

SUPPORTING INFORMATION

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Title: Synthetic, Structural, and Spectroscopic Studies of Bis(porphyrinzinc) Complexes Linked by Two-Atom Conjugating Bridges

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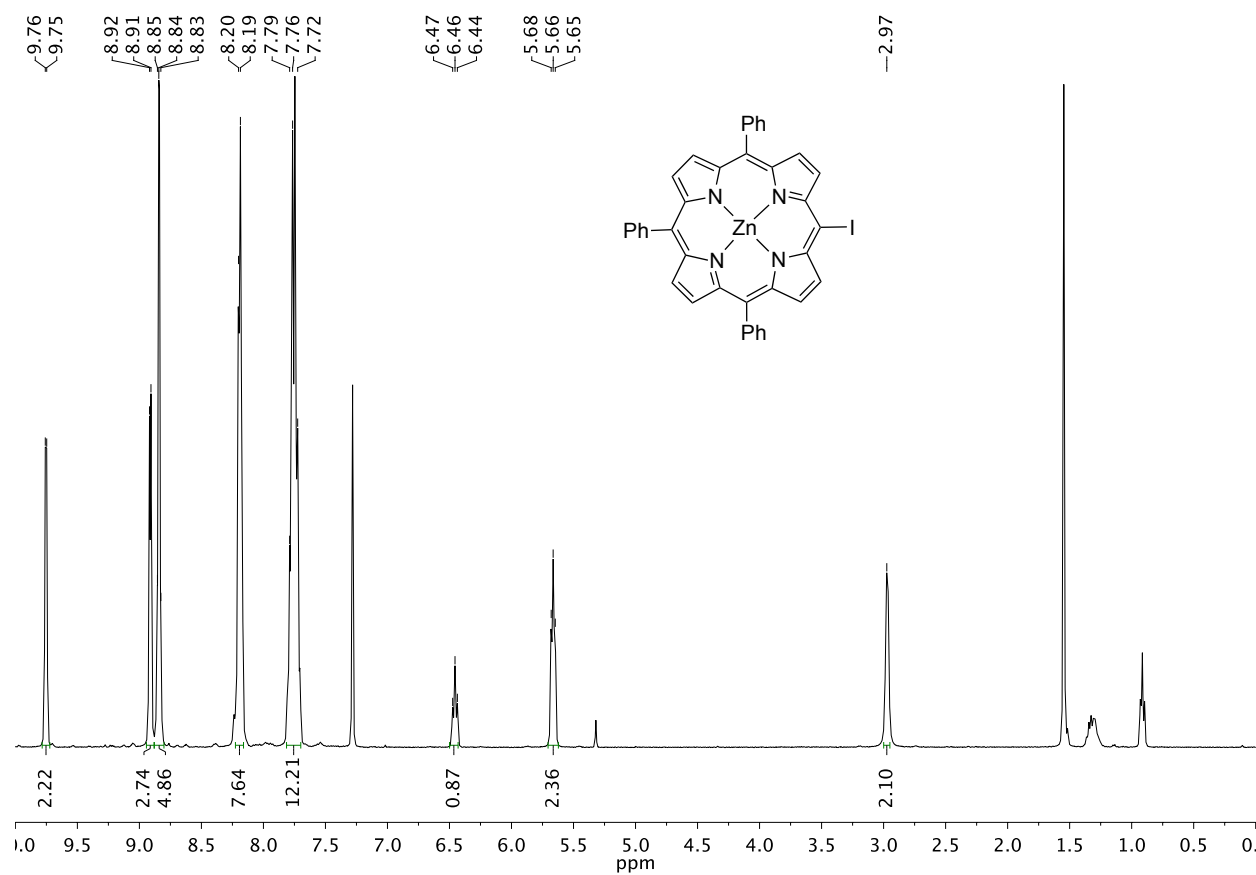
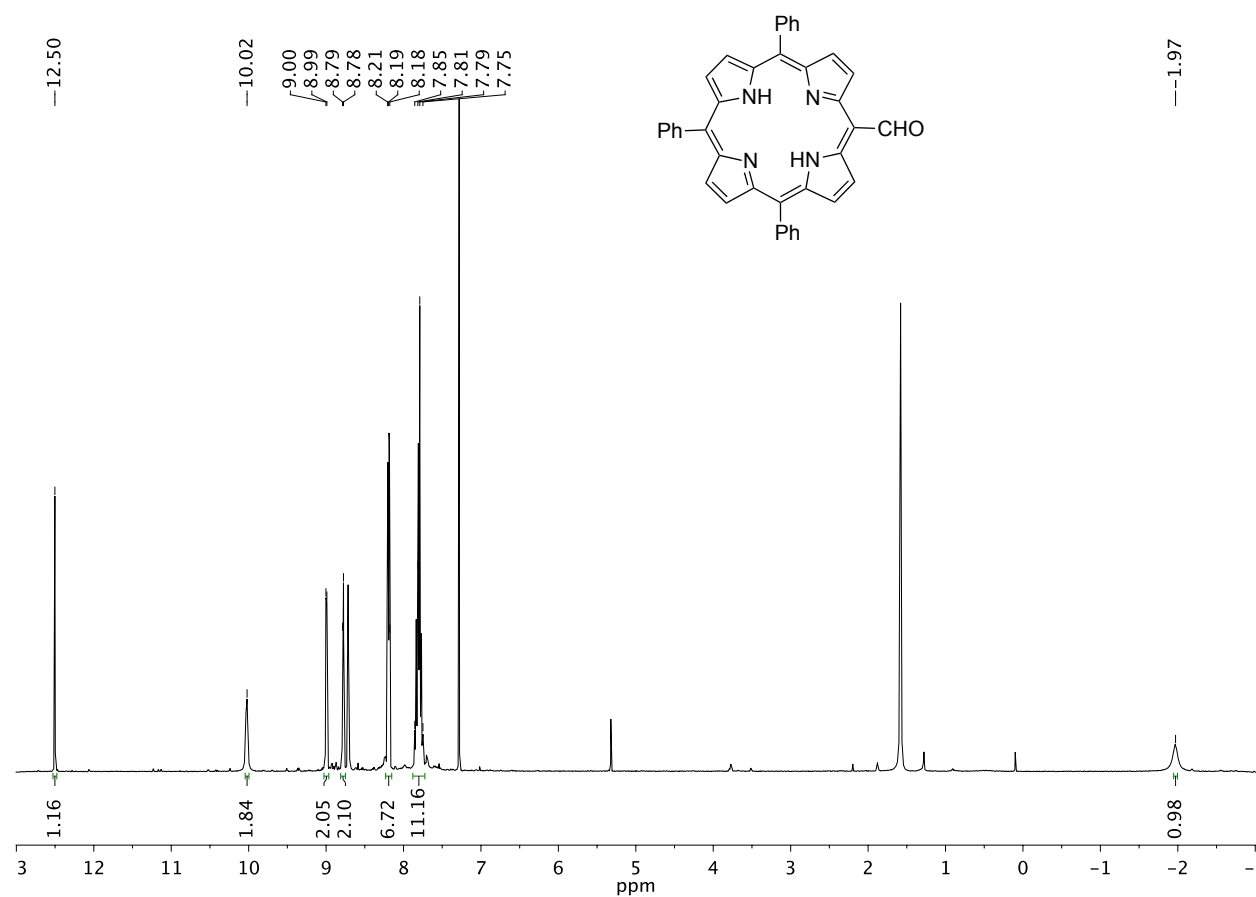
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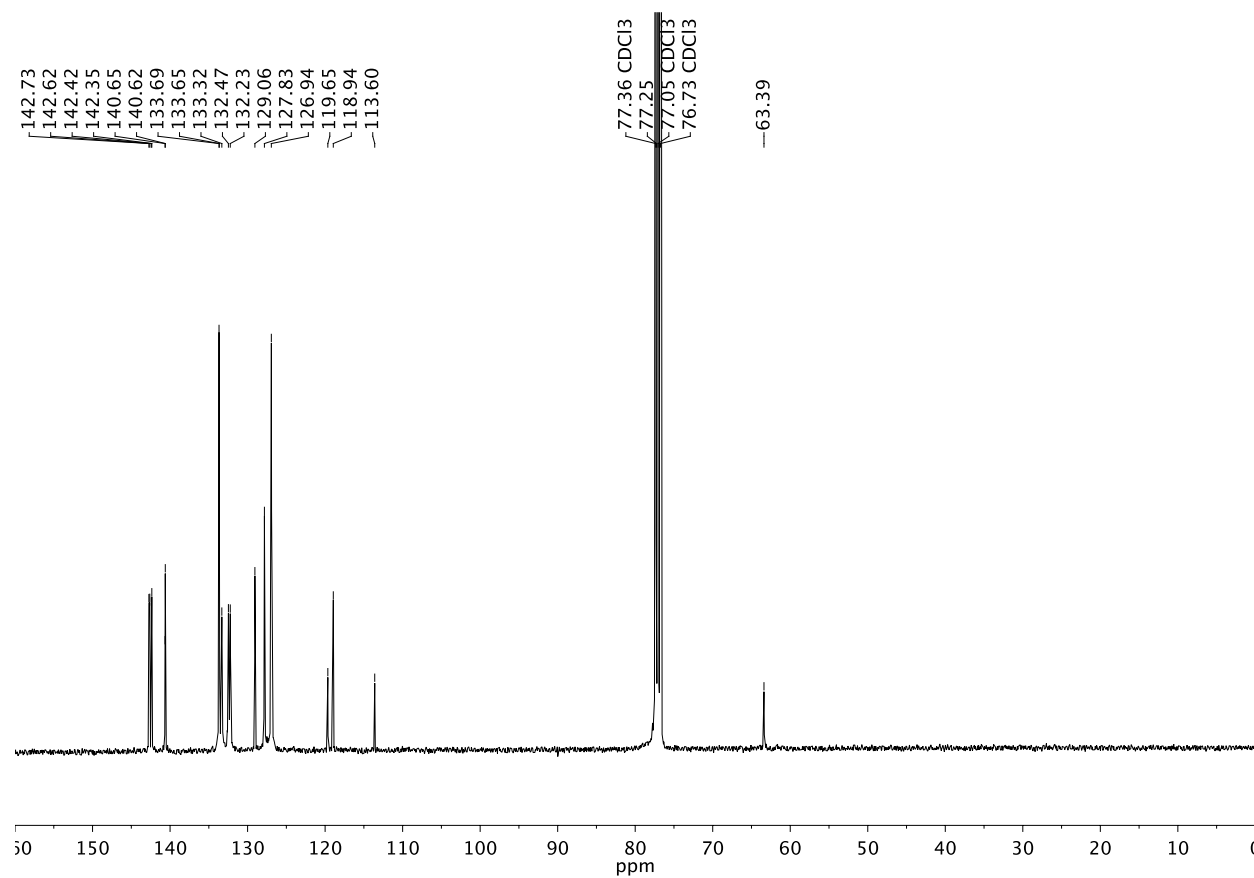
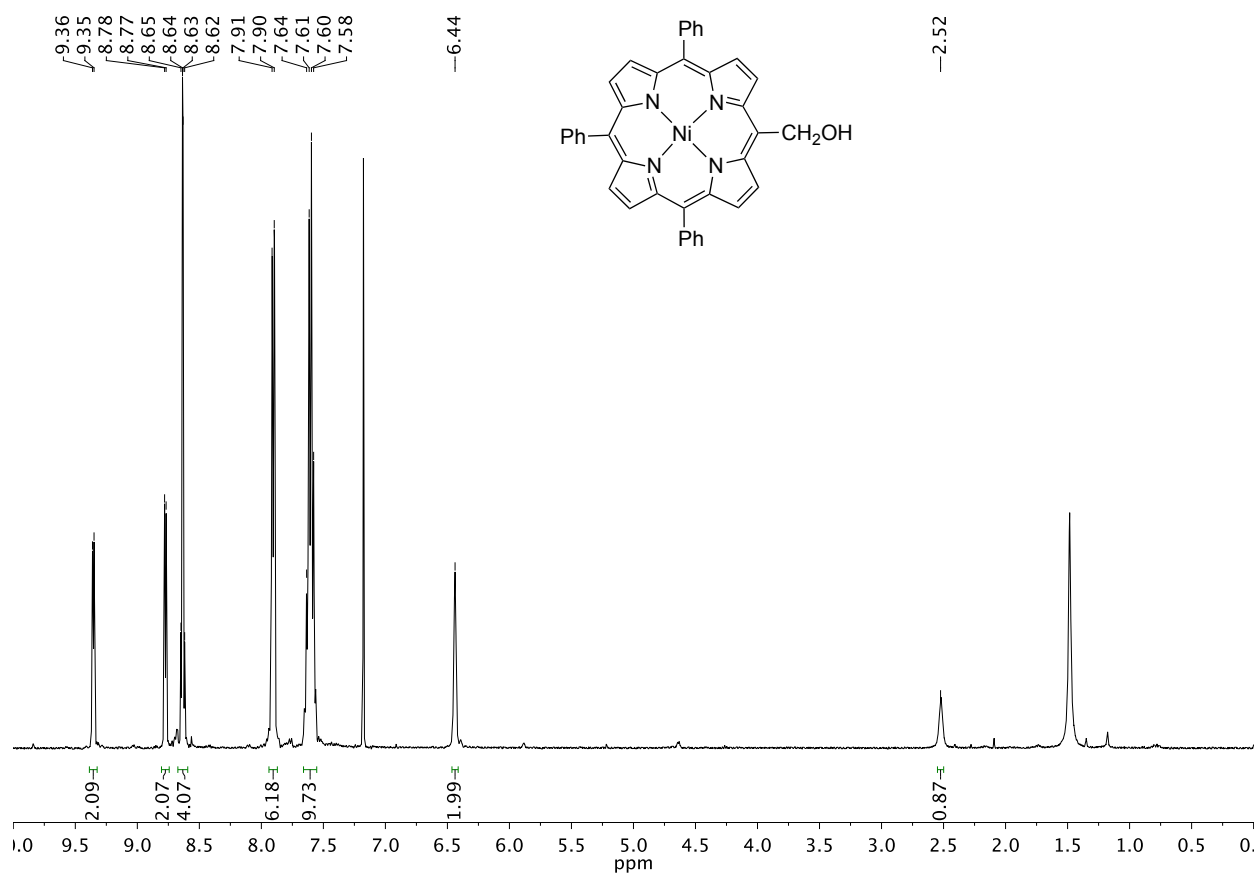
NMR Spectra of novel porphyrins in CDCl_3 (resonances for residual CHCl_3 and water at ca. 1.5 ppm present in all spectra; CH_2Cl_2 residues present in some samples, other impurities noted on spectra) and high resolution mass spectra for novel diporphyrins. S2-S19

Figure S1. The hydrogen bonded cyclic hexamer in the crystal structure of the zinc formylporphyrin **23**. S20

