

Supporting Information for

Positional isomerism of unsymmetrical semirigid ligands toward the construction of discrete and infinite coordination architectures of zinc(II) and cadmium(II) complexes

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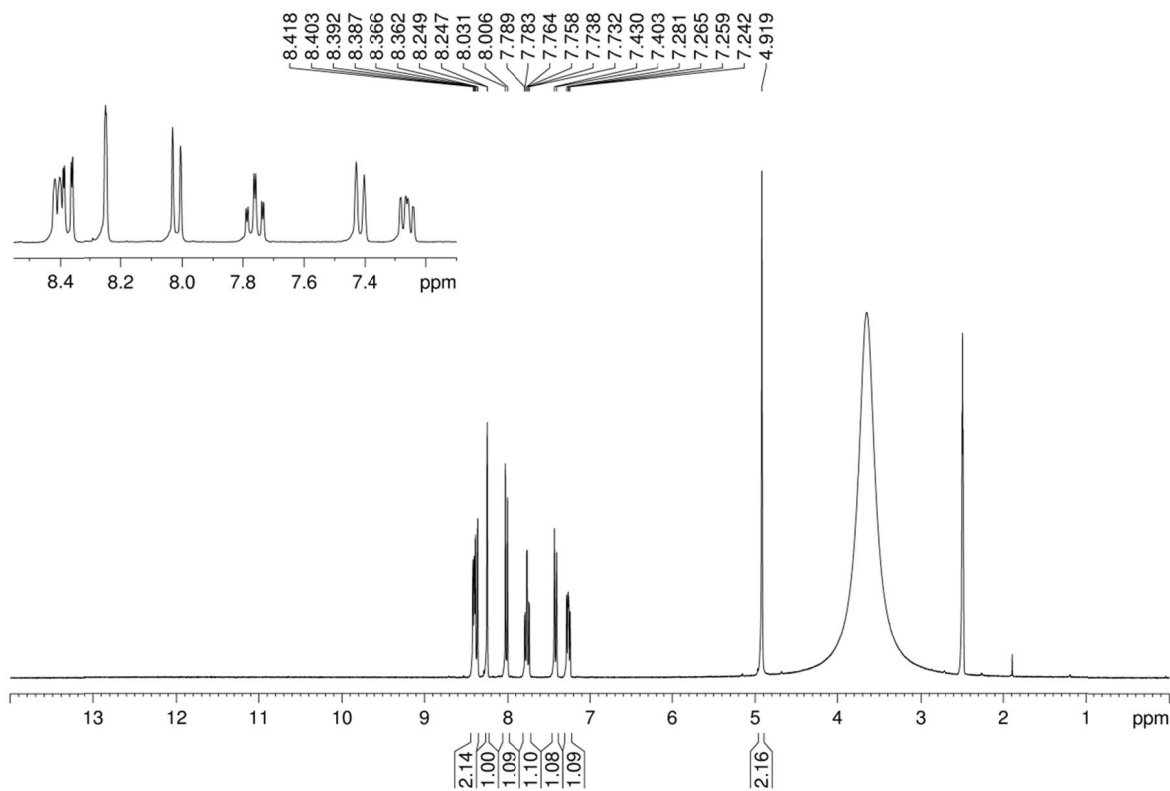


Fig. S1 ^1H NMR spectrum of 2-(2-pyridylmethyl)-1,3-dioxisoindoline-5-carboxylic acid (HInMe-2-py) in $\text{DMSO-}d_6$ at room temperature.

