636	0.862553
H	-1.329033	5.202042	0.941675	H	4.418894	-2.779120	-1.815167	C	12.932820	6.724695	-1.256276
H	1.108513	2.981358	0.490647	C	4.660521	-5.177561	1.219449	H	13.753401	5.737207	3.632557
H	1.369733	5.174756	1.029184	H	2.771140	-4.341567	1.842098	H	14.650657	6.338459	1.037439
N	0.011752	2.175380	0.290856	C	5.726045	-5.139996	0.292896	C	-7.040455	6.105800	0.030093
C	2.457559	2.554635	0.446404	H	4.720919	-5.856109	2.069585	C	-7.607309	6.246095	-1.315726
C	2.879352	1.218105	0.260877	H	-4.755310	6.226263	-1.474437	C	-7.775696	6.523886	1.153647
C	4.237628	0.760051	0.164443	H	6.488495	3.879919	2.317217	C	-8.784744	6.993104	-1.388603

C	-7.311769	5.479359	-2.512774	C	12.525206	-7.087135	-2.014350	H	1.066430	0.097409	0.210827
C	-9.101512	7.085684	1.061510	H	11.472414	-6.857460	-3.866212	C	12.699975	2.159445	-2.984460
H	-7.354539	6.327612	2.139429	C	13.438503	-6.814622	0.430132	C	12.256108	-3.098415	1.646518
C	-9.514749	7.401709	-0.232303	H	13.826583	-6.143441	2.427173	C	12.924925	3.014673	1.442763
C	-9.715674	6.865105	-2.465379	H	13.485213	-6.898685	-2.495701	C	9.428614	3.545489	0.833133
H	-6.405651	4.879027	-2.551322	H	14.463226	-6.602426	0.124338	C	9.129857	-3.537111	-0.988237
C	-8.216628	5.364563	-3.568444	C	-14.353148	-2.388904	-0.554662	C	7.859111	2.116094	-0.446005
C	-10.151378	7.113856	2.060636	C	-14.518431	-1.868910	0.794739	C	7.638924	-0.227507	-1.329927
C	-10.889138	7.521694	-0.590079	C	-13.390905	-3.387710	-0.814608	C	8.335848	1.526887	1.769623
C	-9.522624	5.983311	-3.529344	C	-13.731276	-2.368762	1.837019	C	8.438263	-2.559607	-1.816522
C	-11.013818	7.190871	-1.971430	C	-13.186667	-1.457248	2.830182	C	12.338281	3.729049	-1.259043
H	-7.974011	4.682957	-4.384190	C	-13.504216	-0.066066	2.821118	C	11.817316	3.215276	-2.516063
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H	6.757292	-6.116796	-2.381944	C	-9.163770	3.667925	0.352910	C	9.521511	3.492783	-1.644588
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C	11.206100	-7.522668	-0.064956	C	-12.678365	3.648975	1.076417	C	8.651107	1.578683	-2.719545
H	8.651926	-6.572461	-3.819457	C	-13.230092	3.314512	-0.228261	C	14.632688	-1.016842	-0.009283
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C	11.040695	-6.026705	3.327364	C	-11.336078	4.045939	1.192727				
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C	11.371156	-7.061682	-2.799949	C	-12.461583	2.207369	3.075767				
C	12.469745	-7.218981	-0.571249	C	-13.246102	2.707400	2.031513				
H	11.647270	-5.583450	4.117692	C	-10.517269	3.524650	2.281271				
C	13.071248	-6.552265	1.751145	H	-1.101563	0.111559	0.129629				

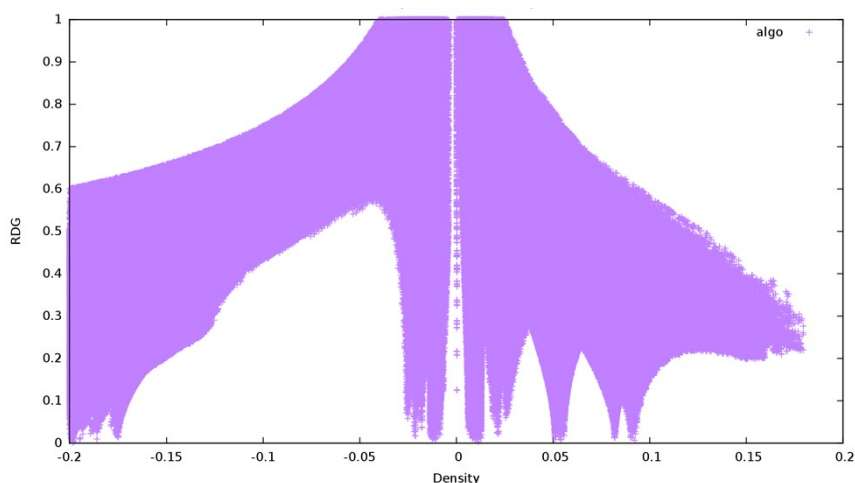


Figure S73: Plot of the reduced density gradient versus the electron density for the complex $C_{60}@2H\text{-}PTetraCor$.

