

Electronic Supplementary Information (ESI) for

Drastic rearrangement of self-assembled hydrogen-bonded tapes in a molecular crystal

Masaki Donoshita,^a Mikihiro Hayashi,^a Ryuichi Ikeda,^a Yukihiro Yoshida,^a
Shota Morikawa,^a Kunihisa Sugimoto,^b and Hiroshi Kitagawa^{*,a}

^aDivision of Chemistry, Graduate School of Science, Kyoto University, Kitashirakawa-Oiwakecho, Sakyo-ku, Kyoto 606-8502, Japan

^bJapan Synchrotron Radiation Research Institute (JASRI), SPring-8, 1-1-1 Kouto, Sayo-cho, Sayo-gun, Hyogo 679-5148, Japan

*E-mail: kitagawa@kuchem.kyoto-u.ac.jp

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Experimental Details

Sample preparation: Chloranilic acid (CA) was obtained from TCI Chemical Industry (Tokyo, Japan) and purified by recrystallization from aqueous solution. 2-Pyrrolidone (2-Py) was obtained from Nacalai Tesque (Kyoto, Japan) and used as received without further purification. Orange plate crystals of High-A were prepared by rapid cooling of a hot acetonitrile solution (120 mL) containing CA (2.12 g, 10.2 mmol) and 2-Py (3.40 g, 39.9 mmol) to room temperature (5.5 h), while stirring. Red rhombic block crystals of High-B were prepared by slow evaporation of an acetonitrile solution (1.01 L) containing CA (2.09 g, 10.0 mmol) and 2-Py (3.41 g, 40.0 mmol) at room temperature.

Single-crystal X-ray structural analysis: Crystallographic data at typical temperatures were collected using the synchrotron X-ray at the BL02B1 beam-line in SPring-8 with a Rigaku Mercury 2 CCD detector ($\lambda = 0.703010 \text{ \AA}$).¹ Variable-temperature experiments were performed on a laboratory X-ray diffractometer (Rigaku XtaLAB P200) with graphite-monochromated Mo K α radiation ($\lambda = 0.71073 \text{ \AA}$). A single crystal was mounted on a glass fibre with epoxy resin and cooled by a stream of cooled nitrogen gas. The structures were solved by direct methods (SHELXT²) and refined by full-matrix least-squares refinement on F^2 (SHELXL³) using the CrystalStructure⁴ software package. The positions of all atoms, including hydrogen atoms, were refined in the general least-squares cycle with no restraints. All non-hydrogen atoms were refined anisotropically, whereas hydrogen atoms were refined isotropically.

(1) Crystallographic data were deposited with the Cambridge Crystallographic Data Centre: deposition nos. CCDC 1838517 (High-A at 298 K), 1838519 (Low-B at 100 K), 1838518 (High-B at 298 K) and 1838516 (High-B at 100 K).

(2) G. M. Sheldrick, *Acta Crystallogr.*, 2015, **A71**, 3.

(3) G. M. Sheldrick, *Acta Crystallogr.*, 2008, **A64**, 112.

(4) CrystalStructure 4.2.5: Crystal Structure Analysis Package, Rigaku Corporation, Tokyo, Japan.

Table S1. Cell parameters and crystallographic information for each form of 2:1 co-crystal of 2-pyrrolidone and chloranilic acid

| | High-A | Low-B ^a | High-B ^a | High-B ^a |
|--|---|---|---|---|
| Formula | C ₁₄ H ₁₆ Cl ₂ N ₂ O ₆ | C ₁₄ H ₁₆ Cl ₂ N ₂ O ₆ | C ₁₄ H ₁₆ Cl ₂ N ₂ O ₆ | C ₁₄ H ₁₆ Cl ₂ N ₂ O ₆ |
| Formula weight | 379.20 | 379.20 | 379.20 | 379.20 |
| Temperature (K) | 298 | 100 | 298 | 100 |
| Wavelength (Å) | 0.703010 | 0.703010 | 0.703010 | 0.703010 |
| Crystal system | Triclinic | Triclinic | Triclinic | Triclinic |
| Space group | <i>P</i> -1 | <i>P</i> -1 | <i>P</i> -1 | <i>P</i> -1 |
| Colour | Orange | Orange | Orange | Orange |
| <i>a</i> (Å) | 8.1575(9) | 4.6904(17) | 4.987(3) | 4.9143(16) |
| <i>b</i> (Å) | 10.3671(12) | 8.921(3) | 7.592(4) | 7.458(2) |
| <i>c</i> (Å) | 11.0921(13) | 9.833(3) | 11.718(6) | 11.698(4) |
| α (°) | 74.290(5) | 103.357(7) | 72.926(6) | 73.270(5) |
| β (°) | 75.655(5) | 96.447(7) | 89.449(6) | 89.528(6) |
| γ (°) | 67.442(5) | 93.984(7) | 71.157(6) | 69.737(5) |
| <i>V</i> (Å ³) | 822.93(17) | 395.8(2) | 399.6(4) | 383.2(2) |
| <i>Z</i> | 2 | 1 | 1 | 1 |
| Calcd density (g cm ⁻³) | 1.530 | 1.591 | 1.576 | 1.643 |
| μ (mm ⁻¹) | 0.410 | 0.426 | 0.422 | 0.440 |
| <i>F</i> (000) | 392 | 196 | 196 | 196 |
| Crystal size (mm ³) | 0.09 × 0.04 × 0.01 | 0.14 × 0.06 × 0.02 | 0.14 × 0.06 × 0.02 | 0.14 × 0.06 × 0.02 |
| Total reflections | 10883 | 5200 | 5228 | 4220 |
| Unique reflections | 3744 | 1799 | 1821 | 1393 |
| Parameters used | 281 | 141 | 141 | 141 |
| <i>R</i> _{int} | 0.0603 | 0.0830 | 0.0658 | 0.0613 |
| Goodness-of-fit | 0.946 | 1.028 | 0.944 | 1.065 |
| <i>R</i> ₁ [<i>I</i> > 2σ(<i>I</i>)] | 0.0491 | 0.0535 | 0.0520 | 0.0434 |
| <i>wR</i> ₂ (all reflections) | 0.1333 | 0.1428 | 0.1302 | 0.1120 |
| max, min Δρ (e Å ⁻³) | 0.25, -0.24 | 0.60, -0.60 | 0.31, -0.23 | 0.38, -0.38 |

^a Data taken from the same piece of single crystal during the temperature-sweeping process.