Host-Guest Systems

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Guest-Induced Instant and Reversible Crystal-to-Crystal Transformation of 1,4-Bis(ferrocenylethynyl)anthraquinone**

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For decades, much attention has focused on crystallization from a mixture of electron-donor (D) and -acceptor (A) molecules, forming D- and/or A-stacked column structures that exhibit unique physical properties such as ferromagnetism,[1] strong absorption of Vis/NIR light,[2] and high electronic conductivity. [3] The tendency toward DA stacking in the crystal indicates that D-A conjugated molecules can form network structures which may manifest interesting physical properties. We are interested in the chemistry of a new class of D-A conjugated systems, namely, ferrocenylethynylanthraquinones (FcAq), from two viewpoints: protonation-induced intramolecular electron transfer resulting in valence tautomers, [4] and the construction of D-A stacked nanoporous network structures, which have attracted much recent attention because of great potential for applications such as selective gas adsorption, [5-7] heterogeneous catalysts, [8,9] and molecular recognition.[10,11] In the present study, we found that the new T-shaped FcAq compound 1,4-bis(ferrocenylethynyl)anthraquinone (1,4-Fc₂Aq (1); Figure 1a) in the presence of guest solvent molecules forms single crystals, which show instant and reversible crystal-to-crystal transformation on heating or treatment with guest vapors. Recently, several reversible crystal-to-crystal transformations have been reported, [5,12] among which examples accompanying a significant conformational change are few.[13,14] The present study demonstrates an instant and reversible trans-

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Figure 1. a) Chemical structure of 1,4-Fc₂Aq (1); b) ORTEP plot of a 1,4-Fc₂Aq molecule in a crystal of α -1-CH₂Cl₂. Hydrogen atoms and disordered guest molecules are omitted for clarity.

formation despite a very large conformational change in the D-A stacked structure, which can be monitored by variable-temperature synchrotron X-ray powder diffraction (VT-XRPD).

We synthesized 1,4-Fc₂Aq (1) by Sonogashira cross-coupling of 1,4-dibromoanthraquinone with 1-ethynylferrocene in 33% yield. Crystals of 1 obtained from various solvents were analyzed by single-crystal X-ray diffraction. An ORTEP diagram of a 1,4-Fc₂Aq molecule in a crystal of α -1·CH₂Cl₂ is shown in Figure 1b. [15] In most of the structures, inclusion of solvent as guest molecule is observed. The crystal structure of α -1·CH₂Cl₂ along the *b* axis is shown in Figure 2a as an example. The crystal was obtained by recrystallization

from dichloromethane/hexane at 263 K; under these conditions, a large crystal $(25 \times 1 \times 1 \text{ mm})$ grew within a week (Figure 2 d). α-1·CH₂Cl₂ has subnanopores with dimensions of $5.7 \times 3.5 \text{ Å}^{[16]}$ along the b axis in which dichloromethane guest molecules are contained with a 1/CH₂Cl₂ molar ratio of 1:1. An infinite one-dimensional columnar structure is formed along the b axis by an alternating arrangement of a π conjugated spacer and a π - π stacking interaction between Fc (D) and Aq (A) moieties (Figure 3a). The shortest distance between a cyclopentadienyl (Cp) plane of Fc and an Aq plane is 3.44 Å. The Aq and Cp planes form angles of about 45° with the b axis. Two kinds of complex columns are almost perpendicular to each other and are arranged alternately to construct the porous framework. Crystallographic data of single crystals with the composition $1 \cdot X$ (X = guest molecule) obtained by recrystallization from different solvents are

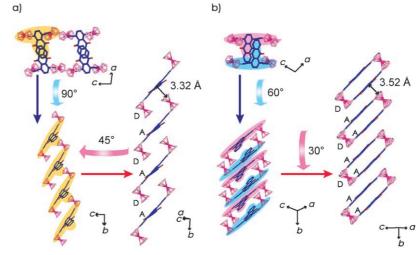


Figure 3. One-dimensional columnar structures of 1-X (a), and solvent-free 1 (b).

