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Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(C-C) = 0.007 \text{ Å}$ R factor = 0.045 wR factor = 0.088 Data-to-parameter ratio = 15.8

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

4'-Bromomethylbiphenyl-2-carbonitrile

In the title compound, $C_{14}H_{10}BrN$, the dihedral angle between the two aromatic rings is 48.34 (14)°. Weak $C-H\cdots Br$ and $C-H\cdots N$ interactions are the principal intermolecular forces.

Comment

The title compound, (I), is well known as a key intermediate in the production of Losartan (Duncia *et al.*, 1991), which is very useful in the treatment of hypertension by inhibiting angiotensin II (Campbell *et al.*, 1995). We are interested in how the molecular conformation of (I) (Fig. 1) affects the reactivity of the Br atom and its influence on the conversion of the cyano group to a tetrazole.



In (I), all the bond lengths and angles are normal. The two aromatic rings are non-coplanar; the dihedral angle is $48.34 (14)^{\circ}$. Weak C-H···Br and C-H···N interactions are the principal intermolecular forces, mediating the formation of the three-dimensional framework (Fig. 2 and Table 1).

Experimental

A solution of bromine (16 g, 0.1 mol) in 1,2-dichloroethane was added dropwise to a mixture of 4'-methyl-2-cyanobiphenyl (19.3 g, 0.1 mol) and azodiisobutyronitrile (0.8 g, 0.005 mol) in 1,2-dichloroethane (150 ml) at 343 (3) K with stirring for 2 h. The reaction mixture was then heated continuously at the same temperature, with monitoring by thin-layer chromatography, until the raw materials disappeared. The mixture was then treated with water (80 ml) three



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The molecular structure of (I), shown with 50% probability displacement ellipsoids.

Received 4 November 2003 Accepted 5 November 2003 Online 15 November 2003 times, dried and concentrated to dryness *in vacuo* to obtain crude (I). Crystallization from 1,2-dichloroethane–hexane yielded the pure solid (25 g, 92%) with a purity of 99.3% (HPLC). Colorless single crystals of (I) suitable for diffraction analysis were obtained from a tetrahydrofuran solution after two weeks at 293 (3) K.

Mo $K\alpha$ radiation

reflections

 $\mu = 3.47 \text{ mm}^{-1}$

T = 293 (2) K

Prism, colorless $0.3 \times 0.2 \times 0.2$ mm

 $\theta = 2.4 - 20.6^{\circ}$

Cell parameters from 1496

 $w = 1/[\sigma^2(F_o^2) + (0.02P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$

Absolute structure: Flack (1983),

 $(\Delta/\sigma)_{\rm max} < 0.001$

 $\Delta \rho_{\rm max} = 0.44 \ {\rm e} \ {\rm \AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.34 \ {\rm e} \ {\rm \AA}^{-3}$

965 Friedel pairs Flack parameter = 0.014 (15)

Crystal data

 $\begin{array}{l} C_{14}H_{10}{\rm BrN} \\ M_r = 272.14 \\ {\rm Orthorhombic}, Fdd2 \\ a = 47.219 \; (12) \; {\rm \AA} \\ b = 24.229 \; (6) \; {\rm \AA} \\ c = 4.1085 \; (10) \; {\rm \AA} \\ V = 4700 \; (2) \; {\rm \AA}^3 \\ Z = 16 \\ D_s = 1.538 \; {\rm Mg \; m^{-3}} \end{array}$

Data collection

Bruker SMART CCD area-detector	2284 independent reflections
diffractometer	1610 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.075$
Absorption correction: multi-scan	$\theta_{\rm max} = 26.0^{\circ}$
(SADABS; Bruker, 2000)	$h = -58 \rightarrow 52$
$T_{\min} = 0.43, T_{\max} = 0.50$	$k = -29 \rightarrow 24$
6379 measured reflections	$l = -5 \rightarrow 4$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.045$ $wR(F^2) = 0.088$ S = 1.052284 reflections 145 parameters H-atom parameters constrained

ers meters constrained

Table 1

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C1-H1B\cdots Br^{i}$	0.93	2.82	3.699 (6)	158
$C1-H1A\cdots N1^{ii}$	1.08	2.61	3.634 (8)	159
C9−H9···Br ⁱⁱⁱ	0.96	3.05	3.750 (6)	131

Symmetry codes: (i) x, y, z - 1; (ii) $\frac{3}{4} - x, \frac{1}{4} + y, \frac{3}{4} + z$; (iii) $\frac{1}{2} - x, 1 - y, \frac{1}{2} + z$.



Figure 2

A packing diagram of (I), viewed down the *c* axis. Intermolecular $C-H\cdots N$ and $C-H\cdots Br$ interactions are shown as dashed lines.

All the H atoms were located in a difference-density maps and refined isotropically as riding. The C-H bond lengths are in the range 0.90–1.08 Å and $U_{\rm iso} = 1.2U_{\rm eq}(\rm C)$.

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

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