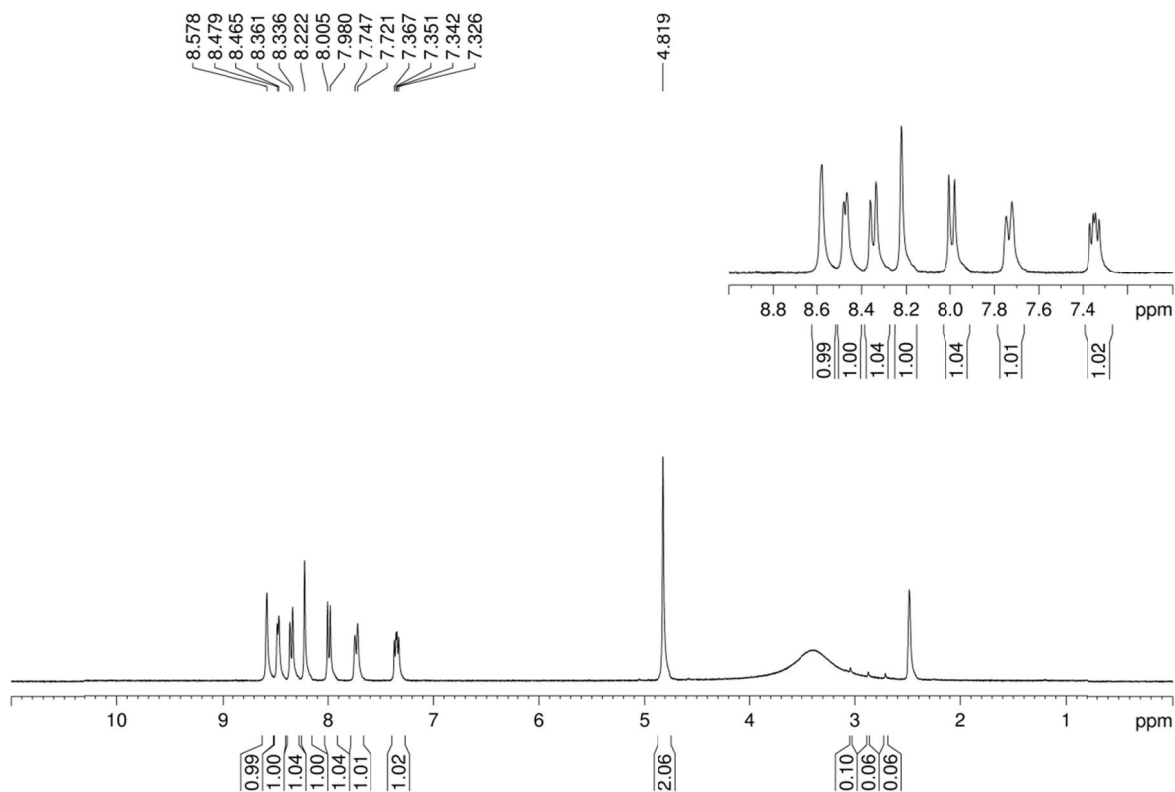


Fig. S2 ^1H NMR spectrum of 2-(3-pyridylmethyl)-1,3-dioxisoindoline-5-carboxylic acid (HInMe-3-py) in $\text{DMSO-}d_6$ at room temperature.

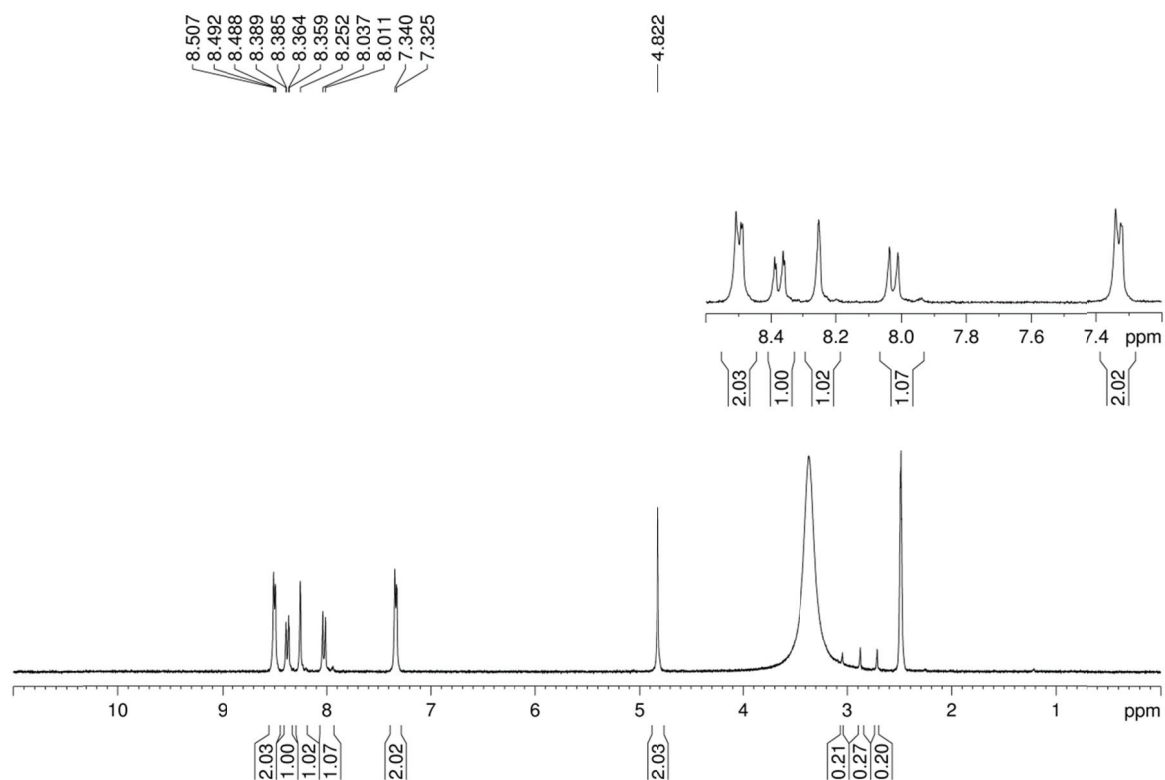


Fig. S3 ^1H NMR spectrum of 2-(4-pyridylmethyl)-1,3-dioxisoindoline-5-carboxylic acid (HInMe-4-py) in $\text{DMSO-}d_6$ at room temperature.

