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Md. Akhtaruzzaman,^a Masaaki Tomura^b* and Yoshiro Yamashita^c

^aDepartment of Structural Molecular Science, The Graduate University for Advanced Studies, Myodaiji, Okazaki 444-8585, Japan, ^bInstitute for Molecular Science, Myodaiji, Okazaki 444-8585, Japan, and ^cDepartment of Electronic Chemistry, Interdisciplinary Graduate School of Science and Engineering, Tokyo Institute of Technology, Nagatsuta, Midori-ku, Yokohama 226-8502, Japan

Correspondence e-mail: tomura@ims.ac.jp

Key indicators

Single-crystal X-ray study T = 296 KMean $\sigma(C-C) = 0.005 \text{ Å}$ R factor = 0.053 wR factor = 0.161 Data-to-parameter ratio = 12.0

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

One-dimensional hydrogen-bonded molecular tapes in 1,4-bis[(4-pyridinio)ethynyl]benzene chloranilate

The structure of the title compound, $C_{20}H_{14}N_2^{2+}C_6Cl_2O_4^{2-}$, contains one-dimensional hydrogen-bonded molecular tapes along the [113] direction. The molecular tapes are connected *via* two N-H···O hydrogen bonds in $R_1^2(5)$ patterns.

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Comment

We have recently shown that the simple combination of chloranilic acid and dipyridyl-type ligands can create a variety of supramolecular architectures involving infinite onedimensional molecular tape structures (Zaman et al., 1999, 2000). In the course of our crystal engineering studies on chloranilic acid, we have obtained the title compound, (I), a 1:1 co-crystal of chloranilic acid and 1,4-bis[(4-pyridyl)ethynyl]benzene. A long rigid and conjugated bridging ligand has received much current interest for the construction of selfassembling macrocyclic architectures (Lehn, 1995; Fujita, 1999) and the development of molecular wires (Tour, 1996). However, we have found only two examples (Lin et al., 1995, 1998) of structures containing a 1,4-bis[(4-pyridyl)ethynyl]benzene unit in the Cambridge Structural Database (Allen & Kennard, 1993). We report here the structure of (I) with hydrogen-bonded molecular tapes.



The molecular structure of (I) is shown in Fig. 1, and both molecules are located on inversion centers. A one-dimensional molecular tape is observed in the structure of (I), as shown in Fig. 2. The molecular tape is nearly flat. The angles between the molecular planes of the chloranilate and the pyridinium ring, and of the pyridinium ring and the benzene ring are 7.3 (2) and 11.8 (4) $^{\circ}$, respectively. The molecular tapes are connected via $R_1^2(5)$ couplings with two intermolecular N- $H \cdots O$ hydrogen bonds (Table 1), where both H atoms of chloranilic acid have transfered to the pyridine rings. In previous work (Zaman et al. 2000), we have shown that the flat molecular tapes form segregated stacks of each molecule. However, overlaps between the chloranilate-pyridinium ringbenzene ring-pyridinium ring-chloranilate are observed in the stacks of the molecular tapes of (I). A short $C-Cl\cdots\pi$ interaction [Cl1...(C7=C8) 3.440 (7), Cl1...C7 3.503 (4), $Cl1 \cdot \cdot \cdot C8$ 3.480 (4) Å; $Cl1 \cdot \cdot \cdot C7 \equiv C8$ 79.1 (3) and

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Figure 1

The molecular structure of (I) with the atomic numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.



Figure 2 Packing diagram of (I). Dotted lines show the intermolecular N-H···O hydrogen bonds.

 $Cl1 \cdots C8 \equiv C7 \ 81.3 \ (3)^{\circ}$ exists between the stacks of the molecular tapes (Reddy et al., 1996; Prasanna & Row, 2000). It is 1.7% shorter than the sum of the van der Waals radii of Cl and Csp^2 (Pauling, 1960).

Experimental

1,4-Bis[(4-pyridyl)ethynyl]benzene was prepared according to the literature (Lin et al., 1995). Chloranilic acid was commercially available and purified by the usual method. A diffusion method for a solution of 1,4-bis[(4-pyridyl)ethynyl]benzene (0.02 mmol) and chloranilic acid (0.02 mmol) in acetonitrile (5 ml) using an H-tube gave crystals of (I) suitable for X-ray analysis.

Crystal data

7 1
Z = 1
$D_x = 1.514 \text{ Mg m}^{-3}$
Cu $K\alpha$ radiation
Cell parameters from 23
reflections
$\theta = 5.9 - 43.0^{\circ}$
$\mu = 3.05 \text{ mm}^{-1}$
T = 296 (2) K
Needle, dark red
$0.7 \times 0.1 \times 0.1 \text{ mm}$

 $R_{\rm int} = 0.041$ $\theta_{\rm max} = 74.3^{\circ}$

 $h = -9 \rightarrow 9$

 $l = 0 \rightarrow 10$

 $k = -10 \rightarrow 10$

3 standard reflections

every 120 reflections

intensity decay: 2.5%

Data collection

Enraf-Nonius CAD-4 diffractometer ω -2 θ scans Absorption correction: ψ scan (North et al., 1968) $T_{\rm min}=0.224,\;T_{\rm max}=0.750$ 2340 measured reflections 2187 independent reflections 1213 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2 182 parameters $R[F^2 > 2\sigma(F^2)] = 0.053$ All H-atom parameters refined $wR(F^2) = 0.161$ $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.29 \text{ e } \text{\AA}^{-3}$ $\Delta \rho_{\rm min} = -0.37 \text{ e } \text{\AA}^{-3}$ S = 0.992187 reflections

Table 1

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$\begin{array}{c} N1 - H1 \cdots O1^{i} \\ N1 - H1 \cdots O2^{i} \end{array}$	1.05 (7)	2.23 (6)	2.897 (4)	120 (5)
	1.05 (7)	1.62 (7)	2.609 (3)	154 (5)

Symmetry code: (i) -x, 1 - y, -z.

All H atoms were located in the Fourier map and were refined isotropically.

Data collection: CAD-4 EXPRESS (Enraf-Nonius, 1992); cell refinement: CAD-4 EXPRESS; data reduction: HELENA (Spek, 1997); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPIII (Burnett & Johnson, 1996); software used to prepare material for publication: SHELXL97.

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