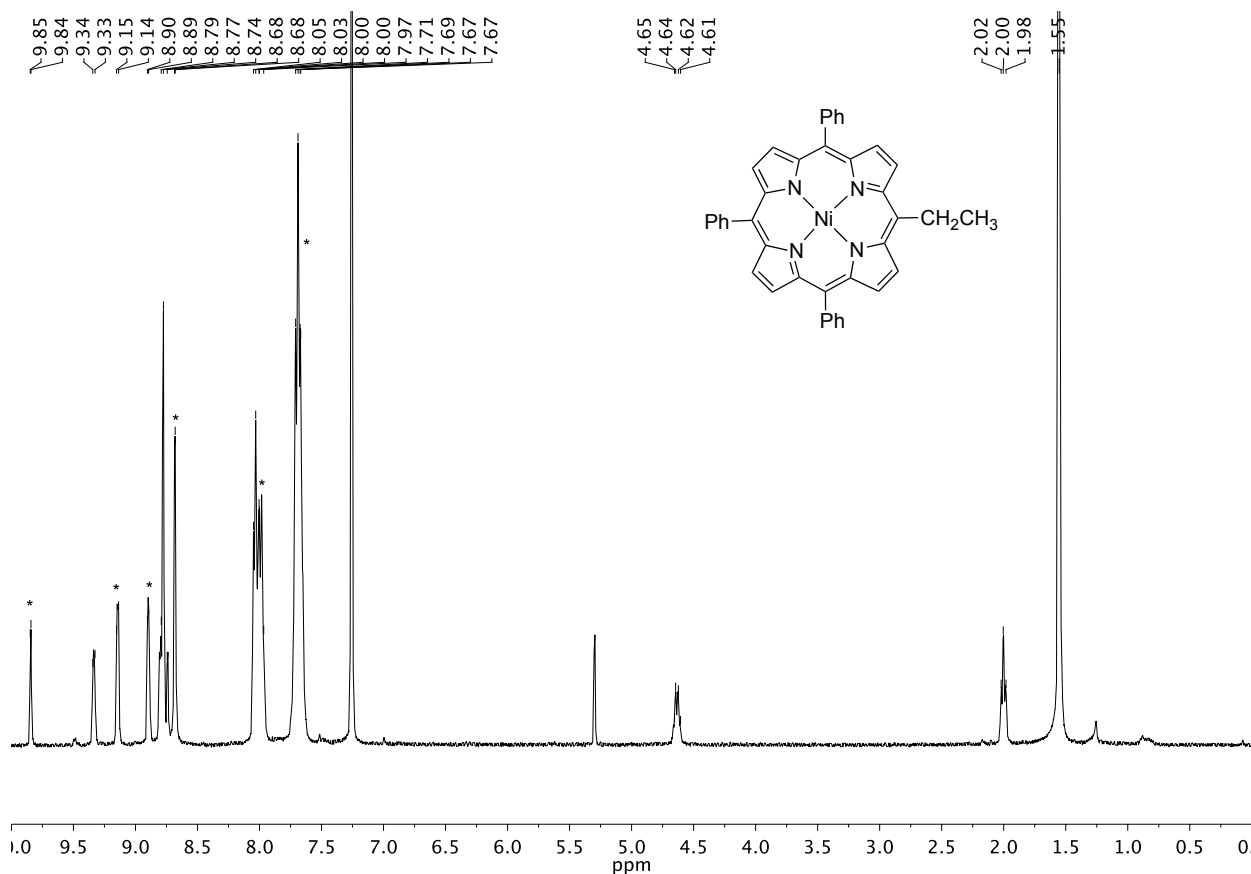
Figure S2. Effects of solvents on the Soret band region of the electronic absorption spectrum of imine diporphyrin **4**: (a) the Soret band region; (b) the Q band region. S21

Figure S3. Fluorescence emission spectra in toluene: (a) ethyne **3** excited at 414 nm (solid), 482 nm (dots), and absorption spectrum scaled for convenience (dashes); (b) imine **4** excited at 427 nm (solid), 449 nm (dots), and absorption spectrum scaled for convenience (dashes). S22

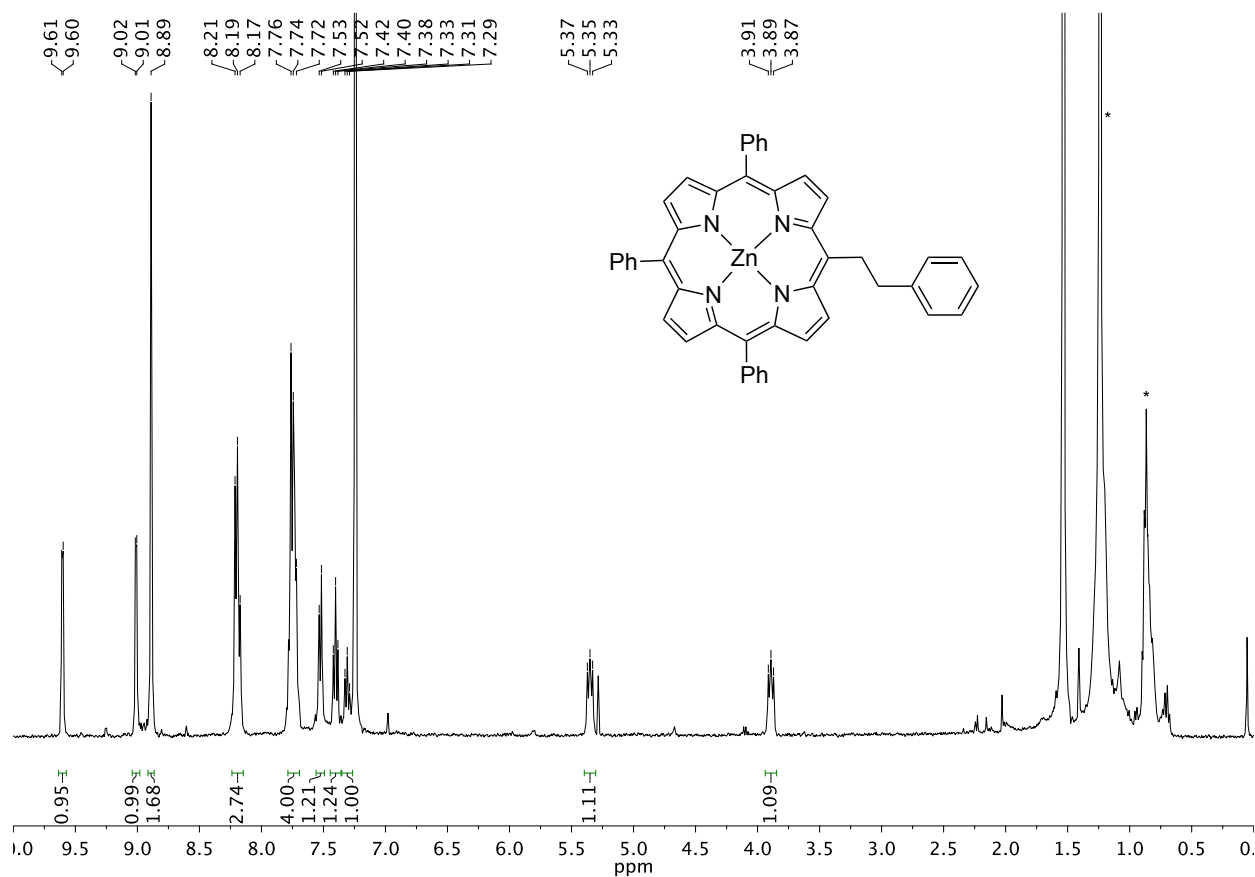
ZnTriPPI 24**H₂TriPPCHO 33**

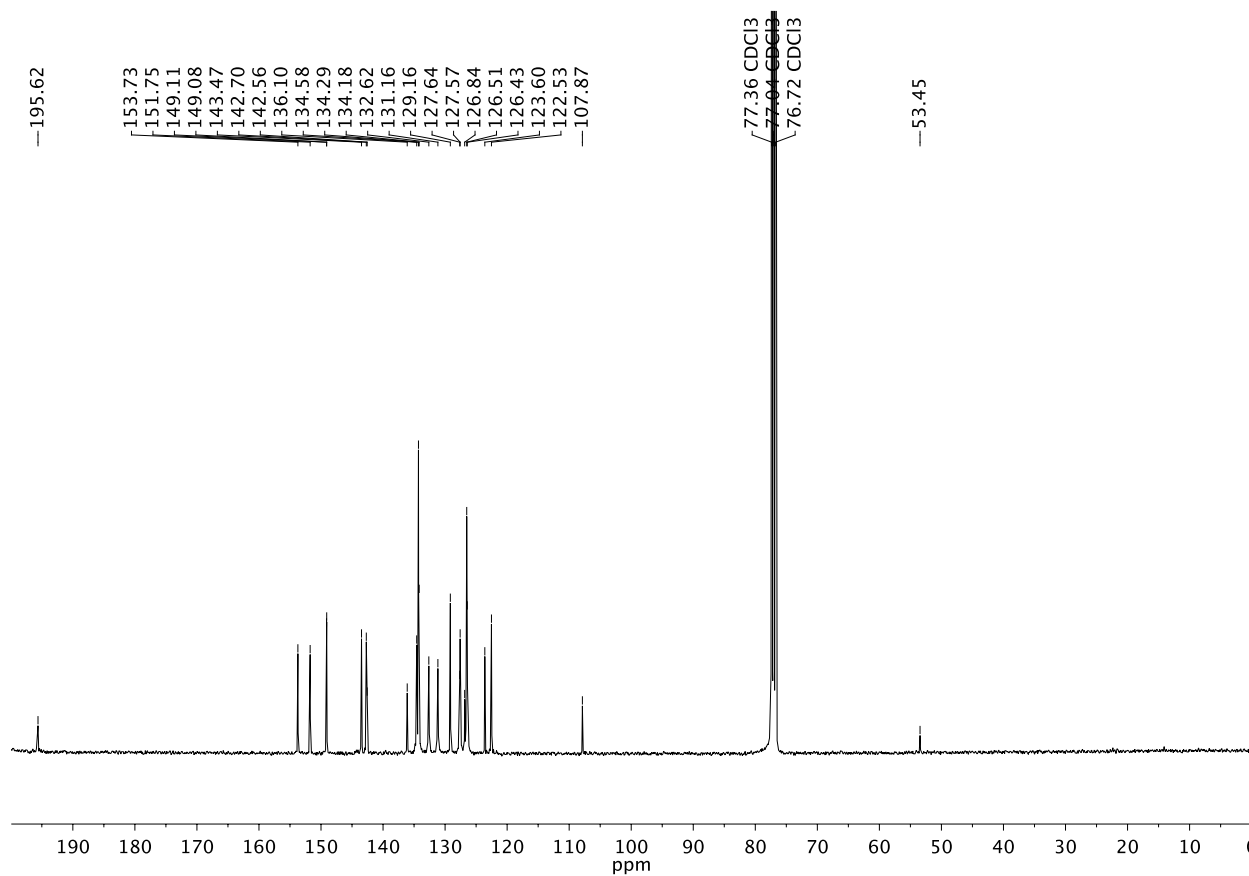
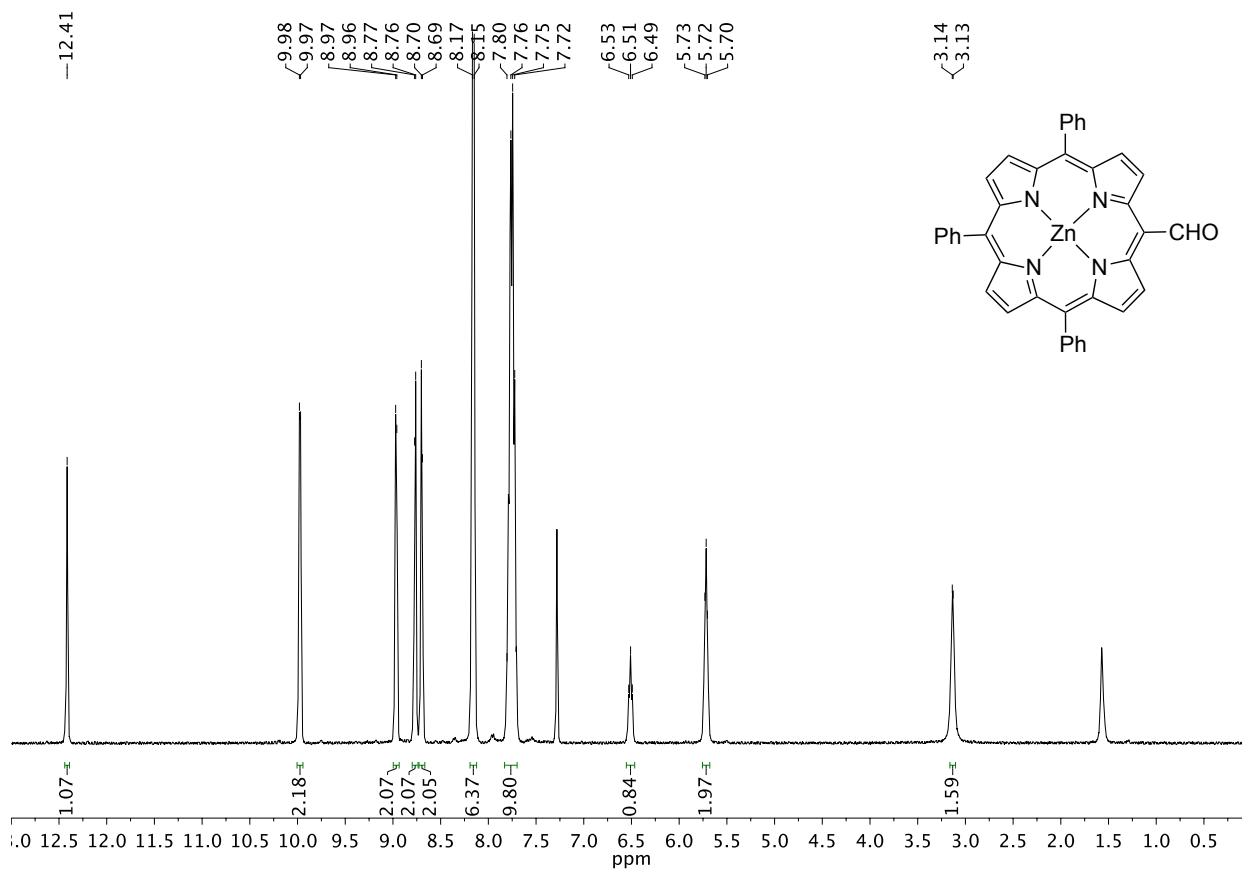
NiTriPPCH₂OH 8