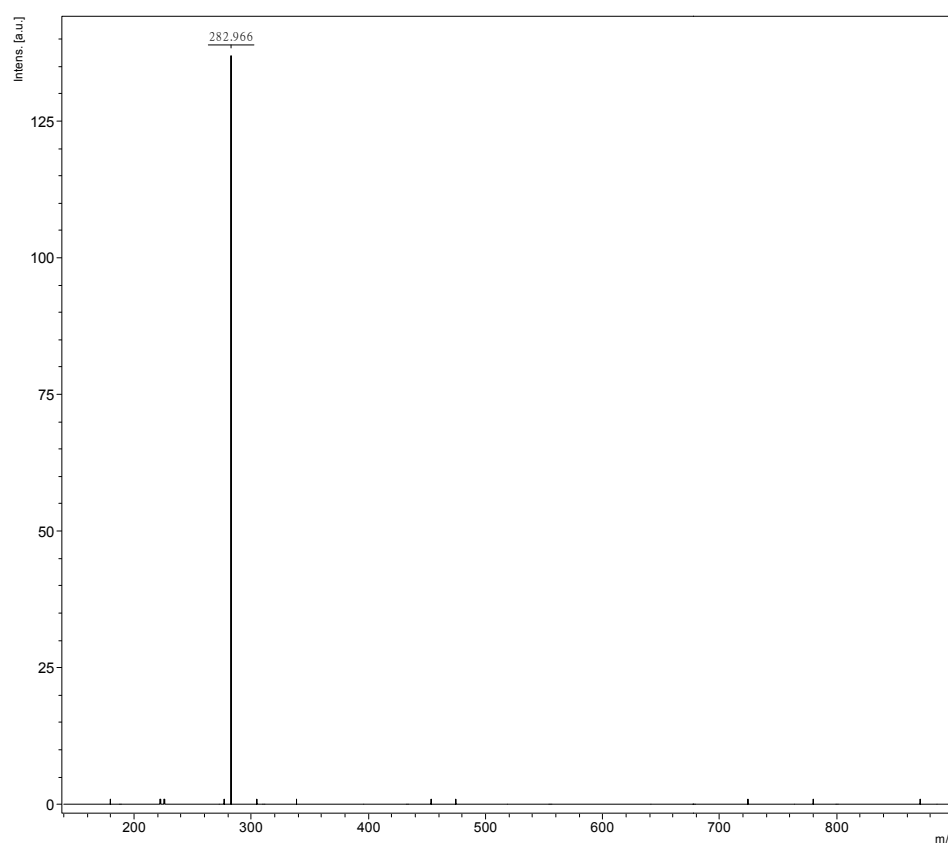


Fig. S4 Matrix-assisted laser desorption/ionization time of flight (MALDI-TOF) mass spectrum of 2-(2-pyridylmethyl)-1,3-dioxisoindoline-5-carboxylic acid (HInMe-2-py).

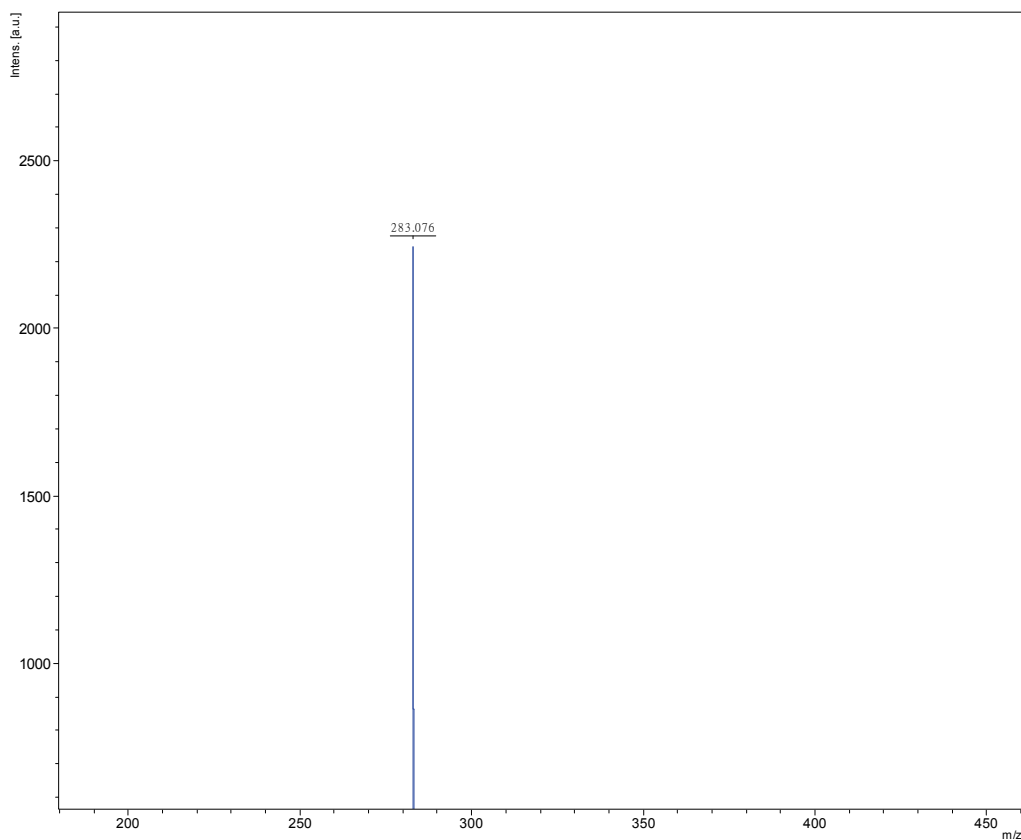


Fig. S5 Matrix-assisted laser desorption/ionization time of flight (MALDI-TOF) mass spectrum of 2-(3-pyridylmethyl)-1,3-dioxisoindoline-5-carboxylic acid (HInMe-3-py).