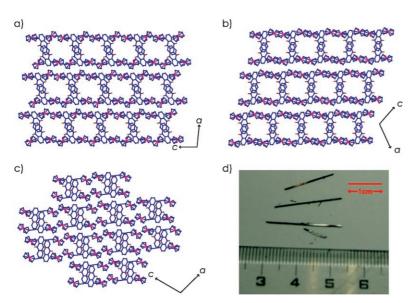


Figure 2. Crystal structures of α -1·CH₂Cl₂ along the *b* axis (a), β-1·CH₂Cl₂ along the *b* axis (b), solvent-free 1 along the *b* axis (c), and a photograph of α -1·CH₂Cl₂ crystals (d). Hydrogen atoms and disordered guest molecules (CH₂Cl₂) are omitted for clarity.

summarized in the Supporting Information.^[15] Their crystal structures are similar to that of α-1·CH₂Cl₂, but the size of the subnanopores changes according to the size of the included guest molecules. Two types of crystals (α and β phases in Figure 2a and b; see the Supporting Information) were obtained when X was dichloromethane, chloroform, or carbon tetrachloride. The guest molecules in these structures are extremely disordered. The α phase has a shorter plane-to-plane distance between Fc and Aq but a larger dihedral angle ϕ between the Cp ring of ferrocene and the Aq plane (C25–C29, and C38) than the β phase (see the Supporting Information). Because of the balance between these two factors, that is, the shorter distance stabilizing the α phase and the large ϕ destabilizing it, both α and β phases may exist under the same conditions. Only one type of structure was formed when X was a larger guest molecule such as THF, trichloroethylene, or hexane.

Recrystallization of **1** from *o*-dichlorobenzene/hexane at 263 K afforded crystals containing *o*-dichlorobenzene and hexane guests in an ordered arrangement

(see the Supporting Information). In contrast, when 1 was recrystallized at 293 K from o-dichlorobenzene/hexane, crystals without pores and solvent molecules were obtained (Figure 2c).[15] In the solvent-free form, a columnar structure is maintained (Figure 3b), whereas the syn conformation of the ferrocene moieties in 1 allows the formation of pairs of adjacent molecules in the crystal and the pattern of the stacking arrangement changes from D-A to D-A-A compared with the porous 1.X form.

Thermal desorption of the guest molecules in 1.X was examined by thermogravimetric analysis (TGA). The TG curve indicated desorption of guest molecules at $T_{\rm de} = 346$ -380 K without chemical decomposition. The $T_{\rm de}$ values do not correlate with the boiling points of the guest molecules (e.g., the most volatile dichloromethane had high $T_{\rm de}$), that is, the host-guest interaction affects T_{de} significantly.

To clarify the crystal structure after desorption of guest molecules, the synchrotron X-ray powder diffraction (XRPD) pattern of a sample prepared by heating 1. THF at 420 K under vacuum for 12 h was measured, and the cell parameters of the sample were determined by the Rietveld method. The XRPD-derived cell parameters were consistent with those of solvent-free 1 (see the Supporting Information). The XRPD patterns of samples of 1·CH₂Cl₂ from which guests had been desorbed thermally were also the same as those of solvent-free 1. Thus, it is confirmed that guest desorption causes crystal-to-crystal transformation from 1:X to solvent-

Reversibility of the crystal-to-crystal transformation on guest desorption and adsorption by 1 was studied by variabletemperature XRPD (VT-XRPD). For this measurement, we prepared a vacuum-sealed 1-mm-diameter glass capillary containing a 0.5-mm-diameter capillary filled with frozen THF and a 0.5-mm-diameter capillary packed with powdered solvent-free 1 (Figure 4a). After 3 h at room temperature, the XRPD pattern had completely changed from that of solventfree 1 to that of 1. THF (Figure 4b, $A\rightarrow B$). Then, THF was frozen at 90 K and the sample was heated from 300 to 400 K. Subsequently, the temperature of the sample capillary T_s was maintained at 300 K, and the temperature of the THF capillary $T_{\rm g}$ was increased from 90 to 400 K. XRPD patterns were measured at every 50 K with an X-ray exposure time of 5 min for each measurement. On heating a sample of 1 THF, THF molecules were released, the XRPD pattern of solventfree 1 emerged, and, at 400 K, the pattern corresponded completely to that of solvent-free 1 (Figure 4b, C). When the capillary was filled with THF vapor by increasing $T_{\rm s}$, THF molecules were adsorbed by the host, and XRPD patterns arising from 1. THF were observed (Figure 4b, D-G). When $T_{\rm s}$ and $T_{\rm g}$ were kept at 300 K, guest adsorption was complete within 5 min (see the Supporting Information).