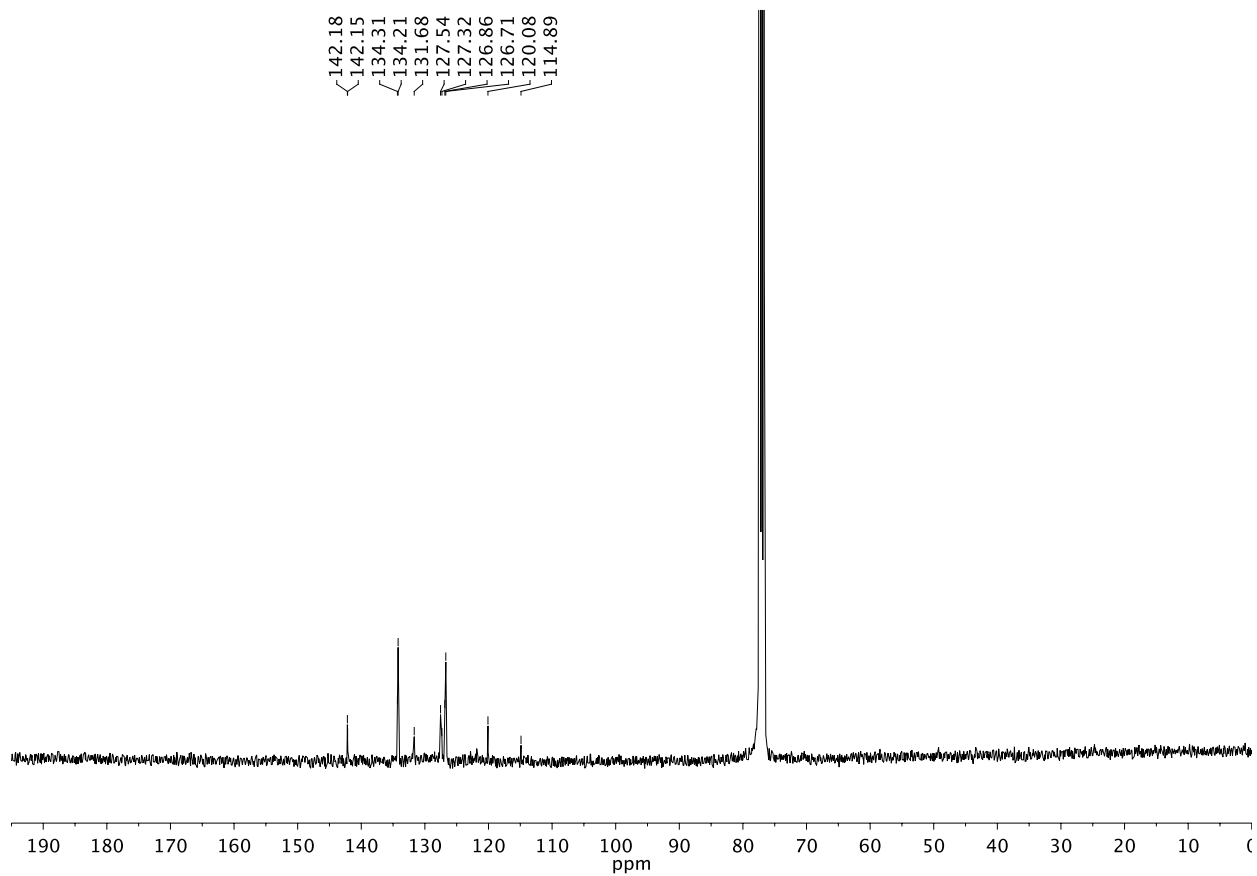
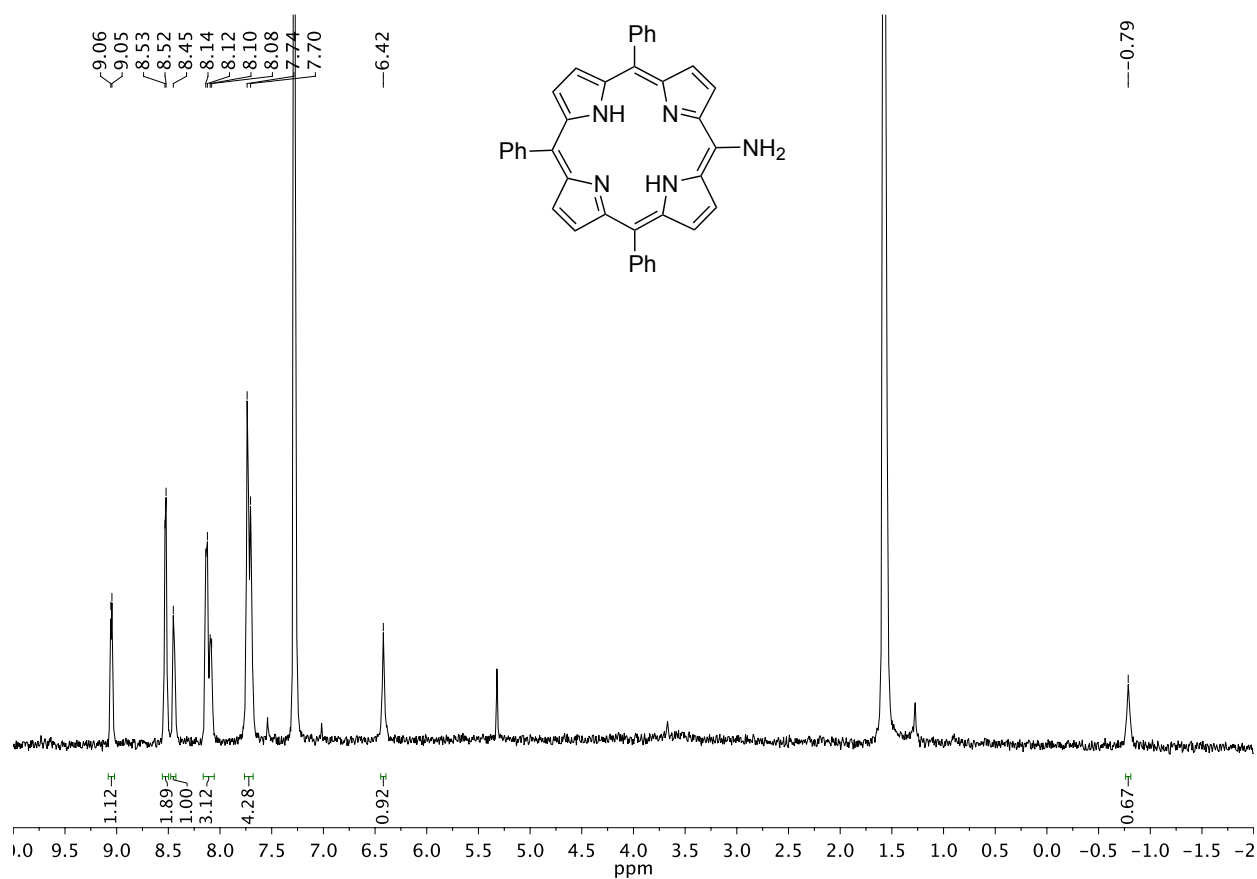
NiTriPPCH₂CH₃ 17 (* = NiTriPP 6)



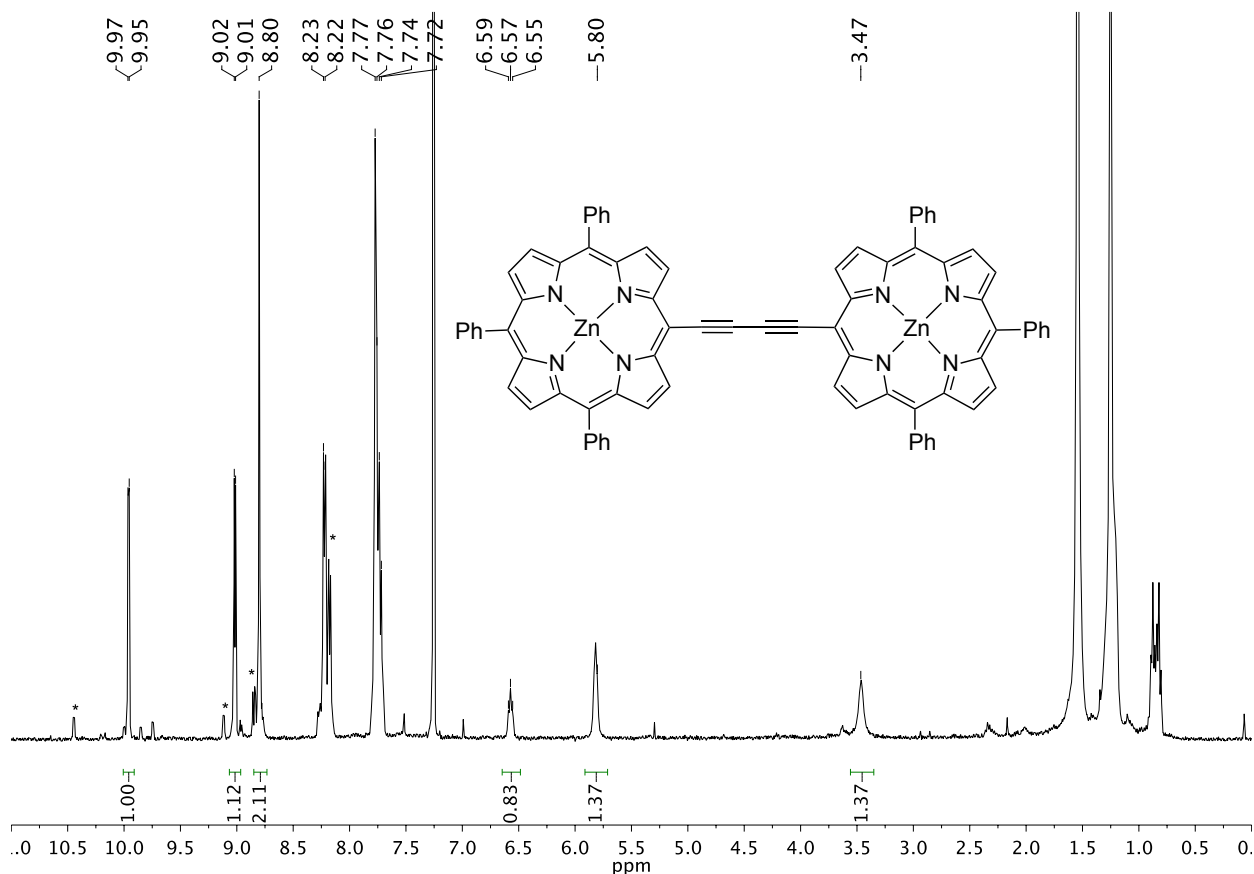
ZnTriPPCH₂CH₂Ph 18 (* = aliphatic hydrocarbon residues)



ZnTriPPCHO 23 (pyridine coordinated)

H₂TriPPNH₂ 30

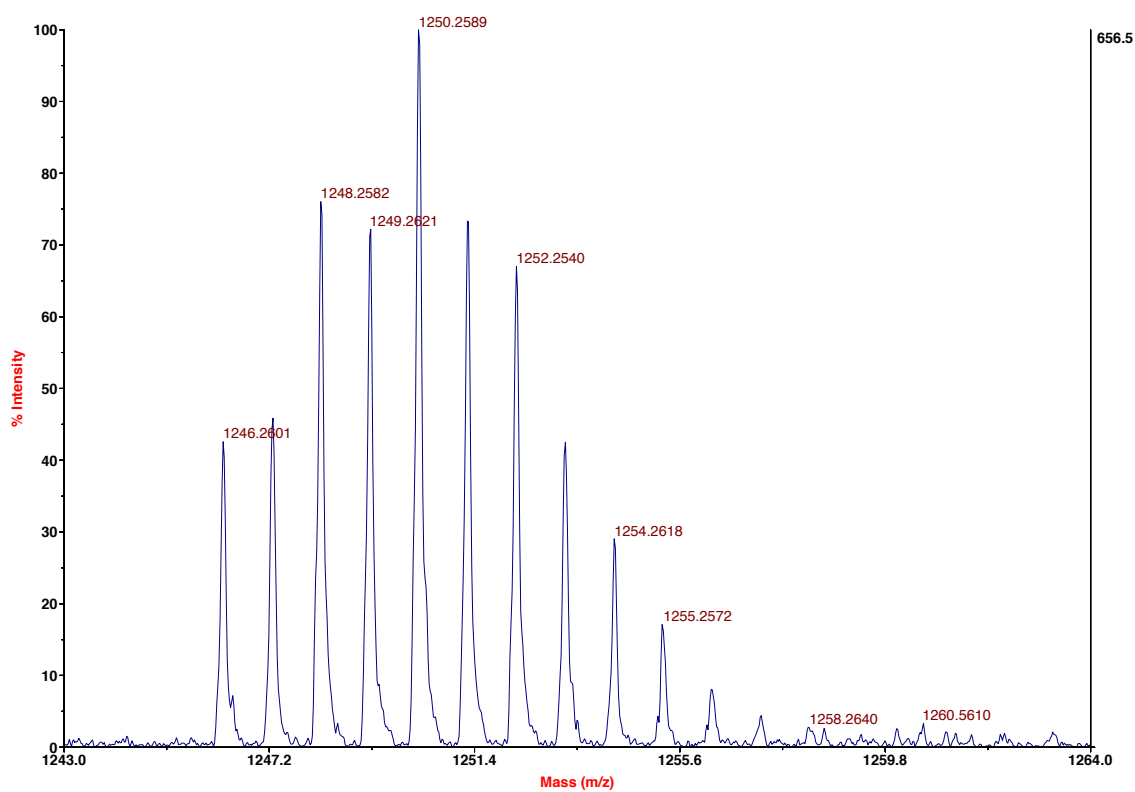
ZnTriPP-C₄-ZnTriPP 26 (pyridine coordinated; * = ZnTriPP-C₂-ZnTriPP 3; aliphatic hydrocarbon residues present)

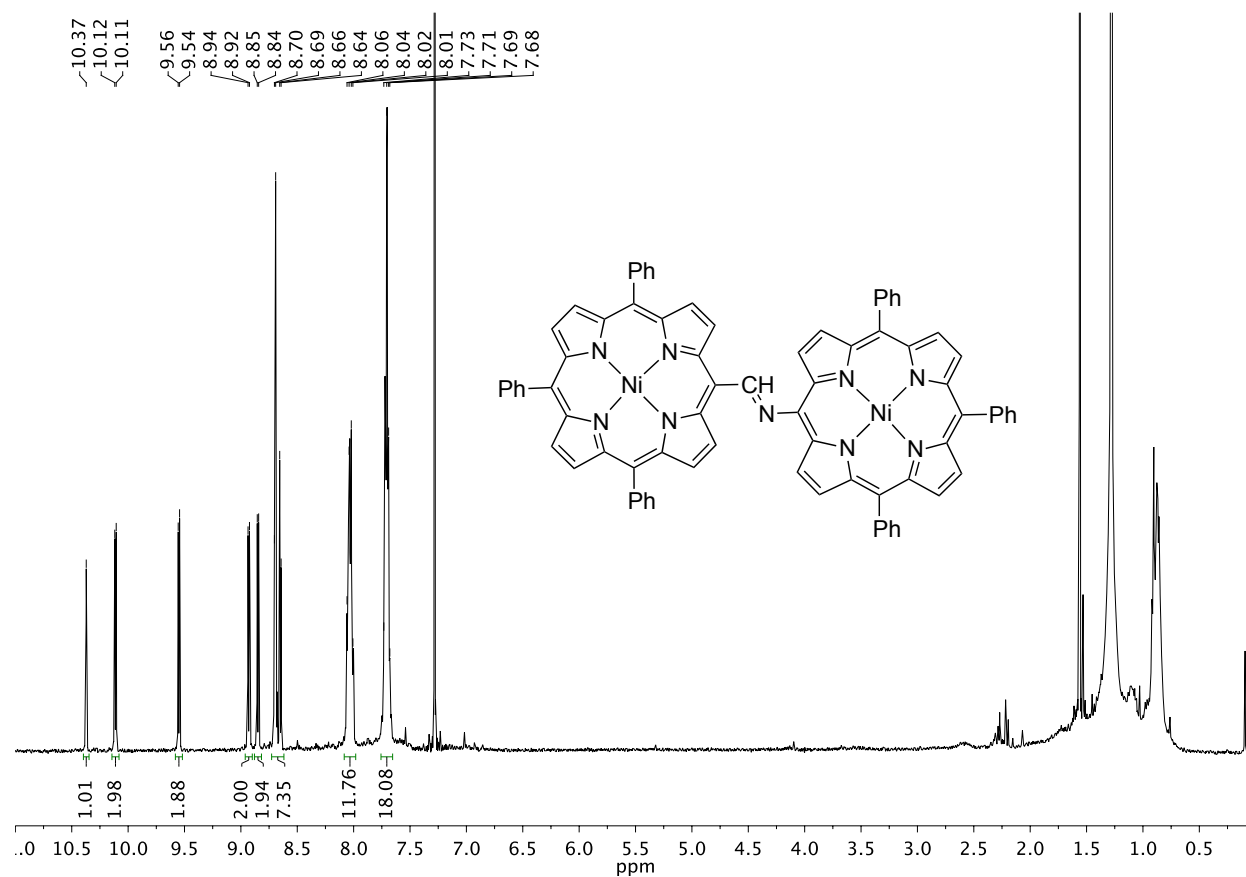


HRMS (MALDI)