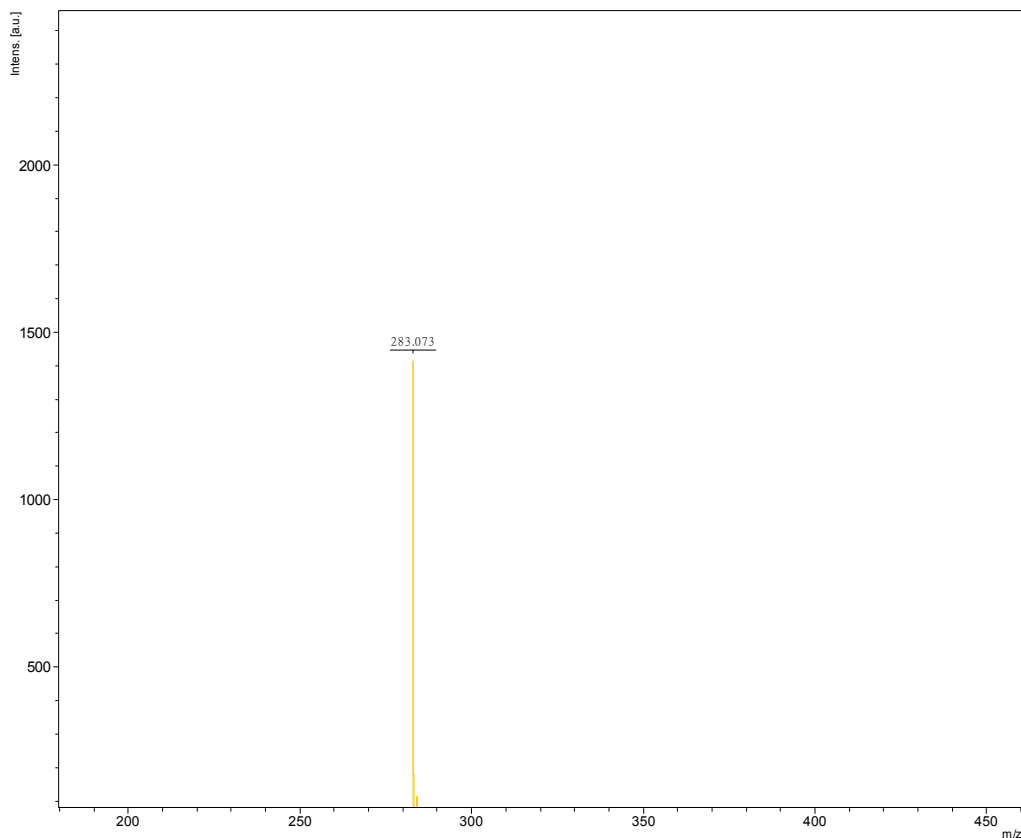


Fig. S6 Matrix-assisted laser desorption/ionization time of flight (MALDI-TOF) mass spectrum of 2-(4-pyridylmethyl)-1,3-dioxisoindoline-5-carboxylic acid (HInMe-4-py).

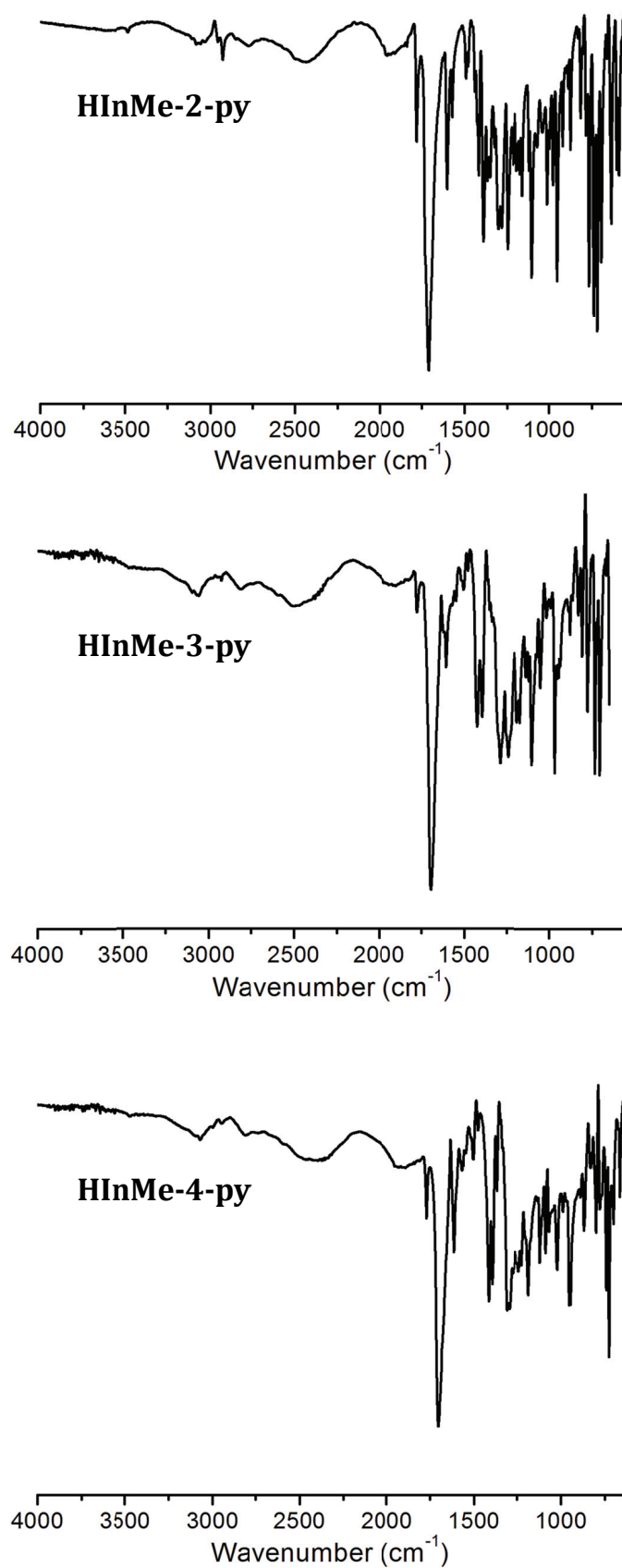


Fig. S7 Infrared (IR) spectra of HInMe-*n*-py (*n* = 2, 3, 4) ligands.

