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#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.005 Å R factor = 0.077 wR factor = 0.233 Data-to-parameter ratio = 16.4

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

# 6-(4-Chlorobenzyl)-3-phenyl-1,2,4,5-tetrazine

In the title compound,  $C_{15}H_{11}ClN_4$ , the tetrazine ring is twisted with respect to the chlorobenzene plane, with a dihedral angle of 69.4 (3)°. Weak  $C-H\cdots N$  hydrogen bonding between neighboring molecules helps to stabilize the crystal structure. Received 18 September 2006 Accepted 19 September 2006

## Comment

1,2,4,5-Tetrazine derivatives possess a wide spectrum of antiviral and antitumor properties and have been widely used in pesticides and herbicides (Sauer, 1996). In a continuation of our work on the structure–activity relationships of 1,2,4,5tetrazine derivatives (Hu *et al.*, 2004; 2005), we have prepared the title compound, (I), and determined its crystal structure.



The molecule structure of (I) is illustrated in Fig. 1. The tetrazine ring is coplanar with the C14-phenyl ring [dihedral angle 2.46 (3)°], but twisted with respect to the C8-benzene ring [dihedral angle 69.4 (3)°]. Weak  $C-H\cdots N$  hydrogen bonding occurs between neighboring molecules (Table 1), which helps to stabilize the crystal structure.

### **Experimental**

Figure 1

With sulfur (1.0 g) as catalyst, 85% hydrazine hydrate (10 ml, 170 mmol) was added dropwise to an anhydrous ethanol solution (15 ml) of 4-chlorobenzyl cyanide (50 mmol) and benzonitrile (50 mmol) at 295 K. After refluxing for 3 h, the mixture was cooled to room temperature and the resulting solid product was filtered off. The solid product was then dissolved in diethyl ether (15 ml) and oxidized by sodium nitrite (14 mmol) and acetic acid (14 mmol) over



The molecular structure of (I), shown with 30% probability displacement

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ellipsoids (arbitrary spheres for H atoms).

a period of 2 h to afford the product as purple crystals which were purified by preparative thin-layer chromatography over silica gel PF254 (2 mm) (cyclohexane–dichloromethane, 1:1) to give red single crystals of (I). The solid product was dissolved in tetrahydrofuran– anhydrous ethanol (4:1  $\nu/\nu$ ) and the solution evaporated gradually at room temperature to afford single crystals of (I).

Crystal data

 $\begin{array}{l} C_{15}H_{11}{\rm CIN_4} \\ M_r = 282.73 \\ {\rm Monoclinic, $P_{21}/c$} \\ a = 6.246 (2) {\rm \AA} \\ b = 16.671 (6) {\rm \AA} \\ c = 14.173 (4) {\rm \AA} \\ \beta = 112.661 (12)^{\circ} \\ V = 1361.9 (8) {\rm \AA}^3 \end{array}$ 

Data collection

Bruker SMART CCD area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: none 6707 measured reflections

#### Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.077$   $wR(F^2) = 0.233$  S = 0.962976 reflections 182 parameters H-atom parameters constrained Z = 4  $D_x = 1.379 \text{ Mg m}^{-3}$ Mo K $\alpha$  radiation  $\mu = 0.28 \text{ mm}^{-1}$ T = 293 (2) K Prism, red  $0.25 \times 0.20 \times 0.20 \text{ mm}$ 

2976 independent reflections 1595 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.061$  $\theta_{\text{max}} = 27.2^{\circ}$ 

$$\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.1387P)^2] \\ &where \ P = (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{max} < 0.001 \\ \Delta\rho_{max} = 0.25 \ e \ \text{\AA}^{-3} \\ \Delta\rho_{min} = -0.40 \ e \ \text{\AA}^{-3} \\ &\text{Extinction correction: SHELXL97} \\ &\text{Extinction coefficient: 0.058 (9)} \end{split}$$

## Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C13-H13\cdots N5^{i}$	0.93	2.55	3.295 (4)	137
Symmetry code: (i) x	- 1. v. z.			

H stoms were placed in colculated position

H atoms were placed in calculated positions, with C–H = 0.93 (aromatic) or 0.97 Å (methylene), and refined in riding mode, with  $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm C})$ .

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1997); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1999); software used to prepare material for publication: *SHELXL97*.

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