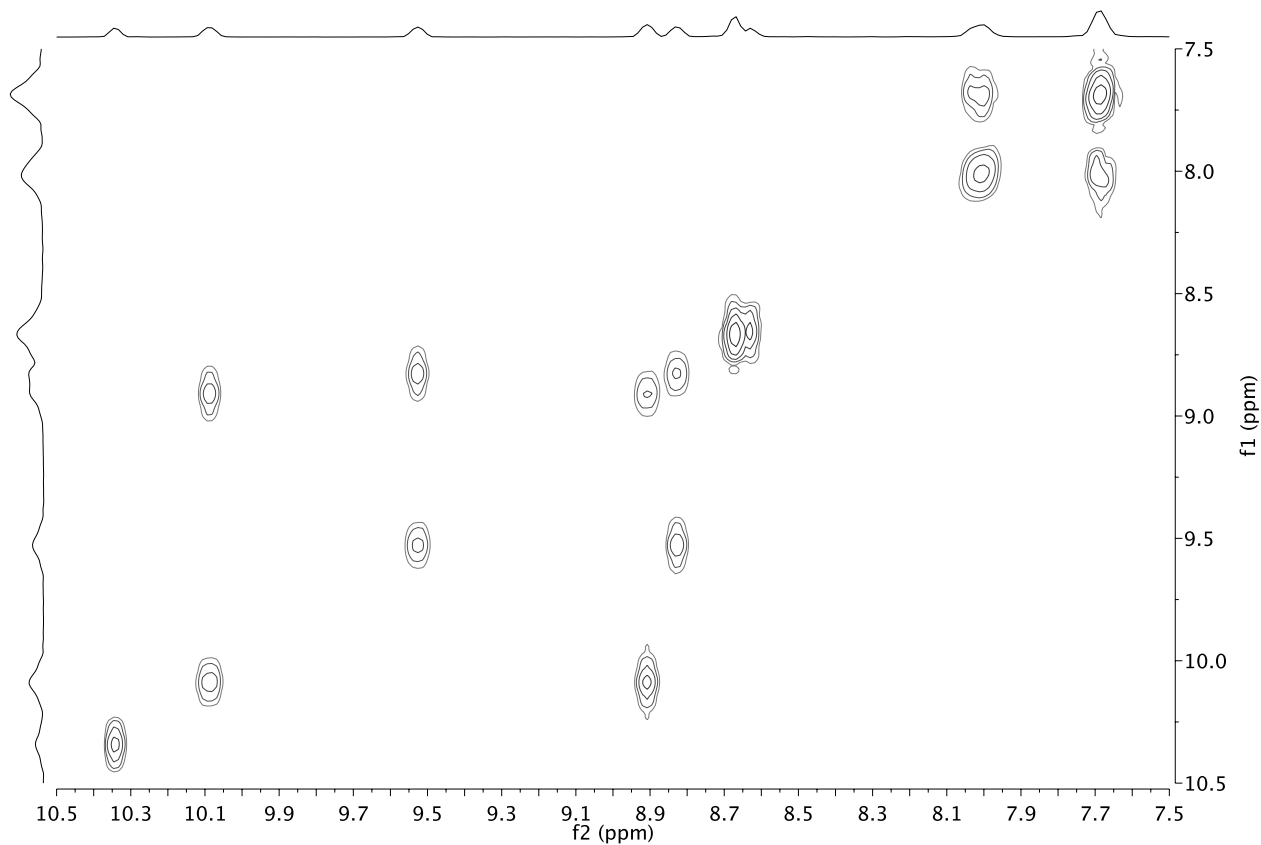
Applied Biosystems 4700 Proteomics Analyzer 66

4700 Reflector Spec #1 MC=>MC[BP = 1250.3, 656]

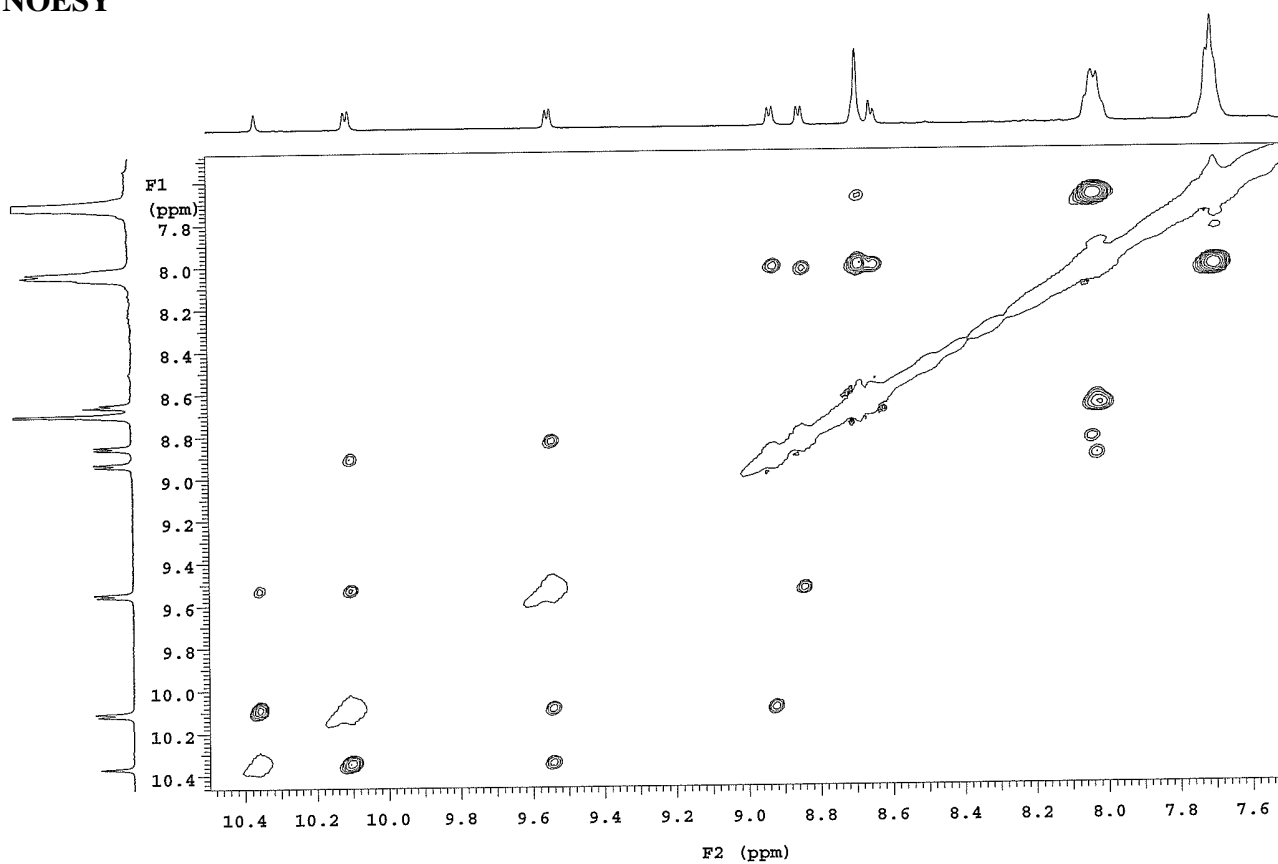


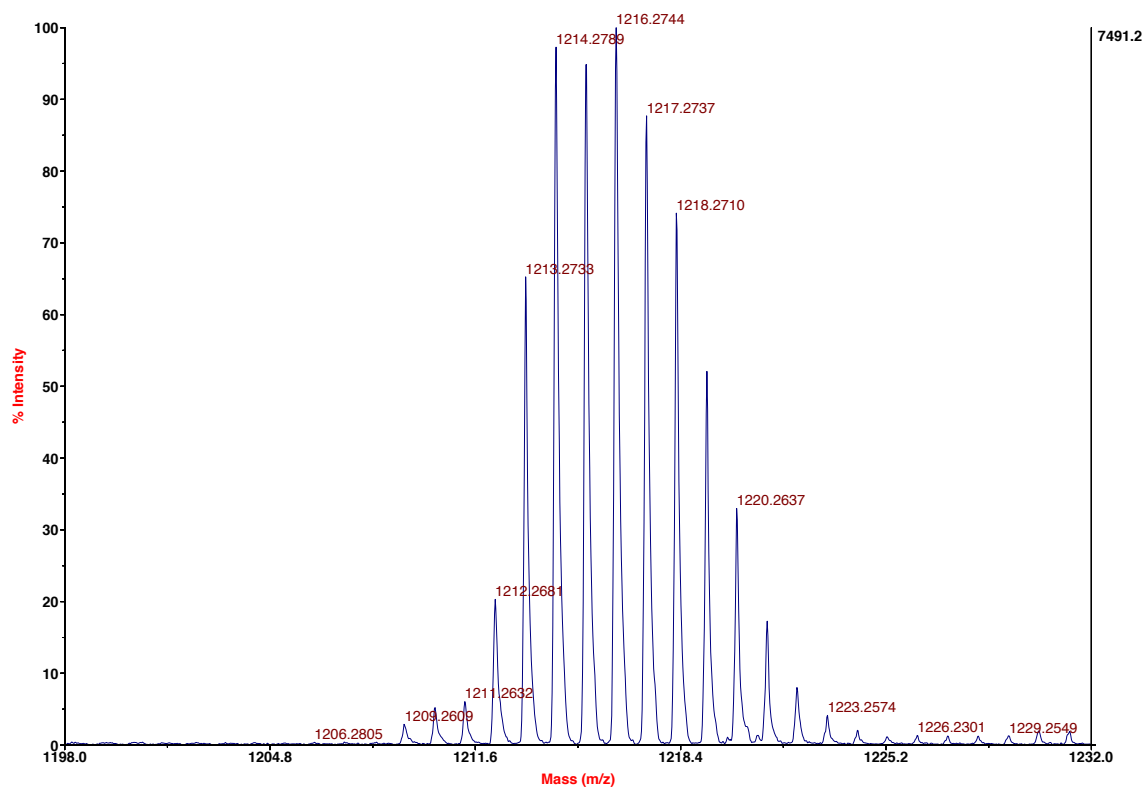
NiTriPP-CH=N-NiTriPP 29 (aliphatic hydrocarbon residues present)