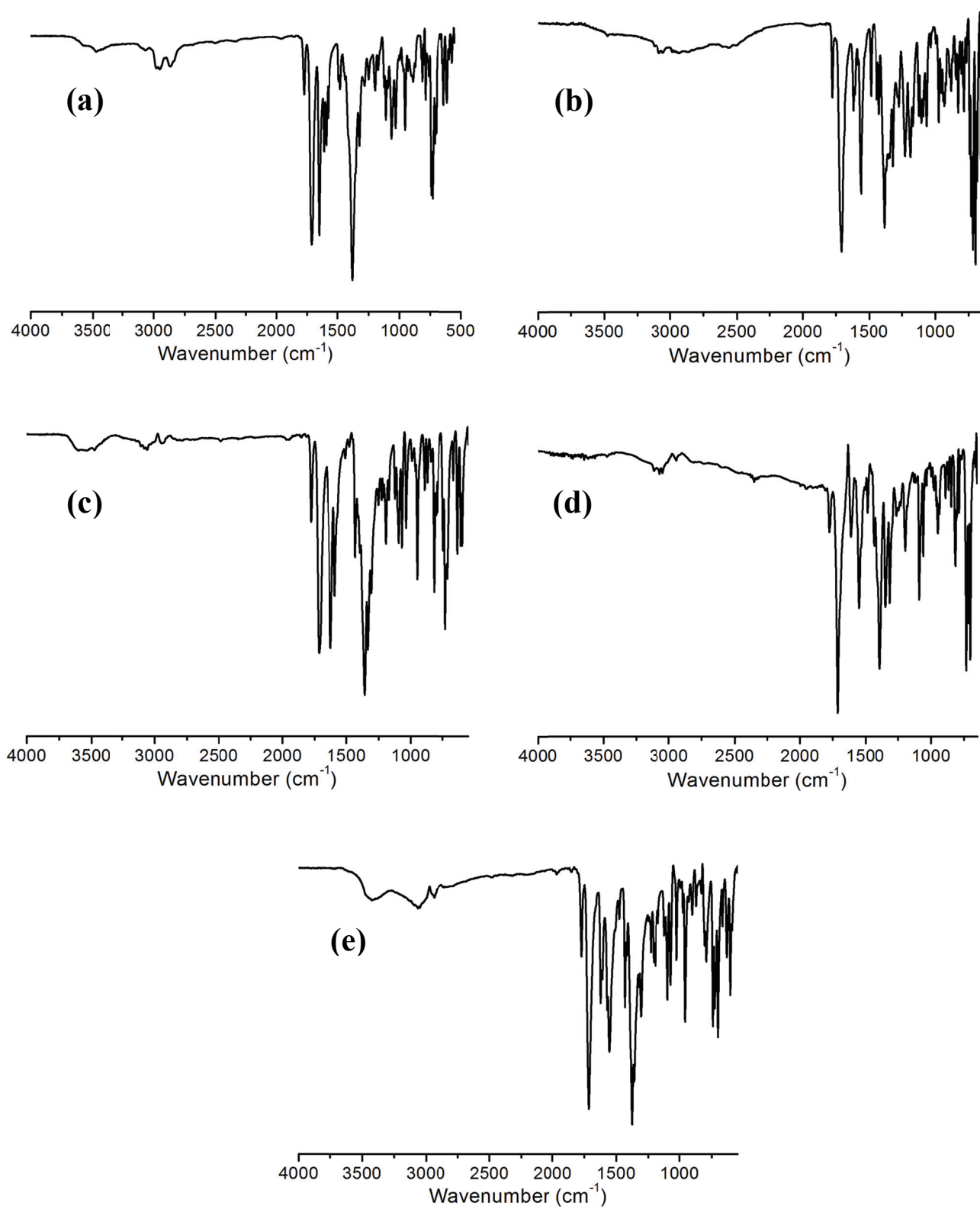


Fig. S8 Infrared (IR) spectra of (a-e) 1-5.

