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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.003 Å R factor = 0.048 wR factor = 0.123 Data-to-parameter ratio = 9.8

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

Bis(imidazol-1-yl)methane

In the title compound, $C_7H_8N_4$, the C atom of the bridging methylene group lies on a twofold rotation axis. The dihedral angle between the symmetry-related imidazole rings is 75.0 (2)°. The crystal packing is stabilized by van der Waals forces. Received 17 March 2006 Accepted 29 March 2006

Comment

Polyazolylmethanes have attracted considerable attention in the fields of biochemistry (Sorrell & Borovik, 1987), materials chemistry (Muller *et al.*, 2001), heterocyclic chemistry (Juliá *et al.*, 1982) and coordination chemistry (Effendy *et al.*, 2003). Although many structures of their metallic complexes have been reported, there are only a few reports on the structures of the ligands.



The title compound, (I), was prepared by using the phasetransfer organic synthesis method (Juliá *et al.*, 1984). The molecule contains two five-membered imidazole rings bridged by a methylene group (Fig. 1). Atom C4 of the bridging methylene group lies on a twofold rotation axis. The back donation of the lone pair of electrons of atom N2 may affect the C–N bond lengths and C–N–C bond angles (Table 1). The dihedral angle between the symmetry-related imidazole rings is 75.0 (2)°. The crystal packing is stabilized by van der Waals forces.

Experimental

The title compound was synthesized by modifying a literature method (Diez-Barra *et al.*, 1993) and was identified by NMR. It was crystallized by slow evaporation of a chloroform–methanol (5:1 v/v) solution (yield 200 mg, 78%; m.p. 441–442 K).

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Crystal data

 $\begin{array}{l} C_7H_8N_4 \\ M_r = 148.17 \\ Orthorhombic, \ P2_12_12 \\ a = 8.0357 \ (18) \ {\rm \AA} \\ b = 10.545 \ (2) \ {\rm \AA} \\ c = 4.3613 \ (10) \ {\rm \AA} \\ V = 369.55 \ (15) \ {\rm \AA}^3 \end{array}$

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\rm min} = 0.757, T_{\rm max} = 0.998$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.048$ $wR(F^2) = 0.123$ S = 1.02508 reflections 52 parameters H-atom parameters constrained

Table 1	
Selected	geometric parameters (Å

Selected	geometric	parameters	(A, `	') .

N1-C1	1.295 (3)	N2-C3	1.375 (2)
N1-C2	1.367 (2)	N2-C4	1.444 (2)
N2-C1	1.351 (2)	C2-C3	1.342 (2)
C1-N1-C2	104.94 (16)	N1-C1-N2	112.50 (18)
C1-N2-C3	106.04 (16)	C3-C2-N1	110.81 (19)
C1-N2-C4	126.83 (14)	C2-C3-N2	105.71 (14)
C3-N2-C4	127.13 (13)	$N2^{i}-C4-N2$	112.1 (2)
C2-N1-C1-N2	-0.1(2)	N1-C2-C3-N2	0.3 (2)
C3-N2-C1-N1	0.3 (2)	C1-N2-C3-C2	-0.3(2)
C1-N1-C2-C3	-0.1 (2)		

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

H atoms were positioned geometrically at distances of 0.93 (CH) and 0.97 Å (CH₂) from the parent C atoms and refined as riding with $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$. Due to the absence of any significant anomalous scatterers, Friedel pairs were merged before the final refinement.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics:



2176 measured reflections 508 independent reflections 461 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.079$ $\theta_{\text{max}} = 27.0^{\circ}$

$$\begin{split} w &= 1/[\sigma^2(F_o^2) + (0.0898P)^2 \\ &+ 0.003P] \\ \text{where } P &= (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\text{max}} &= 0.007 \\ \Delta\rho_{\text{max}} &= 0.19 \text{ e } \text{\AA}^{-3} \\ \Delta\rho_{\text{min}} &= -0.17 \text{ e } \text{\AA}^{-3} \\ \text{Extinction correction: SHELXL97} \\ \text{Extinction coefficient: 0.11 (4)} \end{split}$$



Figure 1

View of (I), with the atom-numbering scheme and displacement ellipsoids drawn at the 50% probability level [symmetry code: (A) -x, 1 - y, z].

SHELXTL (Bruker, 1998); software used to prepare material for publication: SHELXTL.

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