COSY

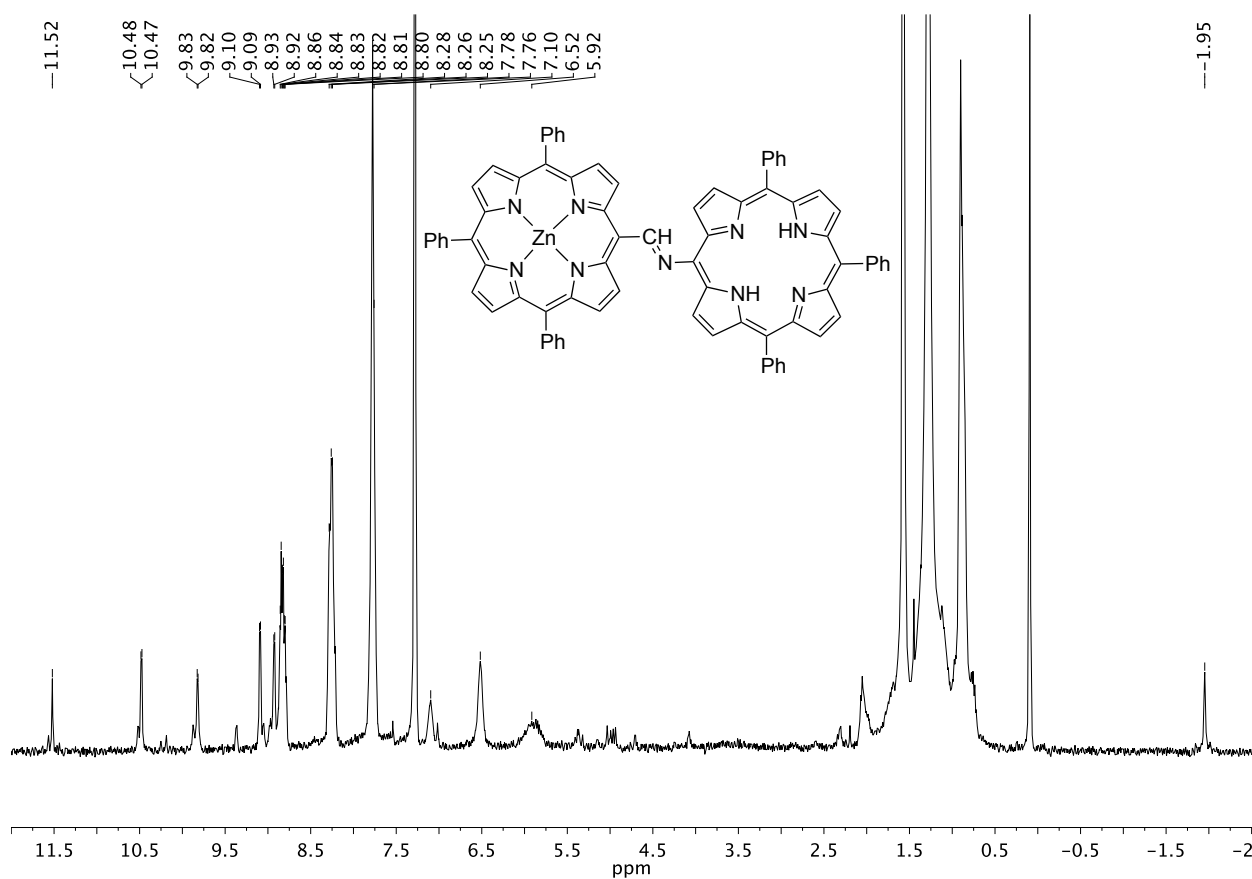


NOESY

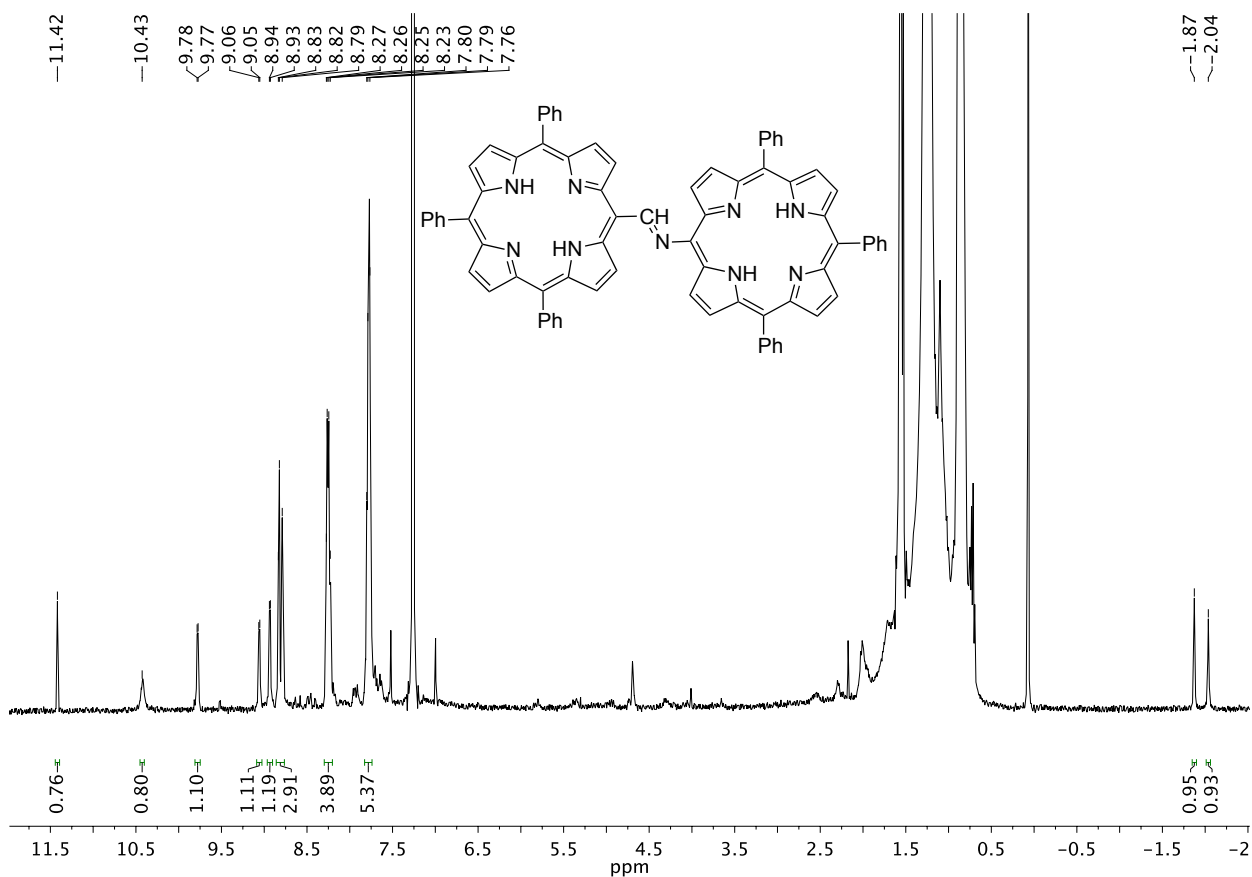


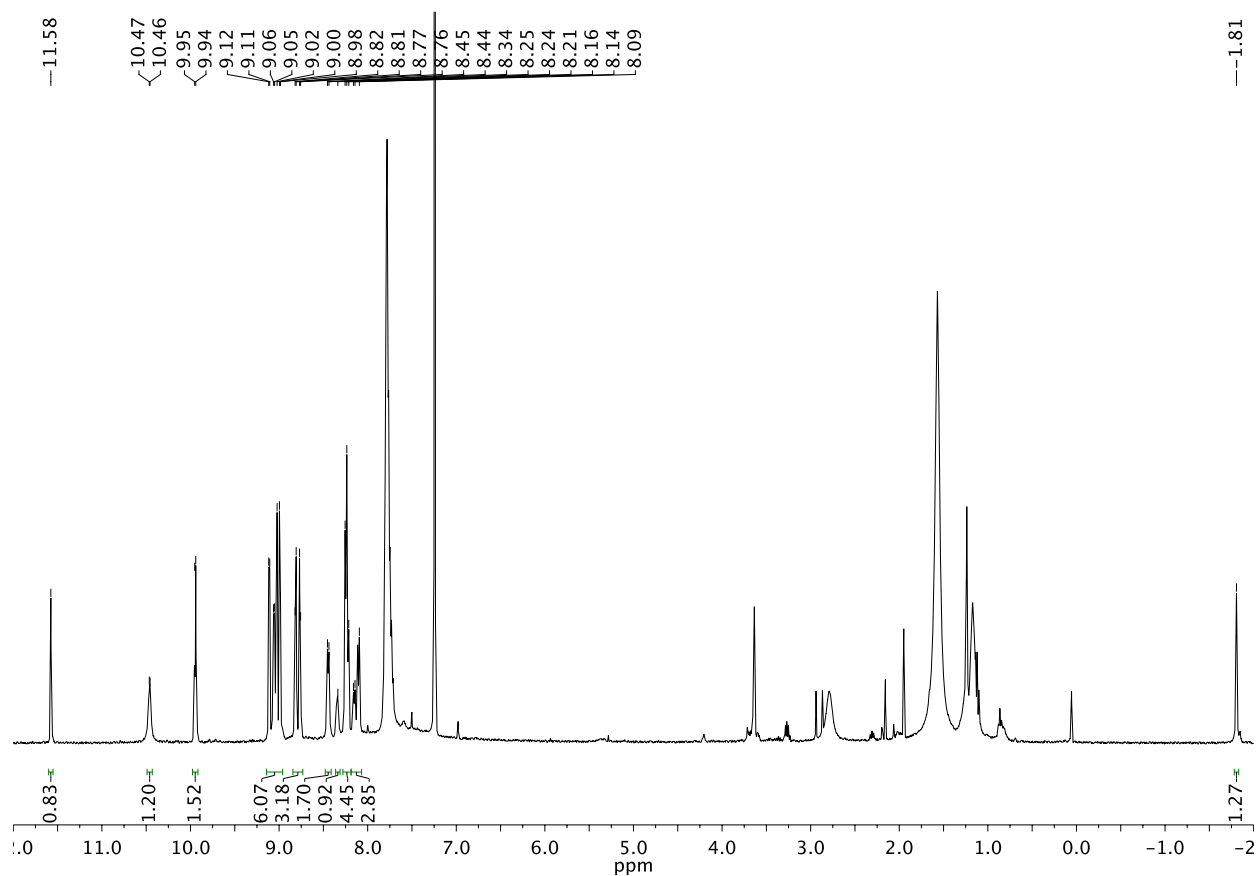
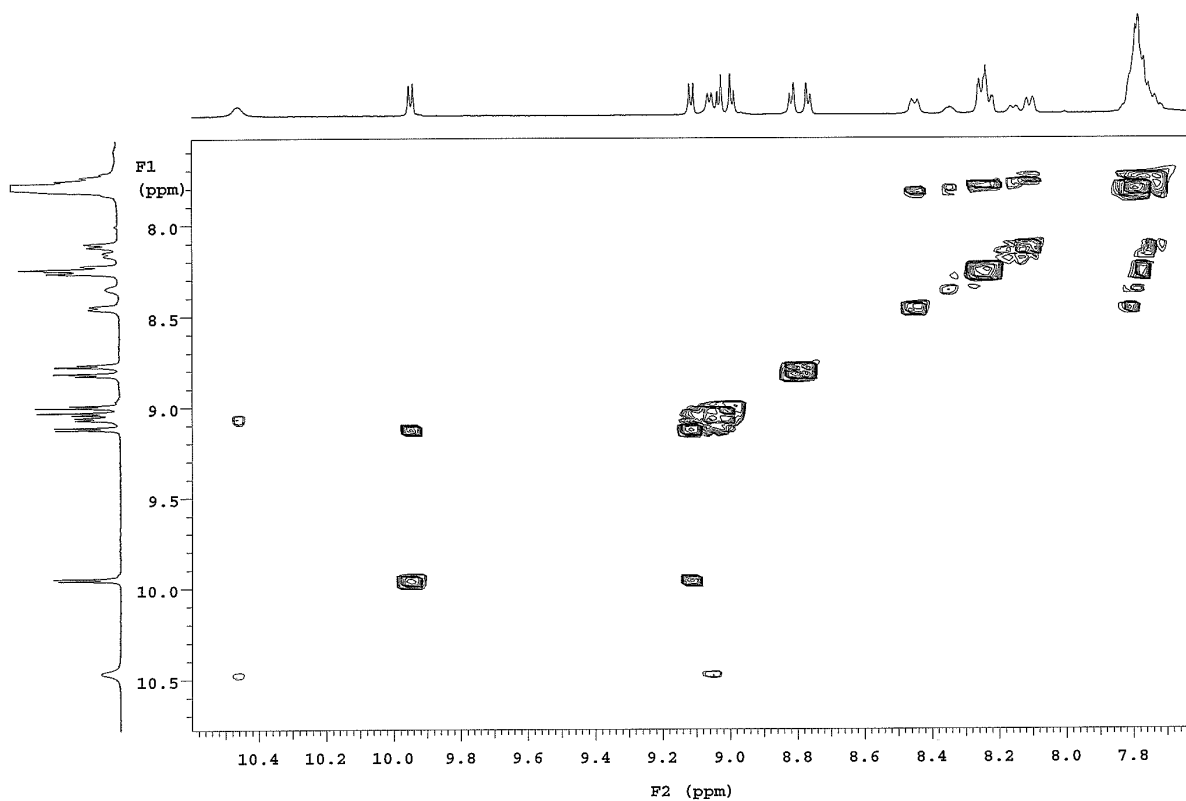
HRMS (MALDI)*Applied Biosystems 4700 Proteomics Analyzer 66***4700 Reflector Spec #1 MC=>MC[BP = 1216.3, 7491]**

ZnTriPP-CH=N-H₂TriPP 31 (minor porphyrin impurities, some coordinated pyridine and aliphatic hydrocarbon and other residues present)

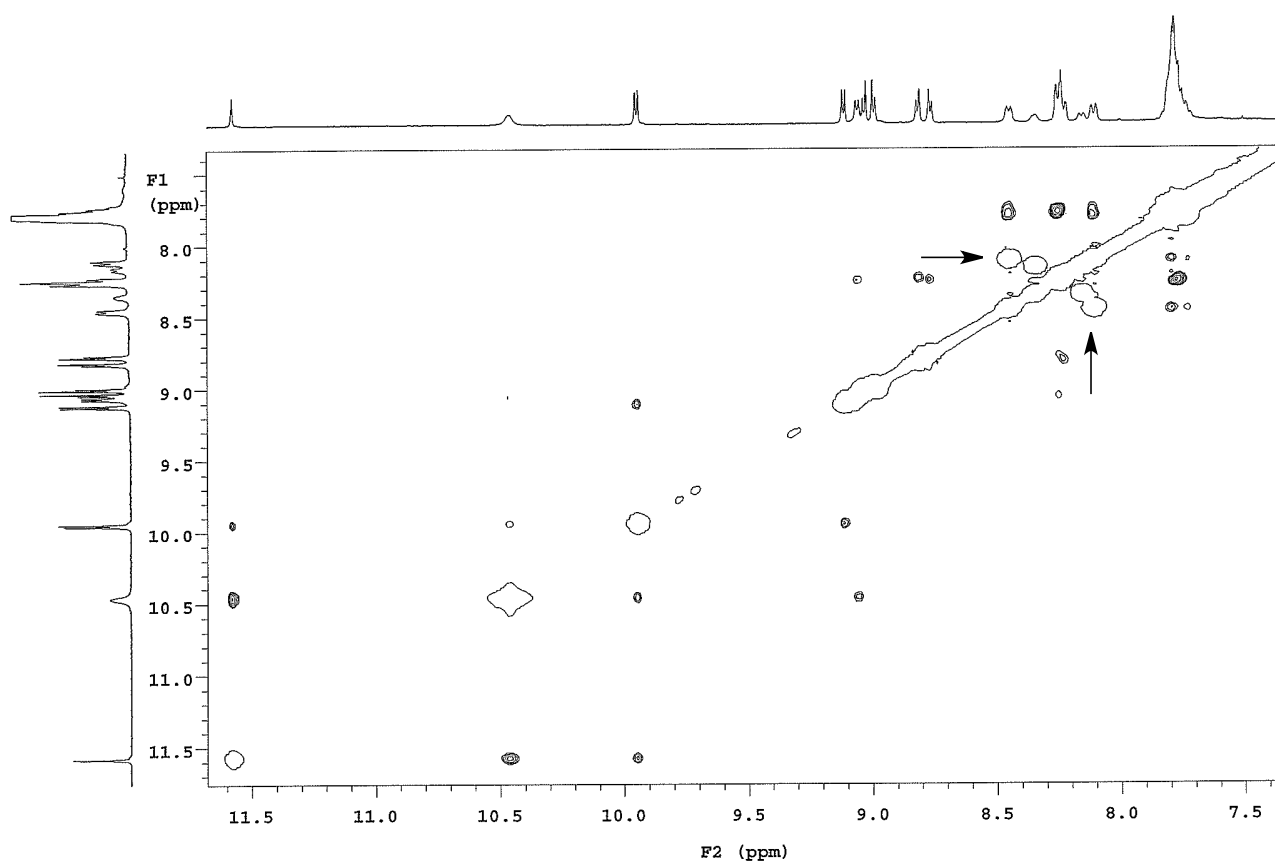


H₂TriPP-CH=N-H₂TriPP 32 (aliphatic hydrocarbon and other residues present)



InXTriPP-CH=N-H₂TriPP 35 (aliphatic hydrocarbon and other residues present)**COSY**

