Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 298 K Mean σ (C–C) = 0.004 Å R factor = 0.058 wR factor = 0.167 Data-to-parameter ratio = 11.8

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

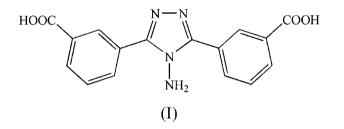
3,3'-(4-Amino-4*H*-1,2,4-triazole-3,5-diyl)dibenzoic acid

The title compound, $C_{16}H_{12}N_4O_4$, has a non-planar molecular structure. An intermolecular hydrogen-bonding network helps to stabilize the crystal structure.

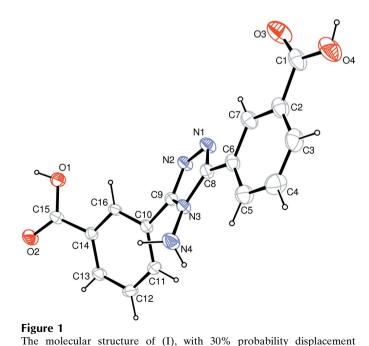
Received 13 May 2006 Accepted 1 June 2006

Comment

Dicarboxylates have been used in preparing coordination polymers (Yaghi *et al.*, 1995). As part of our ongoing investigation of polymeric complexes (Dong *et al.*, 2005), we recently prepared the title dicarboxylic acid, (I), involving a 1,2,4triazole group, and have determined its crystal structure.



The molecular structure of (I) is shown in Fig. 1. The three aromatic rings in (I) are twisted relative to each other: the dihedral angles between the triazole and C2-benzene rings and between the triazole and C10-benzene rings are 16.91 (17) and 46.68 (16)°, respectively.



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ellipsoids (arbitrary spheres for H atoms).

organic papers

An intermolecular hydrogen-bonding network helps to stabilize the crystal structure.

Experimental

3-Cyanobenzoic acid (2.2 g, 15 mmol), hydrazine sulfate (1.95 g, 15 mmol), 80% hydrazine hydrate (2.8 g, 45 mmol) and glycol (8 ml) were mixed under a nitrogen atmosphere and heated at 403 K for 3 h. The mixture was then cooled to room temperature and poured into water (50 ml); 35% HCl solution (2.5 ml) was then added to yield a white precipitate (2.0 g, 90%). The precipitate (16.2 mg, 0.05 mol) and water (2 ml) were sealed in a 6 ml glass tube and heated at 453 K for 3 d, then cooled to room temperature to obtain single crystals of (I).

Z = 8

 $D_x = 1.482 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation $\mu = 0.11 \text{ mm}^{-1}$

Block, colourless

 $0.32 \times 0.12 \times 0.08 \text{ mm}$

T = 298 (2) K

Crystal data

 $C_{16}H_{12}N_4O_4$ $M_r = 324.30$ Monoclinic, C2/c a = 36.92 (1) Å b = 7.466 (2) Å c = 10.868 (3) Å $\beta = 104.038$ (4)° V = 2906.1 (13) Å³

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: none 6351 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.058$ $wR(F^2) = 0.167$ S = 1.062572 reflections 218 parameters $R_{\rm int} = 0.046$ $\theta_{\rm max} = 25.2^{\circ}$

2572 independent reflections

1726 reflections with $I > 2\sigma(I)$

H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.0808P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} = 0.001$ $\Delta\rho_{max} = 0.20 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{min} = -0.19 \text{ e} \text{ Å}^{-3}$

Table 1

Hydrogen-bond geometry (Å, $^{\circ}$).

$D-\mathrm{H}\cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots \mathbf{A}$
	0.82	1.84	2.635 (3)	164
	0.80	1.84	2.639 (4)	174
	0.96	2.27	3.216 (3)	168
	0.86	2.42	3.194 (4)	150

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

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

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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