Received 9 May 2006

Accepted 30 May 2006

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 298 KMean σ (C–C) = 0.003 Å R factor = 0.051 wR factor = 0.111 Data-to-parameter ratio = 11.7

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

The inner salt 4-carboxy-2-(pyridinium-4-yl)-1*H*-imidazole-5-carboxylate monohydrate

In the title compound, $C_{10}H_7N_3O_4$ · H_2O , one carboxyl group is deprotonated and the pyridyl group is protonated. The inner salt molecule has a planar structure, apart from the carboxylic acid group, which is tilted from the imidazole plane by a small dihedral angle of 7.3 (3)°.

Comment

Imidazole-4,5-dicarboxylate has been used to prepare metal complex polymers (Lu *et al.*, 2006). Recently, we prepared the title compound, (I), and we present its crystal structure here.



The molecular structure of (I) is shown in Fig. 1. The C1containing carboxyl group is deprotonated (Table 1) and forms an intramolecular hydrogen bond with the neighboring C4-containing carboxyl group. The molecule displays a planar structure, apart from the C4-carboxyl group, which is tilted from the imidazole plane by a small dihedral angle of 7.3 (3)°.

Intermolecular $O-H\cdots O$ and $N-H\cdots O$ hydrogen bonding occurs (Table 2), helping to stabilize the crystal structure.

Experimental

2-(4-Pyridyl)benzimidazole (1 g, 5.13 mmol) was prepared in accordance with the literature method (Alcalde *et al.*, 1992), and was then added to 98% H_2SO_4 (8.4 ml) in portions. An H_2O_2 solution (30%, 8 ml) was added dropwise to the above solution at 373 K. The resulting solution was then stirred for 1 h at 413 K. After cooling to 313 K, water (200 ml) was added. The precipitated product was filtered off, washed with water and dried, yielding a light-yellow powder (0.68 g). A suspension of the yellow powder (5 mg, 0.02 mmol) in water (2 ml) was sealed in a 6 ml glass tube and heated at 423 K for 3 d, then cooled to room temperature to obtain suitable single crystals of (I).

Crystal data $C_{10}H_7N_3O_4 \cdot H_2O$ $M_r = 251.20$ Monoclinic, $P2_1/c$ a = 5.6348 (15) Å b = 8.457 (2) Å c = 21.605 (6) Å $\beta = 94.889$ (4)° V = 1025.7 (5) Å³

Z = 4 $D_x = 1.627 \text{ Mg m}^{-3}$ Mo K\alpha radiation $\mu = 0.13 \text{ mm}^{-1}$ T = 298 (2) KPlatelet, yellow $0.26 \times 0.24 \times 0.03 \text{ mm}$

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Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Bruker, 2000) $T_{\min} = 0.966, T_{\max} = 0.990$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.051$ $wR(F^2) = 0.111$ S = 1.121912 reflections 163 parameters H-atom parameters constrained 5216 measured reflections 1912 independent reflections 1541 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.034$ $\theta_{\text{max}} = 25.5^{\circ}$

$$\begin{split} w &= 1/[\sigma^2(F_o^2) + (0.0437P)^2 \\ &+ 0.2198P] \\ \text{where } P &= (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\text{max}} &= 0.001 \\ \Delta\rho_{\text{max}} &= 0.24 \text{ e } \text{\AA}^{-3} \\ \Delta\rho_{\text{min}} &= -0.23 \text{ e } \text{\AA}^{-3} \end{split}$$

Table 1

Selected bond lengths (Å).

C1-O3	1.270 (3)	C4-O1	1.213 (3)
C1-O4	1.230 (3)	C4-O2	1.300 (3)

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
N1-H1N···O5 ⁱ	0.98	1.80	2.758 (2)	164
N3-H3N···O4 ⁱⁱ	0.94	1.84	2.779 (2)	174
$O2-H2A\cdots O3$	0.96	1.51	2.462 (2)	173
$O5-H5B\cdots O1^{iii}$	0.90	1.90	2.759 (2)	161
$O5-H5A\cdots O3^{iv}$	0.88	1.90	2.768 (2)	172

Symmetry codes: (i) -x + 1, -y + 2, -z + 1; (ii) -x + 2, -y + 1, -z + 1; (iii) x + 1, y, z; (iv) $-x + 2, y + \frac{1}{2}, -z + \frac{3}{2}$.

H atoms attached to N and O atoms were located in a difference Fourier map and refined as riding in their as-found relative positions, with $U_{iso}(H) = 1.5U_{eq}(O,N)$. Other H atoms were placed in calculated positions with C-H = 0.93 Å and refined as riding, with $U_{iso}(H) = 1.2U_{iso}(C)$.



Figure 1

The molecular structure of (I), with 50% probability displacement ellipsoids (arbitrary spheres for H atoms).

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2000); software used to prepare material for publication: *SHELXTL*.

The authors thank the Natural Science Foundation of China (grant Nos. 20174023 and 20371030) and the Natural Science Foundation of Shandong Province of China (grant Nos. Z2001B01 and Z2004B01) for financial support.

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