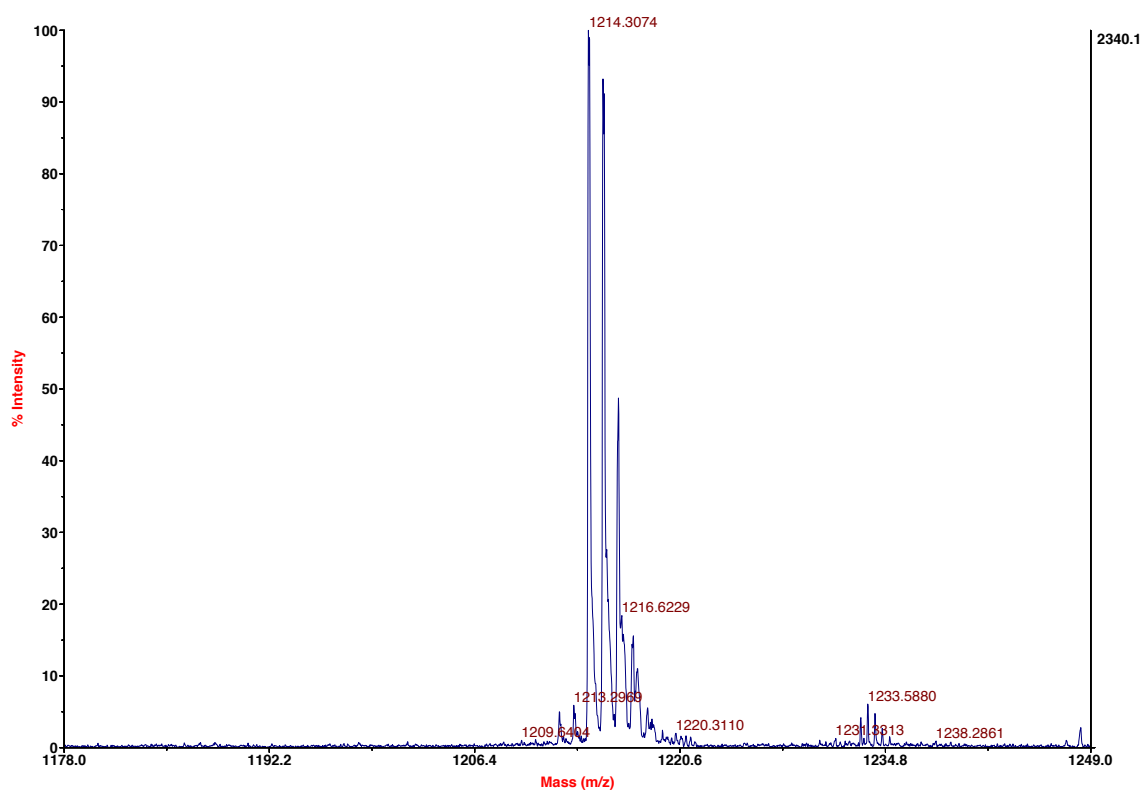
NOESY (the peaks marked with arrows are positive contours, i.e. EXSY cross-peaks showing exchange between *ortho*-phenyl protons on the lateral phenyl groups on the indium(III) ring)

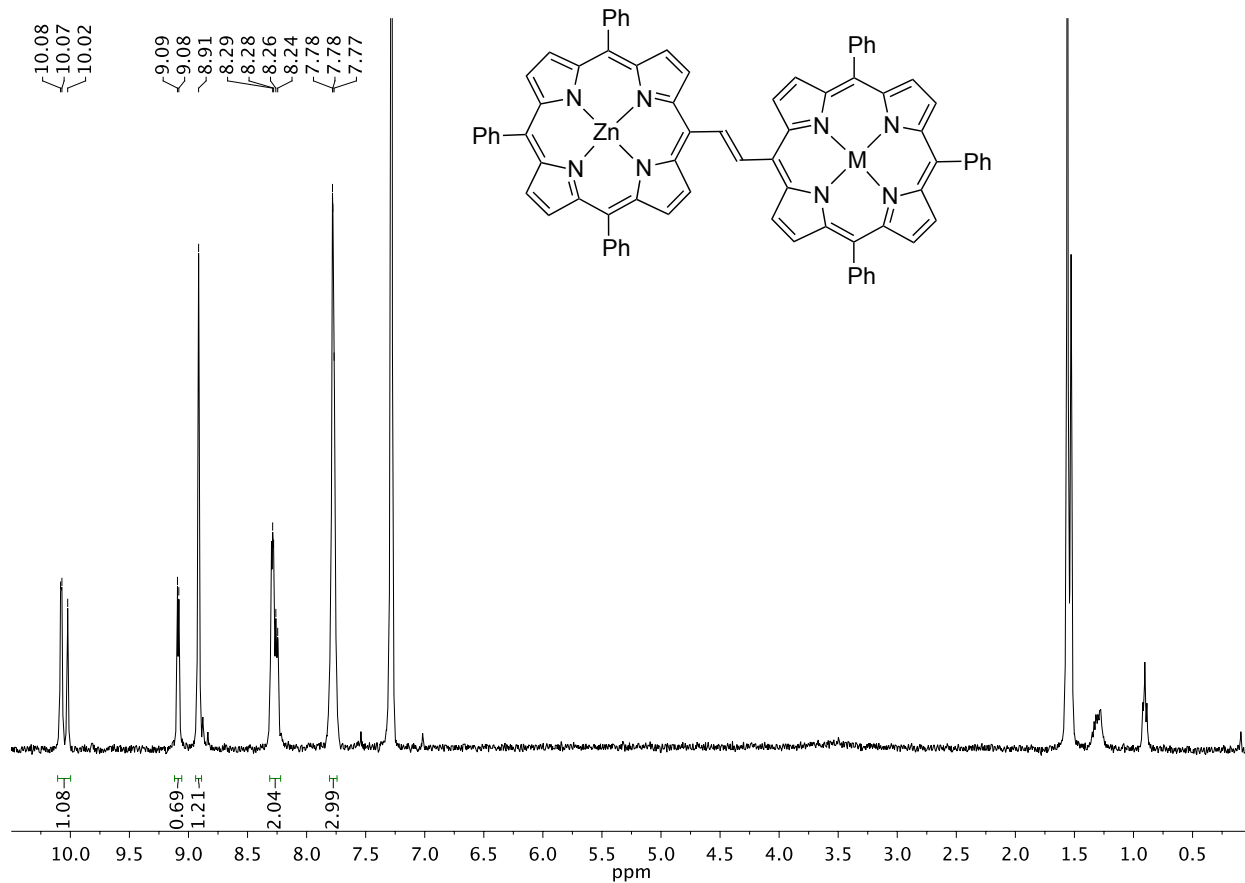


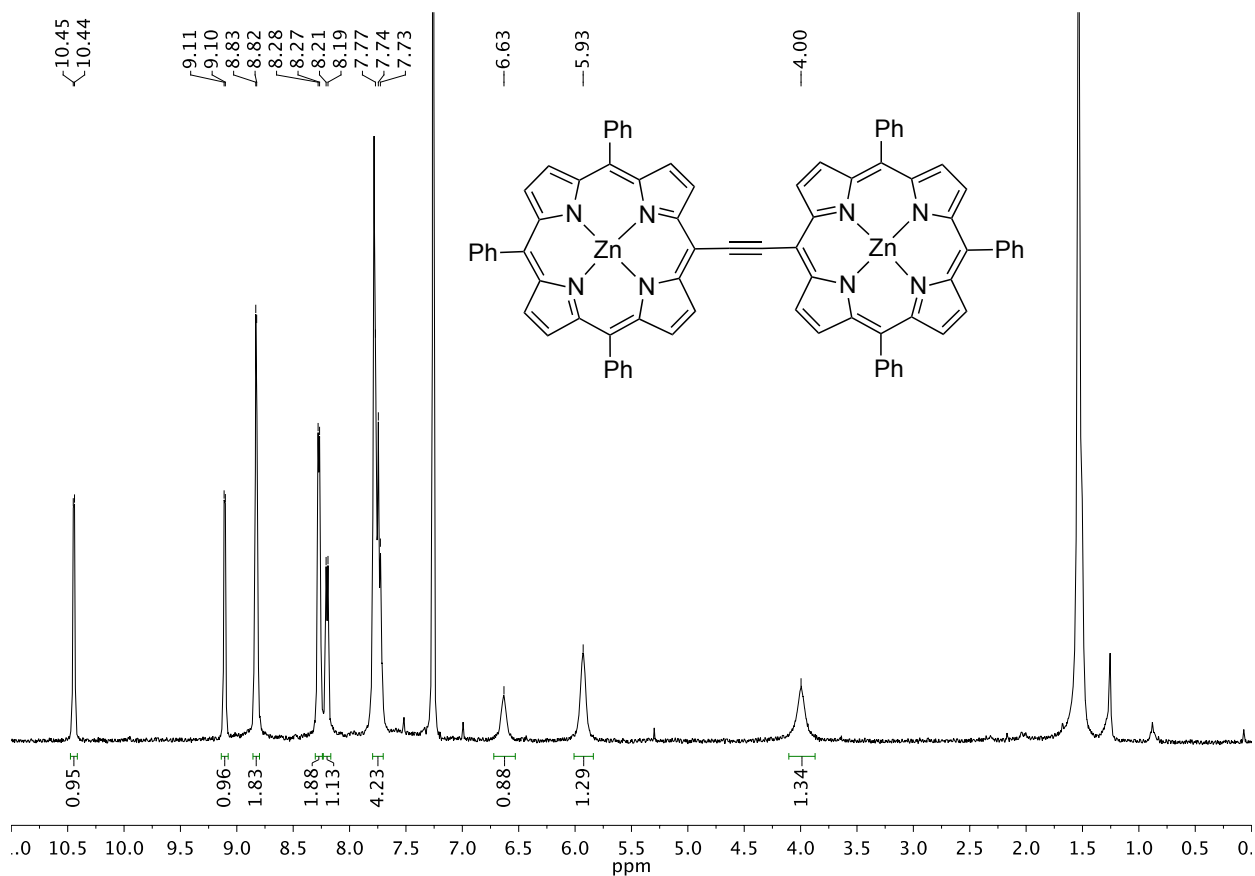
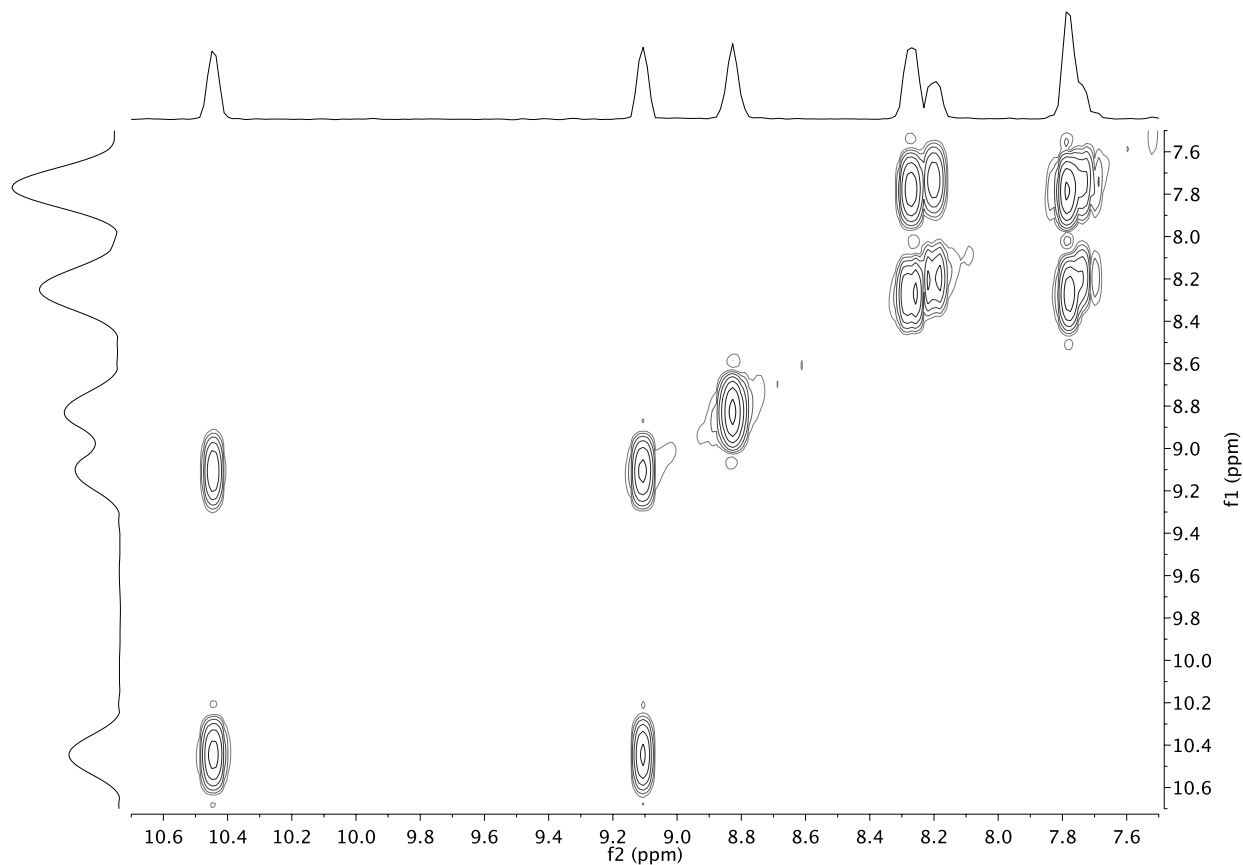
HRMS (MALDI)

Applied Biosystems 4700 Proteomics Analyzer 66

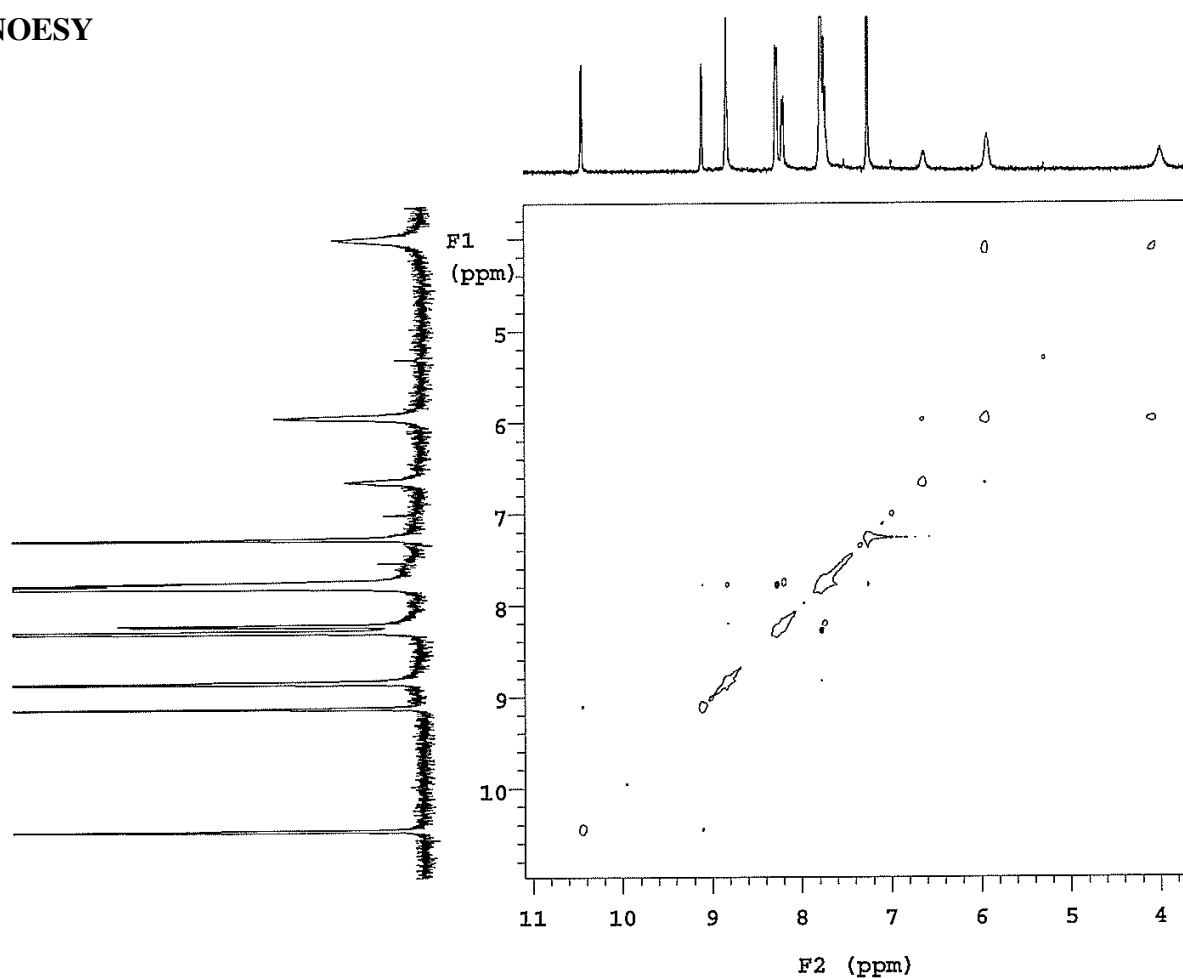
4700 Reflector Spec #1 MC=>MC=>MC[BP = 1214.3, 2340]



