## The influence of anion, ligand geometry and stoichiometry on the structure and dimensionality of a series of Ag(I)bis(cyanobenzyl)piperazine coordination polymers

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**Supporting Information** 

θ

Compound	Torsion Angle θ (°)	Dimensionality
1	57.3(2)	2
2	51.8(4), 45.6(4)	2
3	50.4(2)	1
4	42.4(3)	2
5	176.9(3), 175.9(3)	3

Figure S1 Relationship between Ag-N-CH<sub>2</sub>-C torsion angle and dimensionality of the resulting assembly



**Figure S2** Intermolecular interactions in the structure of L1 and L2; a) Offset face-to-face  $\pi$ - $\pi$  interactions between molecules of L1; b) Offset face-to-face  $\pi$ - $\pi$  interactions between molecules of L2; c) C-H…N hydrogen bonding in the structure of L1.



Figure S3 Crystal packing of adjacent sheets in the structure of 1. Hydrogen atoms omitted for clarity.



Figure S4 Crystal packing of adjacent layers of chains in the structure of 3 viewed parallel to the direction of propagation. Hydrogen atoms omitted for clarity.

## **X-Ray Powder Diffraction**

Data collection details are provided in the X-ray crystallography section. All data collections were carried out at room temperature and compared with single crystal data collected at either 123K or 100K, leading to the expected discrepancies in exact peak positions due to thermal changes in the unit cell parameters.



Figure S5 X-ray powder diffraction pattern for L1



Figure S6 X-ray powder diffraction pattern for L2



Figure S7 X-ray powder diffraction pattern for complex 1



Figure S8 X-ray powder diffraction pattern for complex 2. We ascribe the peak broadening above  $2\theta$ = 20° to partial loss of lattice solvent during the experiment.



Figure S9 X-ray powder diffraction pattern for complex 3



Figure S10 X-ray powder diffraction pattern for the poorly crystalline material obtained from drying complex 4



Figure S11 X-ray powder diffraction pattern for complex 5



**Figure S12** X-ray powder diffraction pattern for the hexafluorophosphate analogue to complex **5**, *poly*-[Ag(L2)]·PF<sub>6</sub>, compared with the calculated pattern for complex **5**.