

Chuan-Ming Jin* and
Chu-Ru GongDepartment of Chemistry and Environmental
Engineering, Hubei Normal University,
Huangshi 435002, People's Republic of ChinaCorrespondence e-mail:
jincm1999@yahoo.com

Key indicators

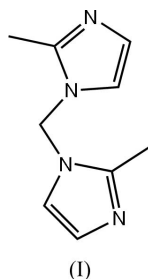
Single-crystal X-ray study
 $T = 298$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.068
 wR factor = 0.169
Data-to-parameter ratio = 13.5For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

Bis(2-methylimidazol-1-yl)methane

In the title compound, $\text{C}_9\text{H}_{12}\text{N}_4$, the asymmetric unit contains one half-molecule. The C atom of the bridging methylene group lies on a twofold rotation axis. The dihedral angle between the symmetry-related imidazole rings is $88.1(2)^\circ$.

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 metal complexes have been reported, there are only a few reports on the structures of the ligands themselves.



The title compound, (I) (Fig. 1), was prepared by using the phase-transfer organic synthesis method (Juliá *et al.*, 1984). The asymmetric unit consists of one half-molecule. The C atom of the bridging methylene group lies on a twofold rotation axis.

The back donation of the lone pair of electrons of atom N1 may affect the C—N bond lengths and C—N—C bond angles. The dihedral angle between the symmetry-related imidazole rings is $88.1(2)^\circ$.

Experimental

The title molecule was synthesized by modifying the literature method of Diez-Barra *et al.* (1993) and was identified by ^1H NMR [300 MHz, CDCl_3 : δ 6.87 (2H, s), 6.77 (2H, s), 5.77 (2H, s), 2.37 (6H, s)] and ^{13}C NMR (δ 144.8, 128.9, 119.1, 55.7, 13.7). It was crystallized from a chloroform–methanol (5:1) mixture by slow evaporation (yield: 50 mg, 72%, m.p. 447–449 K).

Crystal data

$\text{C}_9\text{H}_{12}\text{N}_4$
 $M_r = 176.23$
Monoclinic, $C2/c$
 $a = 12.371(2)$ Å
 $b = 8.6873(16)$ Å
 $c = 9.9175(18)$ Å
 $\beta = 120.599(3)^\circ$
 $V = 917.4(3)$ Å³
 $Z = 4$

$D_x = 1.276$ Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 958
reflections
 $\theta = 6.1$ – 50.4°
 $\mu = 0.08$ mm⁻¹
 $T = 298(2)$ K
Block, colorless
 $0.52 \times 0.24 \times 0.22$ mm

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Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.976$, $T_{\max} = 0.982$
 2281 measured reflections

826 independent reflections
 668 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.018$
 $\theta_{\text{max}} = 25.3^\circ$
 $h = -12 \rightarrow 14$
 $k = -10 \rightarrow 8$
 $l = -11 \rightarrow 11$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.068$
 $wR(F^2) = 0.169$
 $S = 1.08$
 826 reflections
 61 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0801P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.32 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.31 \text{ e } \text{\AA}^{-3}$
 Extinction correction: SHELXL97
 Extinction coefficient: 0.022 (6)

Table 1

Selected geometric parameters (\AA , $^\circ$).

N1—C3	1.362 (2)	N2—C3	1.309 (3)
N1—C1	1.374 (3)	N2—C2	1.368 (3)
N1—C4	1.445 (2)	C3—C5	1.483 (3)
C1—C2	1.341 (3)		
C3—N1—C1	107.29 (15)	C1—C2—N2	111.0 (2)
C3—N1—C4	127.18 (14)	N2—C3—N1	110.62 (16)
C1—N1—C4	125.46 (14)	N2—C3—C5	125.28 (17)
C2—C1—N1	105.35 (17)	N1—C3—C5	124.09 (18)
C3—N2—C2	105.69 (15)	N1—C4—N1 ⁱ	112.4 (2)
C3—N1—C1—C2	−0.3 (2)	C2—N2—C3—N1	−0.1 (2)
N1—C1—C2—N2	0.2 (2)	C1—N1—C3—N2	0.3 (2)
C3—N2—C2—C1	−0.1 (2)		

Symmetry code: (i) $1 - x, y, \frac{1}{2} - z$.

H atoms were positioned geometrically at distances of 0.93 (CH), 0.97 (CH₂) and 0.96 \AA (CH₃) from the parent C atoms; a riding model was used during the refinement process. The $U_{\text{iso}}(\text{H})$ values were constrained to be 1.2 (1.5 for methyl) times U_{eq} of the carrier atom.

Data collection: SMART (Bruker, 1998); cell refinement: SMART; data reduction: SAINT (Bruker, 1998); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1998); software used to prepare material for publication: SHELXTL.

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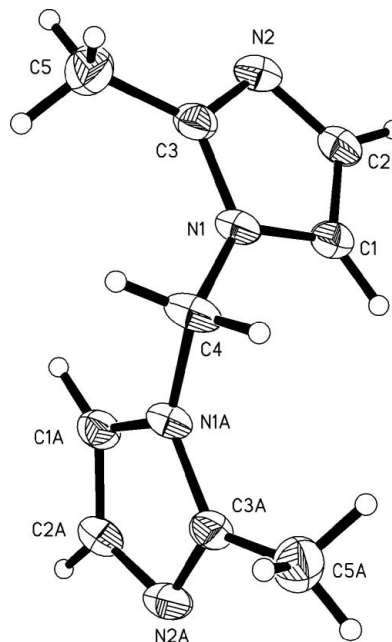


Figure 1

Drawing of the title molecule, showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. [Symmetry code (A): $1 - x, y, \frac{1}{2} - z$.]

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