These results indicate rapid and reversible crystal-tocrystal transformation between 1.THF and solvent-free 1. In view of the conformational change required, the fast transformation is particularly remarkable.^[14] These features are derived from the flexibility of the crystal structure as a result of the D-A interaction, which is weaker than other chemical bonds, and from the higher stability of 1. THF compared with solvent-free **1** ($\Delta G = -32 \text{ kJ mol}^{-1}$ [17]). Diffuse reflectance

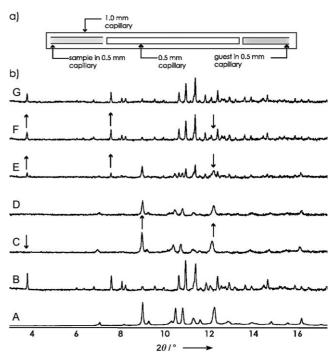


Figure 4. a) Illustration of the capillary used in VT-XRPD experiment. b) XRPD pattern of solvent-free 1 (A) and results of VT-XRPD measurements (B-G). Temperatures of sample (T_s) and guest (T_g) were B: $T_s = 350$, $T_g = 90$ K; C: $T_s = 400$, $T_g = 90$ K; D: $T_s = 300$, $T_g = 200$ K; E: $T_s = 300$, $T_g = 300$ K; F: $T_s = 300$, $T_g = 350$ K; and G: $T_s = 300$, $T_g = 400$ K.

UV/Vis spectra of 1:THF and solvent-free 1 indicated appearance of a CT band that is not present in the spectrum of 1 in solution and a red shift of the CT band for 1. THF $(\lambda_{\text{max}} = 619 \text{ nm})$ from that of solvent-free **1** $(\lambda_{\text{max}} = 601 \text{ nm})$; see the Supporting Information). This result suggests stronger D-A interaction in 1.X, which is consistent with the onedimensional columnar D-A structure and the shorter Cp-Aq distance compared with that in solvent-free 1.

In conclusion, the new D-A conjugated compound 1,4-Fc₂Aq (1) has been synthesized. Crystals of 1 have either a solvent-incorporating porous structure or a solvent-free nonporous structure, which can be instantly and reversibly interconverted by desorption and adsorption of guest molecules, with alternation of D-A and D-A-A arrangements in the one-dimensional columnar structure. A preliminary study has shown that 1 exhibits protonation-induced intramolecular electron transfer. We are currently investigating the combination of physical property changes resulting from the interaction with acid molecules and the crystal structural transformation.

Experimental Section

1: 1,4-Dibromoanthraquinone (2.42 g, 6.6 mmol) and 1-ethynylferrocene (2.831 g, 13.5 mol) were dispersed in triethylamine (120 mL) and stirred. CuI (136 mg, 0.72 mmol) and [PdCl₂(PPh₃)₂] (482 mg, 0.69 mmol) were added. The suspension was heated to reflux for 3 h. The color of the reaction mixture turned from orange to dark red.

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The solvent was removed under vacuum. The residue was dissolved in 100 mL of dichloromethane, washed with water, and then dried over anhydrous Na₂SO₄. The filtrate was subjected to chromatography on an alumina column (activity II–III, 6×20 cm) with dichloromethane/hexane (1:3 \rightarrow 1:1). The third brown band was collected. The product was recrystallized from dichloromethane/hexane, and black crystals were collected. Yield: 1.3442 g (32.6 %). ¹H NMR (400 MHz, CDCl₃): δ = 4.34 (t, 4 H, J = 1.8 Hz; Cp), 4.35 (s, 10 H; Cp), 4.67 (t, 4 H, J = 1.8 Hz; Cp), 7.78 (m, 2 H; Ph), 7.81 (s, 2 H; Ph), 8.34 ppm (s, 2 H; Ph). ¹³C NMR (125 MHz, CD₂Cl₂): δ = 65.4 (Cp), 70.2 (Cp), 70.8 (Cp), 72.4 (Cp), 86.9 (C=C), 98.1 (C=C), 123.9 (Aq), 127.4 (Aq), 134.1 (Aq), 134.8 (Aq), 139.0 (Aq), 182.4 ppm (C=O). MS (ESI-TOF-MS): m/z 623.9541; calcd: 624.29 [M⁺].

X-ray diffraction measurements: The crystals were mounted in a loop. Data was collected with a Rigaku AFC8 diffractometer with the Rigaku Mercury CCD system equipped with a rotating-anode X-ray generator producing graphite-monochromated $Mo_{K\alpha}$ radiation (λ = 0.7107 Å). An empirical absorption correction using equivalent reflections and Lorentzian polarization were performed with the program Crystal Clear 1.3.5. The structure was solved with the program SHELXS-97^[18] and refined against F^2 using SHELXL-97^[19] Selected crystallographic data and experimental details are listed in the Supporting Information.

The in situ synchrotron powder diffraction experiment on THF adsorption was performed on a large Debye–Scherrer camera installed at SPring-8 BL02B2 with an imaging plate as detector. The wavelength of the incident X-rays was 1.001 Å. VT-XRPD measurements were performed with a nitrogen gas-flow system.

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