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Key indicators

Single-crystal X-ray study T = 291 KMean σ (C–C) = 0.002 Å R factor = 0.037 wR factor = 0.103 Data-to-parameter ratio = 12.1

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

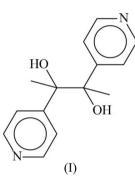
2,3-Di-4-pyridylbutane-2,3-diol

The molecule of the title compound, $C_{14}H_{16}N_2O_2$, lies on a centre of inversion located at the mid-point of the central C-C bond. O-H···N hydrogen bonds link adjacent molecules into a layer motif.

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Comment

Following the study of a copper complex of di(4-pyridyl)glycol (Kitaura *et al.*, 2002), we have used the related title ligand, (I), in an unsuccessful attempt to synthesize its adduct with cadmium iodide. An earlier study describes the isolation of the 1,4-(4*H*-1,2,4-triazol-1-methyl)benzene adduct (Niu *et al.*, 2005).



The molecule of (I) lies on a centre of inversion at the midpoint of the C2–C2ⁱ bond (Fig. 1). O–H···N hydrogen bonds $[O1-H··N1^i = 2.829 (2) \text{ Å}$; symmetry code: (i) $x, \frac{3}{2} - y, z - \frac{1}{2}]$ link adjacent molecules into a layer motif (Fig. 2). The central C–C single bond is surprisingly long [1.591 (3) Å]; the geometry-optimized structure, as calculated by *GAUSSIAN98* (Frisch *et al.*, 1998), also has a similarly long bond [1.603 Å]. On the other hand, the C–C_{methyl} distances in the crystal structure and geometry-optimized structure are normal (Table 1).

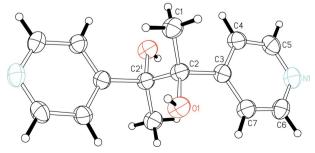


Figure 1

A plot of (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as spheres of arbitrary radii [Symmetry code (i) and for unlabelled atoms: 1 - x, 1 - y, 1 - z.]

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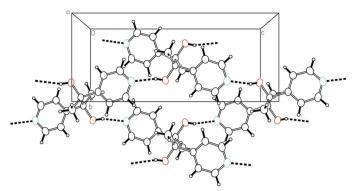


Figure 2

A plot showing the hydrogen-bonded layer structure in (I). Hydrogen bonds are shown as dashed lines.

Experimental

The title compound, which was purchased from Fluka, remained unreacted in an attempt to synthesize the adduct with cadmium iodide. Cadmium iodide (0.037 g, 0.1 mmol) and the reagent (0.024 g, 0.1 mmol) were dissolved in dimethyl sulfoxide (10 ml). Colourless crystals of (I) deposited from the solution after several days. For the density functional theory quantum-chemical calculations, the ground state electronic state was calculated at the B3LYP/6–31G* level using the *GAUSSIAN98* program suite (Frisch *et al.*, 1998).

Crystal data

 $\begin{array}{l} C_{14}H_{16}N_2O_2\\ M_r = 244.29\\ Orthorhombic, Pbca\\ a = 12.408 \ (1) \ \text{\AA}\\ b = 6.4174 \ (6) \ \text{\AA}\\ c = 15.004 \ (2) \ \text{\AA}\\ V = 1194.8 \ (2) \ \text{\AA}^3\\ Z = 4\\ D_x = 1.358 \ \text{Mg m}^{-3} \end{array}$

Data collection

Bruker APEX-II area-detector diffractometer φ and ω scans Absorption correction: none 6611 measured reflections 1376 independent reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.037$ $wR(F^2) = 0.103$ S = 1.021376 reflections 114 parameters All H-atom parameters refined Mo $K\alpha$ radiation Cell parameters from 1618 reflections $\theta = 2.7-26.3^{\circ}$ $\mu = 0.09 \text{ mm}^{-1}$ T = 291 (2) K Block, colourless $0.26 \times 0.14 \times 0.06 \text{ mm}$

1049 reflections with $I > 2\sigma(I)$ $R_{int} = 0.030$ $\theta_{max} = 27.5^{\circ}$ $h = -11 \rightarrow 16$ $k = -8 \rightarrow 7$ $l = -18 \rightarrow 19$

$$\begin{split} &w = 1/[\sigma^2(F_o^{-2}) + (0.0495P)^2 \\ &+ 0.2502P] \\ &\text{where } P = (F_o^{-2} + 2F_c^{-2})/3 \\ (\Delta/\sigma)_{\text{max}} = 0.001 \\ \Delta\rho_{\text{max}} = 0.21 \text{ e } \text{ Å}^{-3} \\ \Delta\rho_{\text{min}} = -0.15 \text{ e } \text{ Å}^{-3} \end{split}$$

Table 1		
Selected geometric parameters	(Å,	°)

1.421 (2)	$C2-C2^{i}$	1.591 (3)
1.329 (2)	C3-C7	1.386 (2)
1.335 (2)	C3-C4	1.392 (2)
1.526 (2)	C4-C5	1.386 (2)
1.532 (2)	C6-C7	1.381 (2)
116.2 (1)	C7-C3-C4	116.8 (1)
110.8 (1)	C7-C3-C2	120.8 (1)
107.1 (1)	C4-C3-C2	122.4 (1)
110.6 (1)	C5-C4-C3	119.3 (1)
108.0 (1)	N1-C5-C4	124.0 (1)
111.6 (1)	N1-C6-C7	124.1 (1)
108.6 (1)	C6-C7-C3	119.5 (1)
	1.329 (2) 1.335 (2) 1.526 (2) 1.532 (2) 116.2 (1) 110.8 (1) 107.1 (1) 110.6 (1) 108.0 (1) 111.6 (1)	$\begin{array}{cccccccccccccccccccccccccccccccccccc$

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

The diffraction measurements were of a quality that allowed all H atom parameters to be refined, albeit with O-H = 0.85 (1) and C-H = 0.95 (1) Å distance restraints. Their displacement parameters were freely refined.

Data collection: *SMART* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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