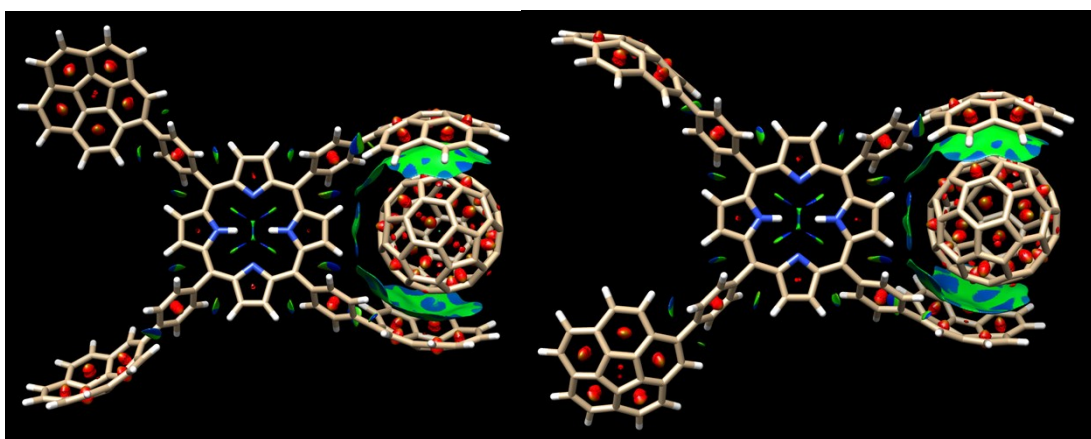


Figure S74: Non-covalent interaction isosurfaces for the assembly $C_{60}@2H\text{-PTetraCor}$.

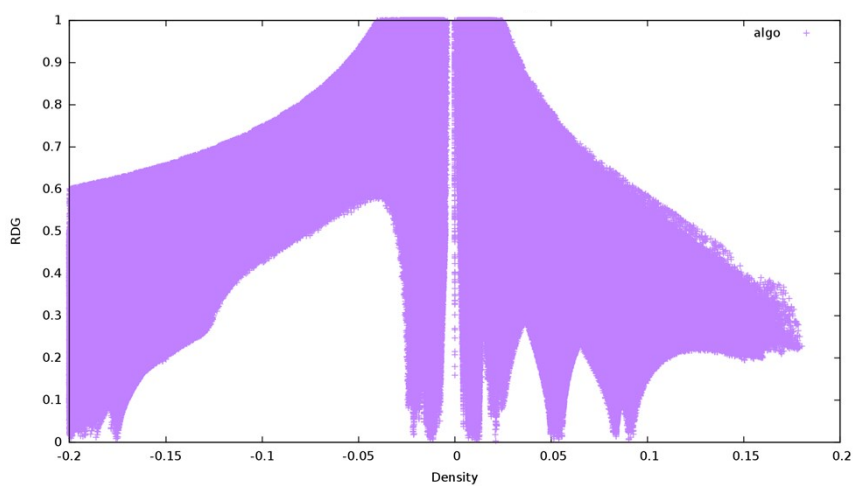


Figure S75: Plot of the reduced density gradient versus the electron density for the complex $C_{70}@2H\text{-PTetraCor}$.

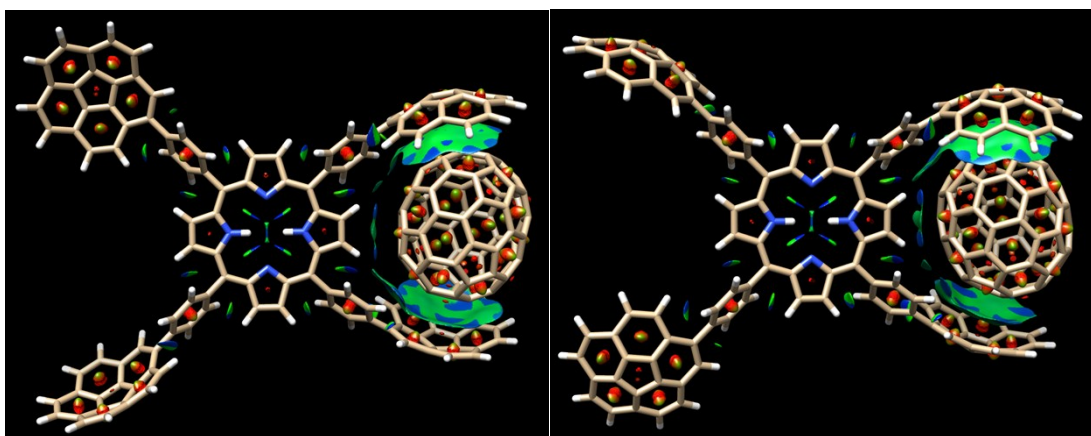


Figure S76: Non-covalent interaction isosurfaces for the assembly $C_{70}@2H\text{-PTetraCor}$

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