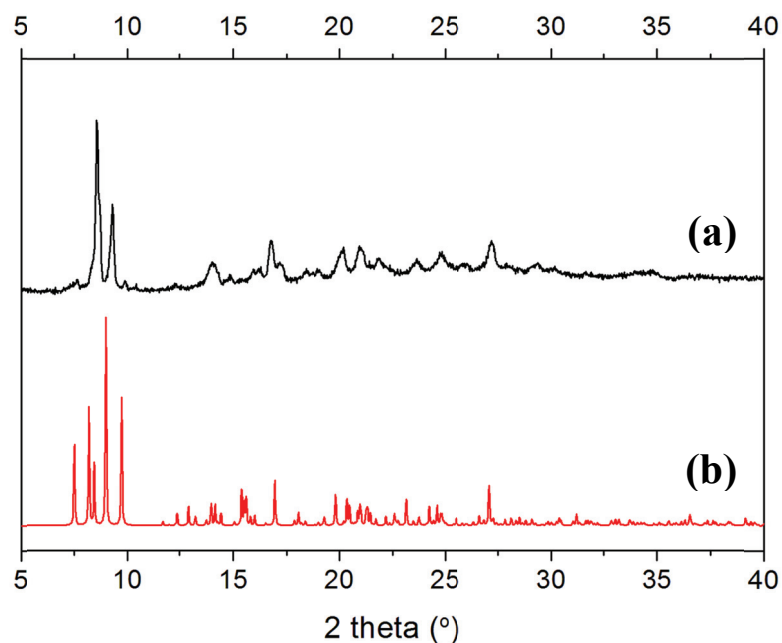


Fig. S9 X-Ray powder diffraction (XRPD) patterns of **1**: (a) a freshly grounded sample at room temperature; (b) simulated from the single-crystal data. For **1**, it is difficult to obtain a well-resolved XRPD pattern that is well-matched with the simulated XRPD patterns since lattice solvent molecules are gradually released during the XRPD measurement.

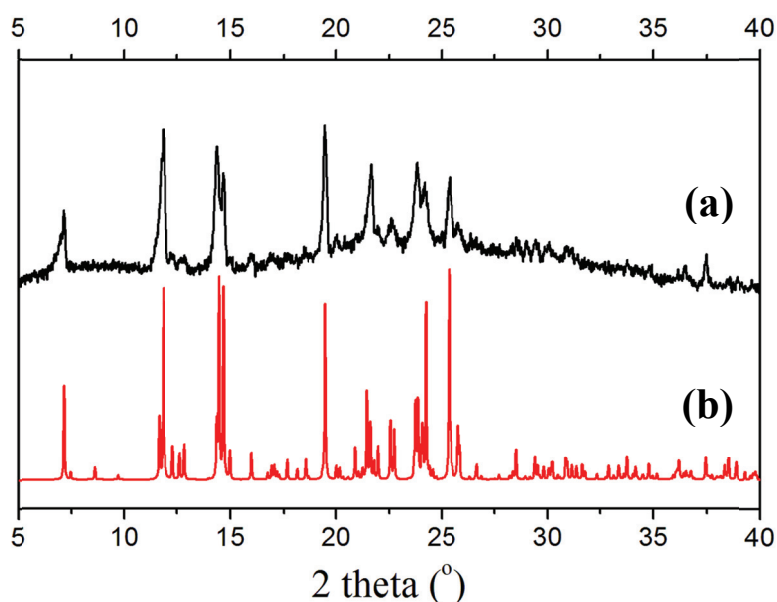


Fig. S10 X-ray powder diffraction (XRPD) patterns of **2**: (a) a freshly grounded sample at room temperature; (b) simulated from the single-crystal data.

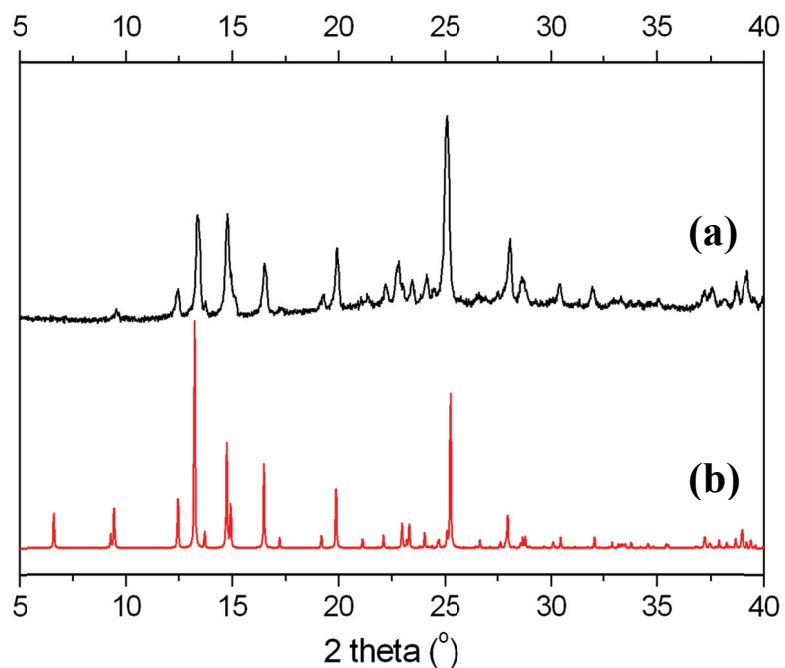


Fig. S11 X-Ray powder diffraction (XRPD) patterns of **3**: (a) a freshly grounded sample at room temperature; (b) simulated from the single-crystal data.

