Guest Induced Transformations of Assembled Pyridyl Bis-Urea Macrocycles

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Guest induced transformations of assembled pyridyl bis-urea macrocycles†‡

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Pyridine macrocycles with no cavities assembled into close packed columns yet absorbed guests including hydrogen, carbon dioxide, and iodine.

Many different approaches have been used to design materials with open channels that display permanent porosity.1 Recent examples indicate that a permanent pore or channel is not necessarily a requirement for the absorption of gases.2 Dynamic structural changes in coordination polymers induced new channels into which guest molecules were incorporated.3 Atwood et al. recently demonstrated that non-porous organic solids could expand by gas induced transformations.4 In this manuscript, we report a pyridyl bis-urea host (1, Fig. 1) that assembled into columns with high fidelity. Close packing of these columns gives a high density structure with no obvious channels. Co-crystallization of host 1 with trifluoroethanol (TFE) formed a crystalline inclusion complex host 1-TFE with TFE separating the columns of assembled bis-ureas. Heating induced desorption of the TFE regenerated the initial host 1 extended structure. While there are examples of crystals that can expand to absorb guests, we were surprised to observe type I gas adsorption isotherms with H2 and CO2, which usually indicate permanent channels. Furthermore, these crystals displayed reversible absorption of iodine to afford a stable crystalline host 1-I2 complex. Closer analysis suggested guest induced transformations that converted a higher density ‘guest free’ form of bis-urea host 1 to a lower density form. We investigated these structures by X-ray diffraction, X-ray photoelectron spectroscopy (XPS) and UV-vis spectroscopy. These studies suggest that the organic solid state can be dynamic and can expand to encapsulate guests.

We have previously shown that ureas, preorganized within rigid frameworks, guided columnar assembly through the strong 3-centered urea hydrogen bonding motif.5 Macrocycles with sizeable cavities afford open columnar channels that are accessible to gases and guests; however, we expected macrocycles that lacked a cavity to be nonporous. For example, previously reported m-xylene 2 assembled into columns through typical 3-centered urea interactions,6 but the inward facing aryl hydrogens were nearly within van der Waals contact. We recently synthesized pyridine 1, which contains basic interior nitrogens with no central cavity. Macrocycle 1 was synthesized in two steps from 2,6-dibromomethylpyridine and triazinanone (ESI†). Colorless crystals of 1 were obtained by vapor diffusion of methanol into a DMSO solution of 1 (10 mg/2 mL). X-Ray analysis revealed the expected bis-urea macrocycle (Fig. 1a). Accounting for van der Waals radii, the interior of the macrocycle was ~1.1 × 1.8 Å (Fig. 1a). In comparison, gases used to measure porosity have kinetic diameters of 2.9 (H2) to 3.6 Å (N2).7 The individual macrocycles 1 were connected into columns by hydrogen bonds between the urea NH’s and two different acceptors: the urea carbonyl oxygen and the pyridine nitrogen (Fig. 1b). The carbonyl oxygen formed slightly shorter interactions with an N–O distance of 2.904(2) Å vs. the pyridine N, which displayed an N–N distance of 3.082(2) Å. This hydrogen bonding motif gave a repeat distance of ~4.7 Å, similar to the typical 3-centered urea motif.6 The chains of hydrogen bonds run along the crystallographic b axis and the columns are densely packed with no obvious channels (Fig. 1c).

Host 1 also crystallized through the vapor diffusion of TFE into a DMSO solution of 1 (10 mg/2 mL) to afford a 1-TFE complex. This structure showed columns with similar distances from the urea NHs to the heteroatoms.

Fig. 1 Macroyclic bis-ureas 1 and 2: (a) structure of host 1 with intramolecular distances from C1 to C1* = 4.645 Å and N3 to N3* = 5.066 Å; (b) view along a single column of host 1 illustrating the hydrogen bonding pattern; (c) view depicting the packing of the columns.