ZnTriPP-CH=CH-ZnTriPP 2

ZnTriPP-C₂-ZnTriPP 3 (pyridine coordinated)**COSY**

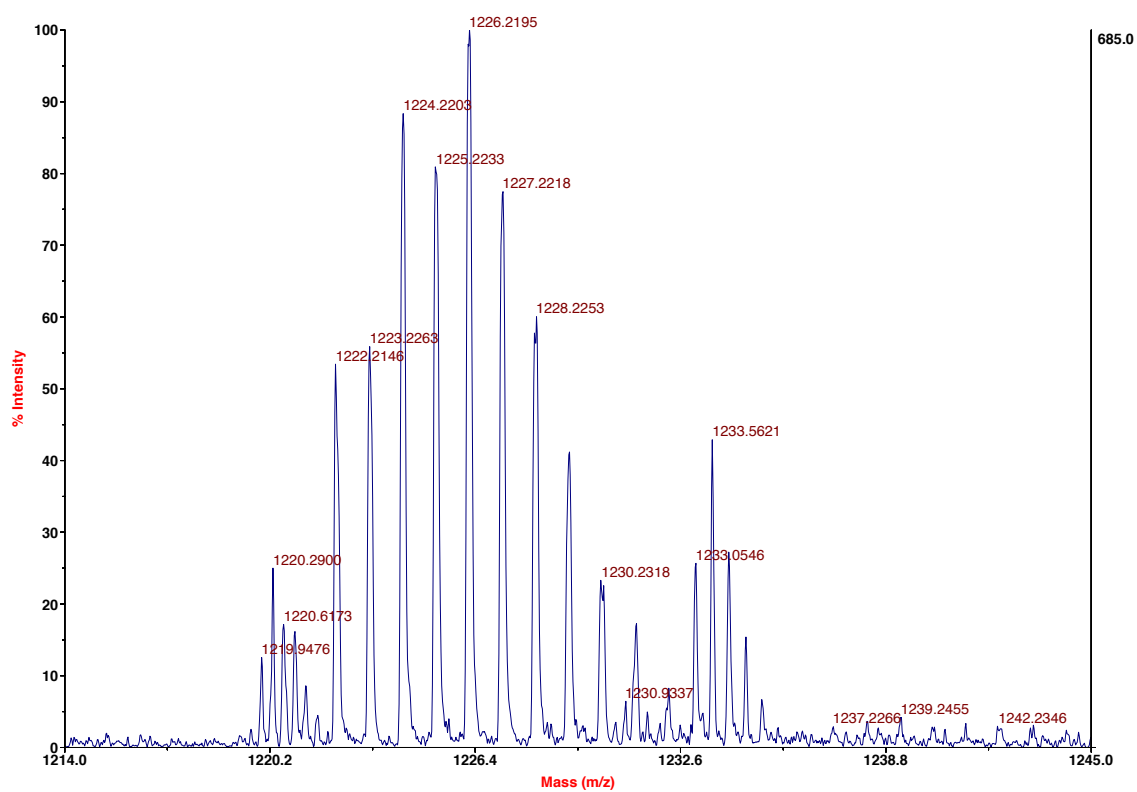
NOESY

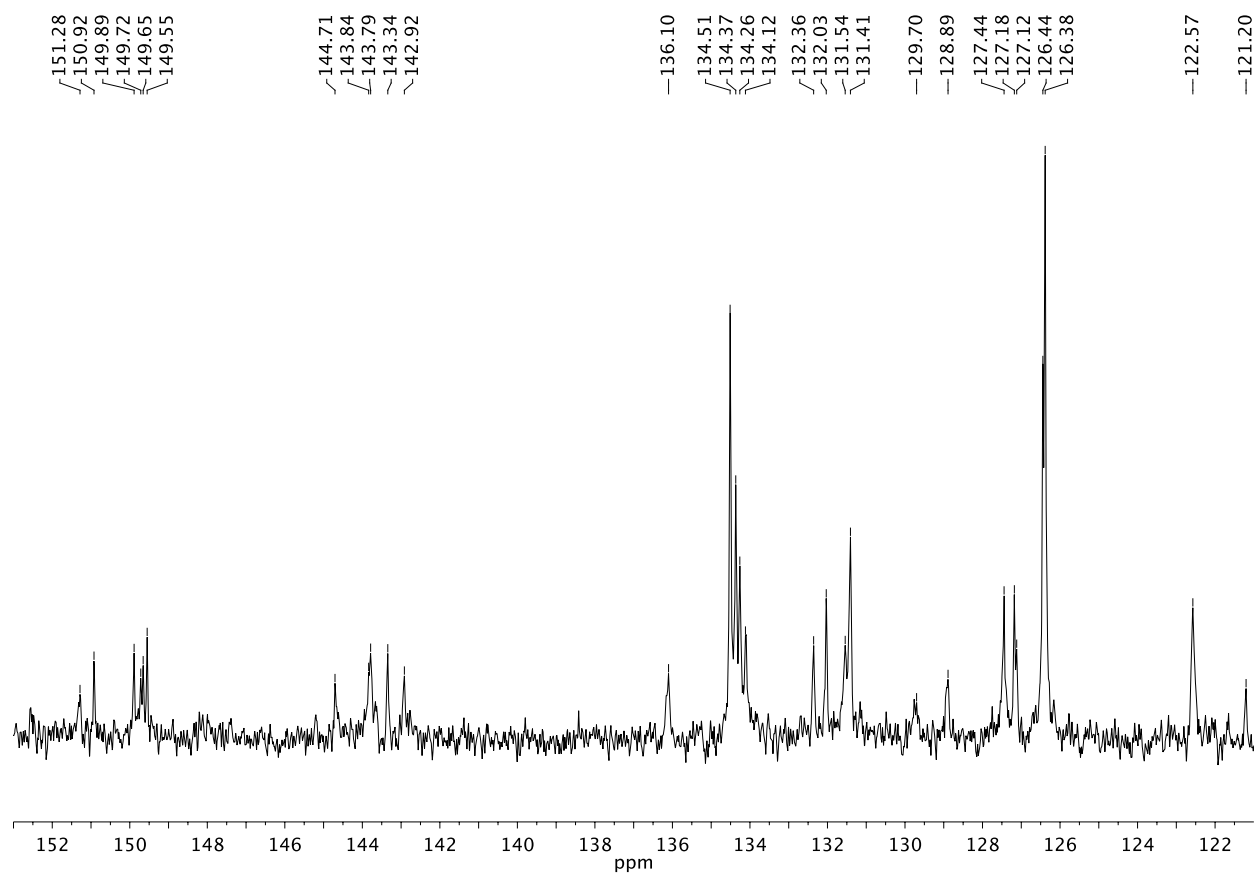
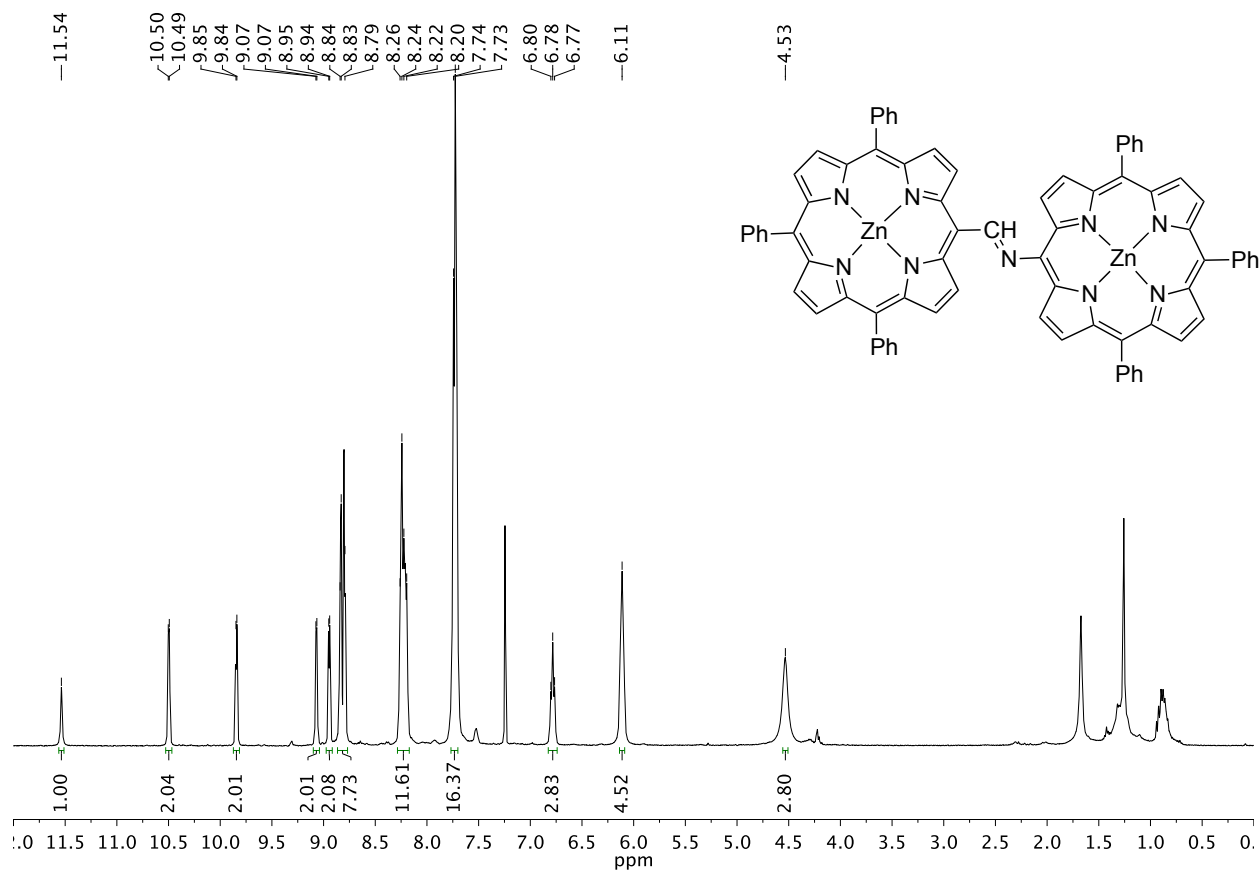


HRMS (MALDI)

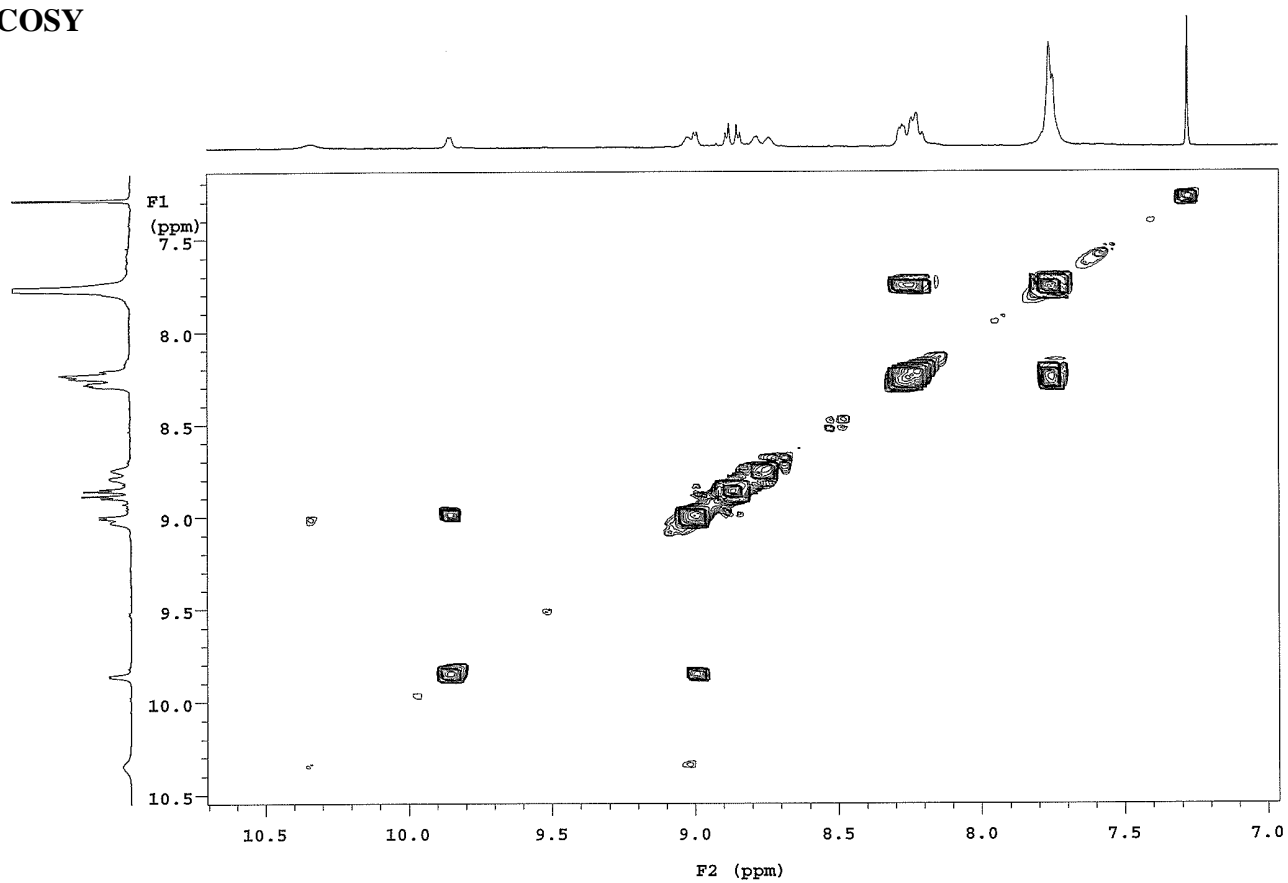
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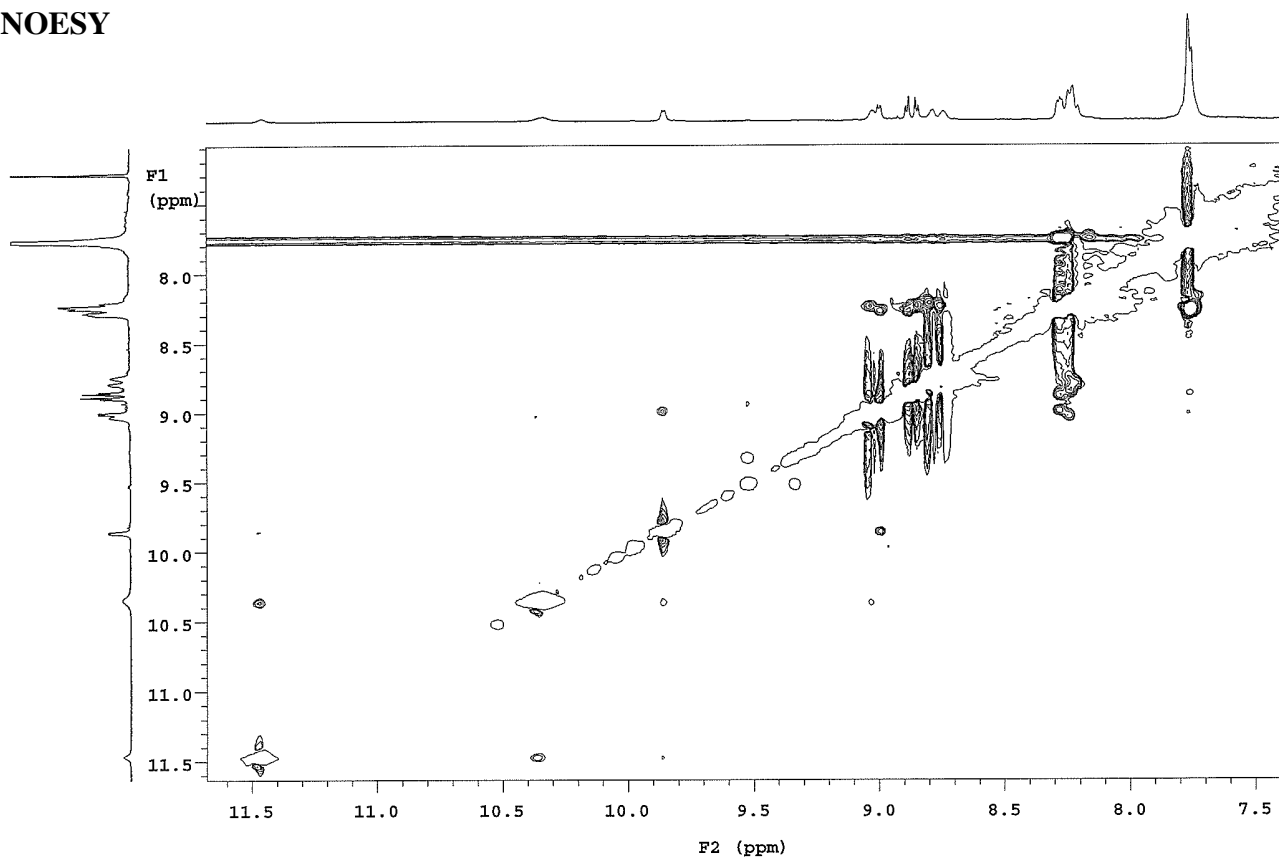