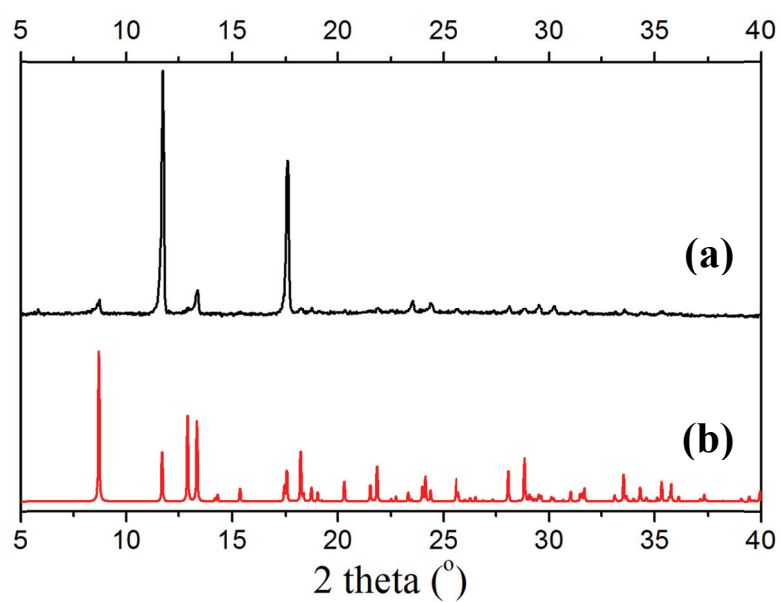


Fig. S12 X-Ray powder diffraction (XRPD) patterns of **4**: (a) a freshly grounded sample at room temperature; (b) simulated from the single-crystal data.

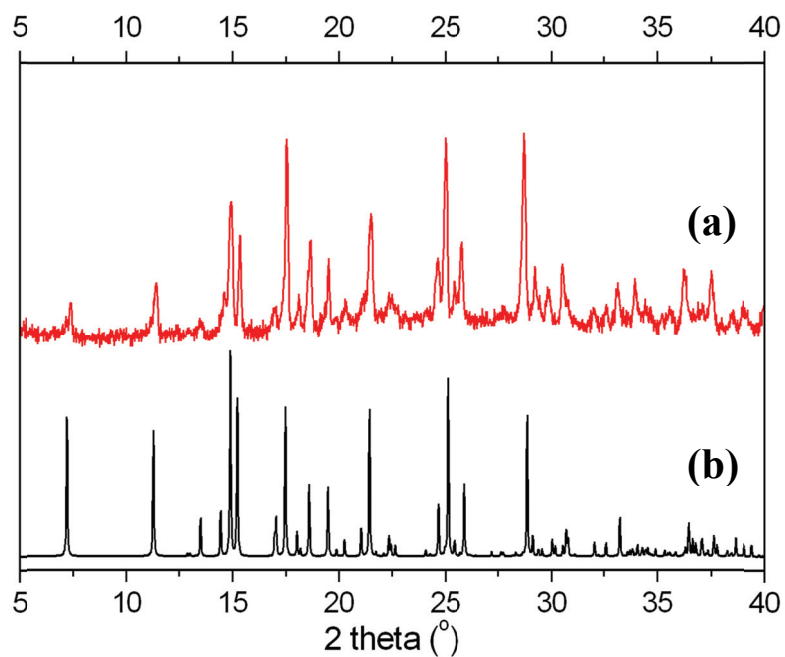


Fig. S13 X-Ray powder diffraction (XRPD) patterns of **5**: (a) a freshly grounded sample at room temperature; (b) simulated from the single-crystal data.

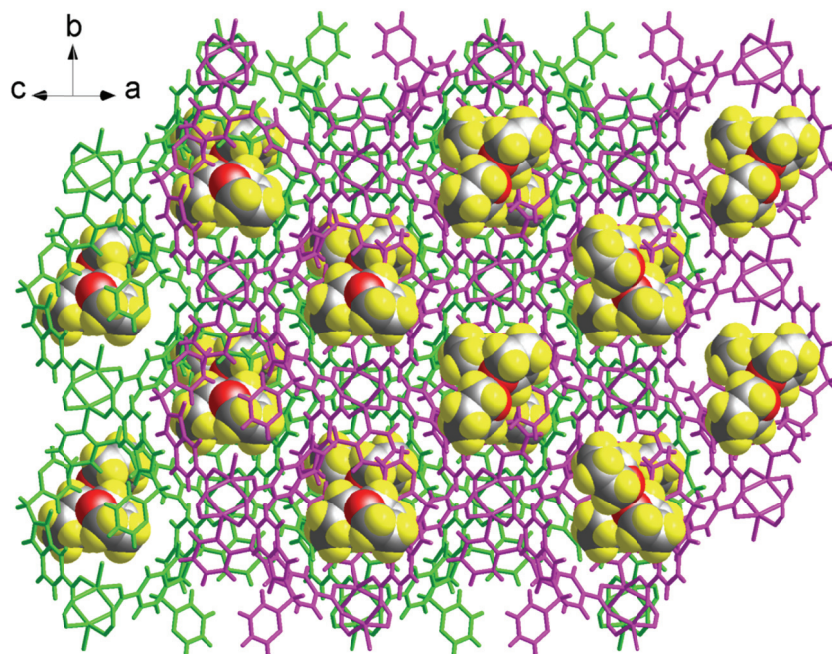


Fig. S14 Crystal packing of **1**, showing the lattice THF molecules (space-filling model) are accommodated within the cavities.