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isotherm with CO$_2$ (g) at 273 K consistent with microporous

Surprisingly, host porosity and were examined as controls for gas adsorption.

and intensities were similar to the original host structures, indicating that these bulk powders had the same overall structure as their crystalline solids (ESI). The PXRD pattern calculated from the respective single crystal observed PXRD patterns closely matched the theoretical volume by measuring the change in length. After 7 days a 14% increase in volume was observed (ESI).

These crystalline solids were ground to powders and examined by powder X-ray diffraction. In each case, the observed PXRD patterns closely matched the theoretical PXRD pattern calculated from the respective single crystal structures, indicating that these bulk powders had the same overall structure as their crystalline solids (ESI†). The solvent was removed from host I-TFE powder by heating and the powder re-examined by PXRD (Fig. 3a). The peak positions and intensities were similar to the original host I structure. Host I and 2 were not expected to display permanent porosity and were examined as controls for gas adsorption. Surprisingly, host I crystals displayed a type I gas adsorption isotherm with CO$_2$ (g) at 273 K consistent with microporous materials (Fig. 3b) as did host 2 (ESI†). The Brunauer–Emmet–Teller method was applied to the CO$_2$ isotherm for host I at $P/P_0$ between 0.016 and 0.028. The calculated surface area of 190 M$^2$ g$^{-1}$ corresponds to a total pore volume of 3.852 × 10$^{-3}$ cm$^3$ g$^{-1}$ for pores smaller than 6.3 Å in diameter. Host I also displayed a similar type I adsorption isotherm with H$_2$ (g) (Fig. 3b, inset). Both these gases have kinetic diameters that were larger than any void in the hosts. Thus, our hypothesis is that these hosts undergo gas induced transformations where the columns are pushed apart to accept the gas as observed in the host I-TFE inclusion complex.

The ability of host I to form inclusion complexes with TFE through hydrogen bonding interactions with the unsatisfied oxygen lone pair, led us to the consider other guests whose structures would be easier to monitor. In host 2, the lone pairs were occupied in the 3-centered urea hydrogen bonds, thus this compound was tested as a control. We next investigated if hosts I and 2 absorbed I$_2$, which was well-known to form charge transfer complexes with pyridine. The colorless crystals of hosts I and 2 were exposed to iodine vapors in a sealed container for 1–9 days. The vapor loading of iodine in host I to afford red crystals was kinetically slow, requiring ~7 days to reach equilibrium. The desorption of iodine from these crystals was followed by thermogravimetric analysis (TGA). A one step desorption with 19.18% weight loss was observed between 60 °C to 140 °C from the host I-I$_2$ complex (Fig. 4a). Assuming that this weight loss corresponded to I$_2$, we calculated a 3 : 1 host : guest ratio in the host I-I$_2$ complex. No weight loss was observed for the iodine treated host 2 samples, indicating that I$_2$ was not absorbed.

Iodine could also be loaded reproducibly from an ethanolic solution and reached a saturation absorption of 280 (±20) mg I$_2$ per gram of I. This corresponds to the same 3 : 1 binding ratio (1 : I$_2$). To further investigate this binding phenomenon, host I (55 mg) was loaded in a capillary and equilibrated with iodine vapor for 7 days. We monitored the daily increase in volume by measuring the change in length. After 7 days a 14% increase in volume was observed (ESI†). The expansion of I was presumably due to the formation of a clathrate (host I-I$_2$).

Powder X-ray diffraction (PXRD) was used to compare the structure of the host I and its I$_2$ complex. The host I-I$_2$ complex exhibited a PXRD pattern (Fig. 4b top) that was different from the free host. We observed a shift in the diffraction peak positions towards smaller 2$\theta$ values for the lower angle (102) peak from 14.1° to 12.55° and the (202) peak (21.3° to 19.6°). These changes suggested that the cell was expanding in the $a$–$c$ direction as expected for iodine bound between the columns. The iodine was removed by heating and the PXRD positions and intensities mirrored those of the initial host I structure (ESI†).