ZnTriPP-CH=N-ZnTriPP 4 (pyridine coordinated)

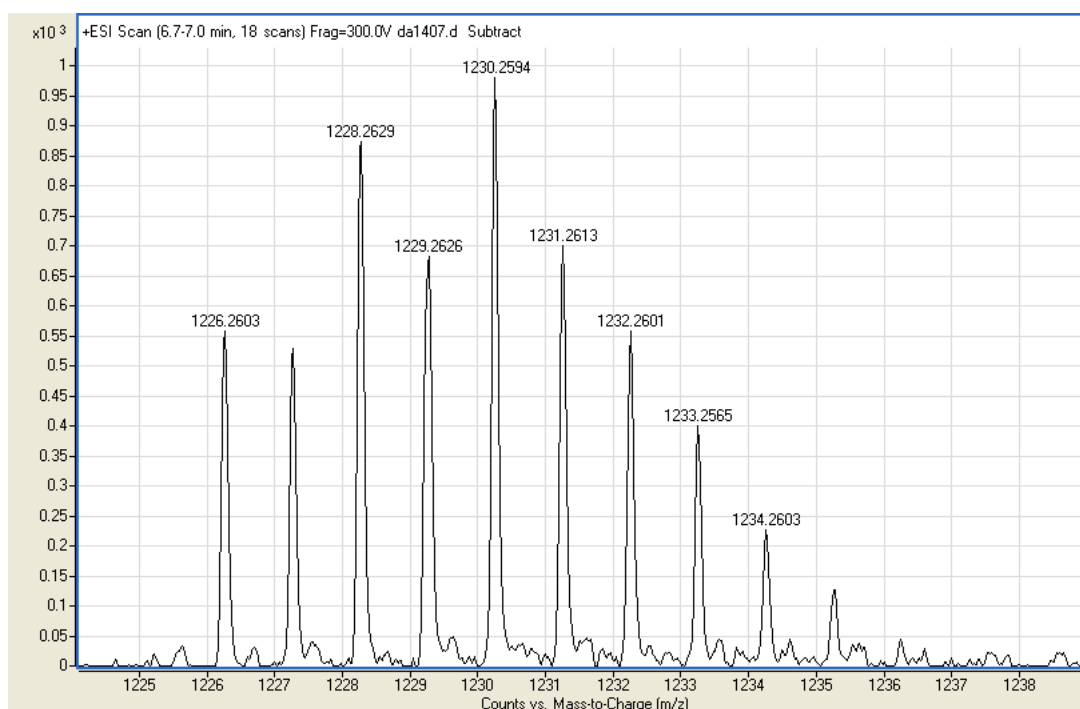
COSY



NOESY



HRMS (ESI)



ZnTriPP-N=N-ZnTriPP 5 (in CDCl₃ + C₅D₅N; * = water in exchange with pyridine)

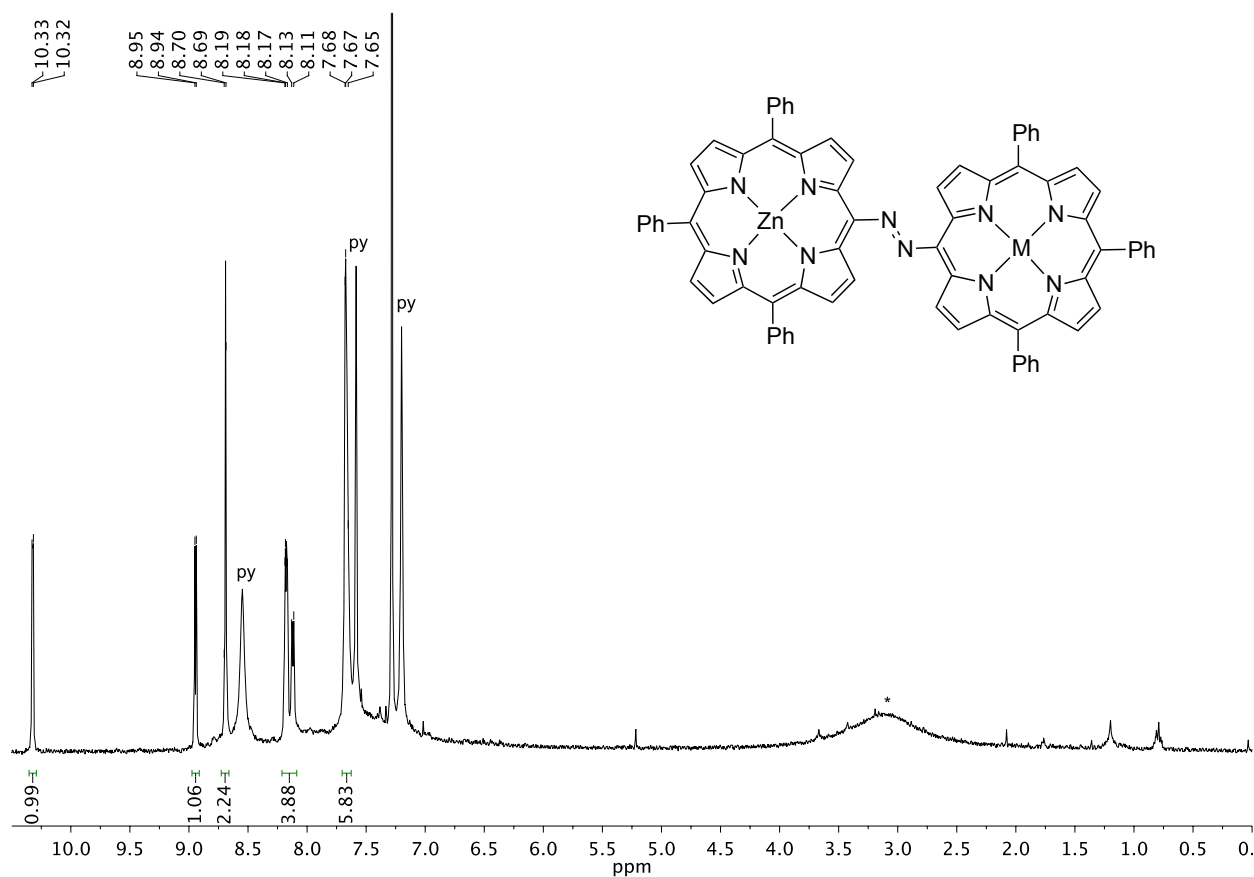


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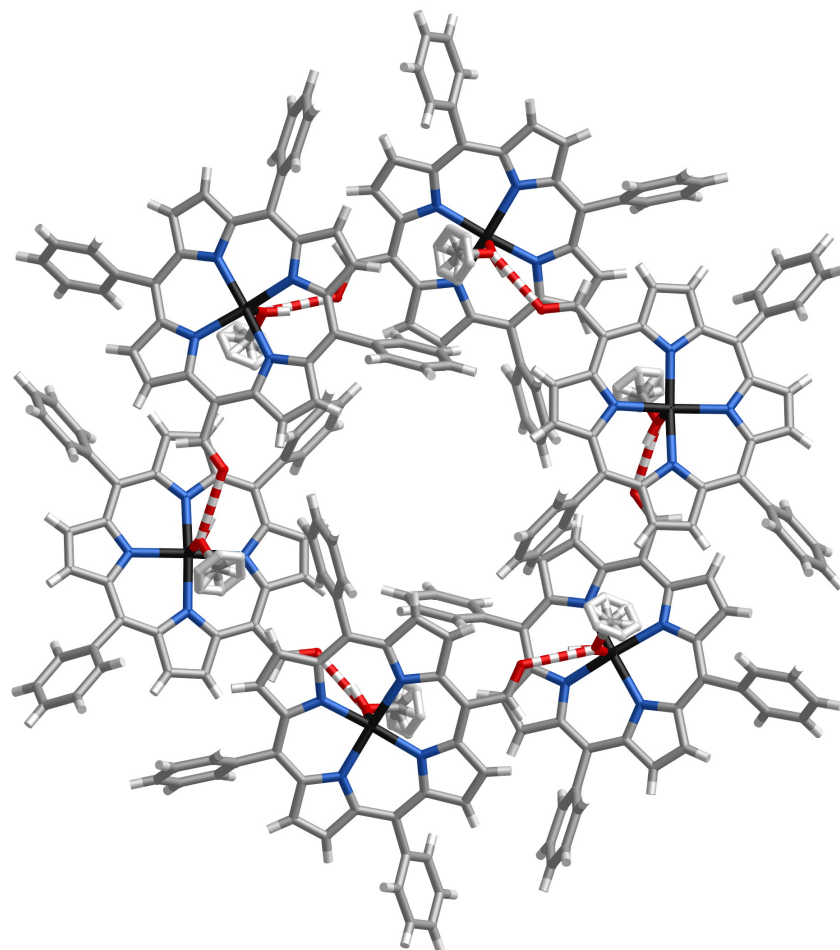


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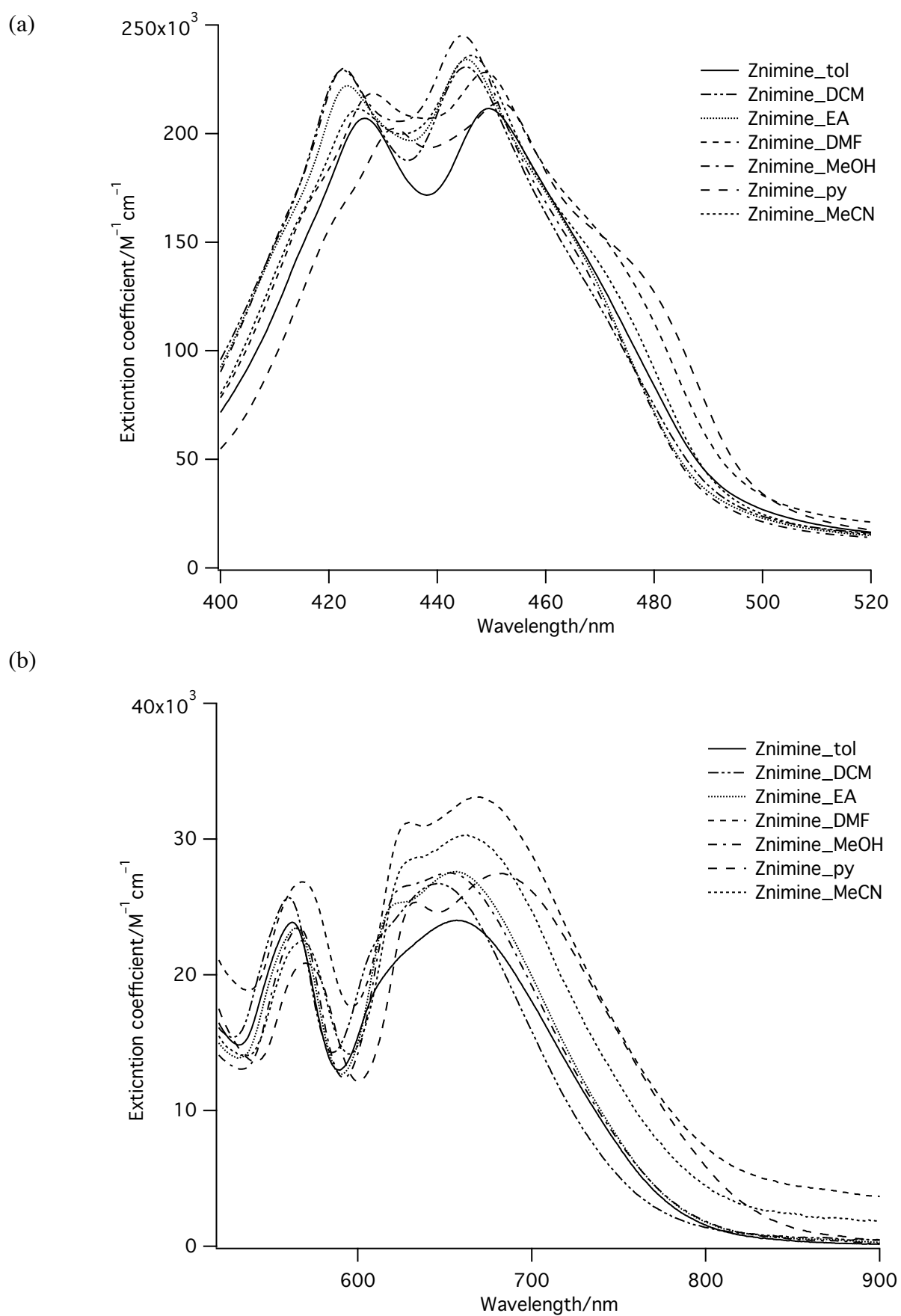


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