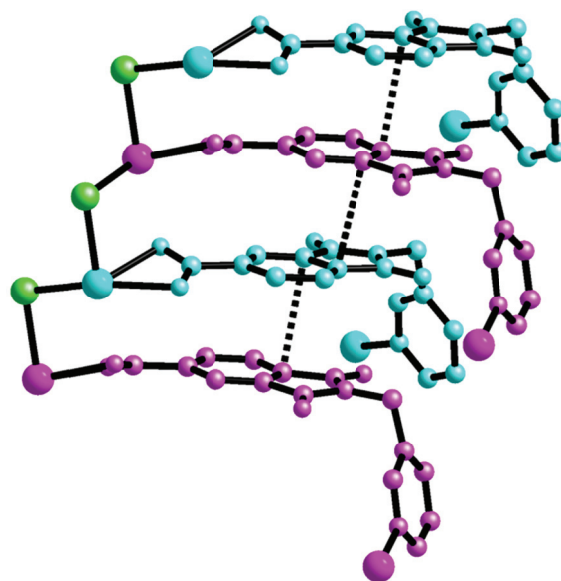


Fig. S15 Representation of the inter-helix $\pi\cdots\pi$ interactions (dashed lines, the shortest atom \cdots atom distance: 3.3106(17) Å in **4**. Carbon-bound hydrogen atoms are omitted for clarity.

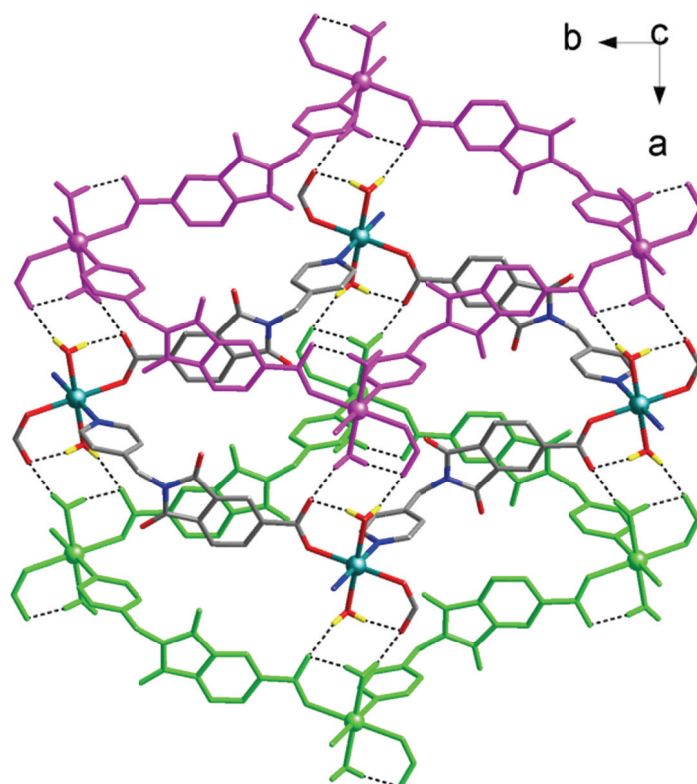


Fig. S16 Crystal packing of **5**, showing the three-dimensional hydrogen-bonded supramolecular net formed by net-to-net aqua-carboxylate hydrogen bonds (dashed lines). Carbon-bound hydrogen atoms are omitted for clarity.

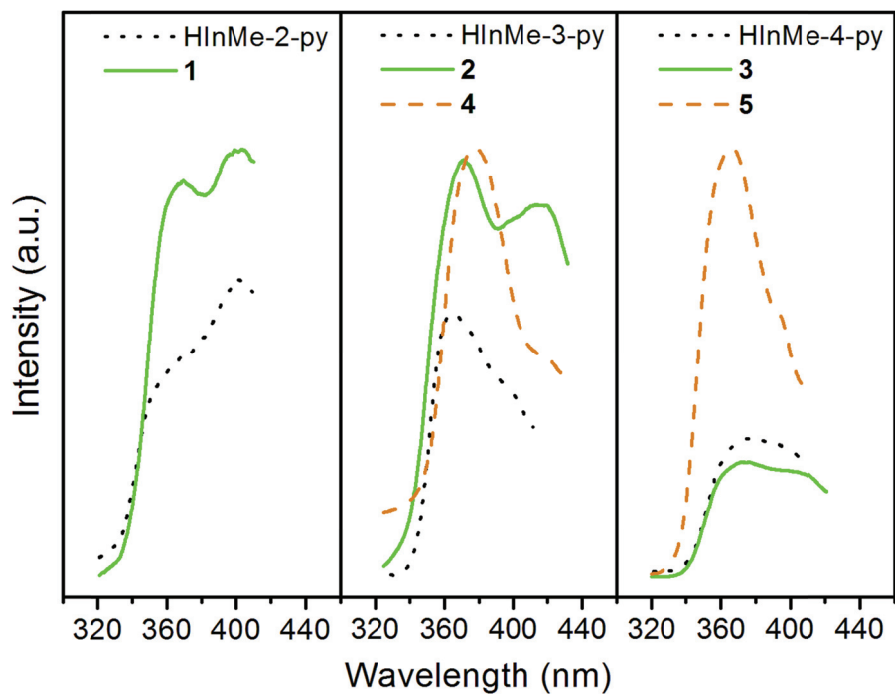


Fig. S17 Solid-state excitation spectra of HInMe- n -py ($n = 2, 3, 4$) ligands and compounds **1–5** at room temperature. *Note:* $\lambda_{em} = 450$ nm for HInMe- n -py ($n = 2, 3, 4$) ligands and **1**, $\lambda_{em} = 470$ nm for **2** and **4**, $\lambda_{em} = 460$ nm for **3**, and $\lambda_{em} = 425$ nm for **5**.