We next investigated the nature of the bound iodine species. The empty host I exhibited an absorbance maximum at

![Fig. 2](image1.png)

**Fig. 2** Views from the X-ray structure of host I-TFE: (a) the hydrogen bonding environment of a single macrocycle was nearly identical to the solvent-free structure except for a TFE molecule that was hydrogen bonded to the carbonyl; (b) view depicting the packing of the columns.

(N-O = 2.928(4) Å, N-N = 3.038(4) Å). The important difference between the two structures was the hydrogen bonding interaction between the alcohol of TFE and urea carbonyl oxygen (Fig. 2a). The TFE forms an interstitial layer between a layer of columns (Fig. 2b). There is a change in symmetry between the solvent free and solvent complex crystals from monoclinic $z$ = 2 to triclinic $z$ = 1. To compare their volumes, we divided the solvent free cell volume by 2. The host I-TFE had a larger volume (563.69 Å$^3$) vs. host I (364.39 Å$^3$), which reflected the space occupied by solvent.

![Fig. 3](image2.png)

**Fig. 3** (a) Comparison of PXRD patterns for host I (bottom), host I-TFE (middle) and host I-TFE after TFE removal (top); (b) carbon dioxide adsorption isotherms at 273 K for host I (adsorption in blue, desorption in red) and hydrogen adsorption isotherms at 273 K for host I (inset).

![Fig. 4](image3.png)

**Fig. 4** (a) TGA desorption curves for the iodine treated host I and host 2; (b) comparison of the PXRD patterns of host I and host I-I$_2$. 

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§ Crystallographic data for I: C_{16}H_{24}N_{6}O_{2}, M = 326.36, colorless block crystal, 0.24 × 0.22 × 0.10 mm, T = 150 K. Monoclinic, space group P21/c, a = 9.5207(5), b = 4.6948(3), c = 16.3045(9) Å, \( \beta = 90.20(1)^{\circ} \). \( U = 728.77(7) \) Å\(^2\), \( Z = 2 \), \( D_0 = 1.487 \text{ g cm}^{-3} \), \( F(000) = 344 \). Bruker SMART APEX CCD-based diffractometer, \( \lambda = 0.71073 \) Å (Mo-K\( \alpha \)), \( \theta_{\text{max}} = 26.37 \), 8848 reflections measured, 1489 unique, 1214 with \( I > 2\sigma(I) \), \( R_{int} = 0.0502 \). Final R-values: \( R_1 (F) \) = 0.0352; \( wR_2 (F^2, \text{all data}) = 0.0875 \). For 1-TFE: C\(_8\)H\(_8\)N\(_2\)O\(_2\)-2CF\(_2\)(CH\(_2\)OH) \( M = 526.45 \), colorless block crystal, 0.60 × 0.06 × 0.04 mm, \( T = 150 \) K. Triclinic, space group \( P1 \), \( a = 4.6955(12) \), \( b = 9.632(3) \), \( c = 12.775(3) \) Å, \( \alpha = 93.970(5) ^{\circ} \), \( \beta = 100.219(5) ^{\circ} \), \( \gamma = 95.594(5) ^{\circ} \). \( U = 563.72(2) \) Å, \( Z = 1 \), \( D_0 = 1.551 \text{ g cm}^{-3} \), \( F(000) = 272 \). Bruker SMART APEX CCD-based diffractometer, \( \lambda = 0.71073 \) Å (Mo-K\( \alpha \)), \( \theta_{\text{max}} = 24.94 ^{\circ} \), 5796 reflections measured, 1970 unique, 1372 with \( I > 2\sigma(I) \), \( R_{int} = 0.0583 \). Final R-values: \( R_1 (F) \) = 0.0684; \( wR_2 (F^2, \text{all data}) = 0.1908 \).

### Notes and references


