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# H. S. Yathirajan,<sup>a</sup> K. K. Vijaya Raj,<sup>b</sup> B. Narayana,<sup>b</sup> B. K. Sarojini<sup>c</sup> and Michael Bolte<sup>d</sup>\*

<sup>a</sup>Department of Studies in Chemistry, University of Mysore, Manasagangotri, Mysore 570 006, India, <sup>b</sup>Department of Chemistry, Mangalore University, Mangalagangotri 574 199, India, <sup>c</sup>Department of Chemistry, P. A. College of Engineering, Nadupadavu, Mangalore 574 153, India, and <sup>d</sup>Institut für Anorganische Chemie, J. W. Goethe-Universität Frankfurt, Max-von-Laue-Strasse 7, 60438 Frankfurt/Main, Germany

Correspondence e-mail: bolte@chemie.uni-frankfurt.de

#### **Key indicators**

Single-crystal X-ray study T = 173 K Mean  $\sigma$ (C–C) = 0.004 Å R factor = 0.037 wR factor = 0.087 Data-to-parameter ratio = 13.7

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

# 2-(2-Bromo-5-methoxyphenyl)-6-phenyl-1,3-thiazolo[3,2-*b*][1,2,4]triazole

The molecule of the title compound,  $C_{17}H_{12}BrN_3OS$ , is essentially planar. Geometric parameters are in the normal ranges. There are two intramolecular  $C-H\cdots N$  hydrogen bonds. Received 29 August 2006 Accepted 4 September 2006

# Comment

The reaction of  $\alpha$ -haloketones with 2-substituted 5-mercapto-1,2,4-triazoles may result in the formation of either the 2,5disubstituted-thiazolo[3,2-*b*]-*s*-triazole or the 3,5-disubstituted thiazolo[3,2-*b*]-1,2,4-triazole or both (Berk *et al.*, 2001; Potts & Husain, 1971). The synthesis of thiazolo[3,2-*b*]-1,2,4-triazoles and the isomeric thiazolo[2,3-*c*]-1,2,4-triazoles, and their diuretic, antibacterial and antifungal activities, have been studied by Jag Mohan & Kiran (1988). In the present study, 5-(2-bromo-5-methoxyphenyl)-4*H*-1,2,4-triazole-3-thiol, (1), was refluxed with phenacyl bromide, (2), in ethanol to obtain 2-(2-bromo-5-methoxyphenyl)-6-aryl-1,3-thiazolo[3,2-*b*]-[1,2,4]triazole, (3) (see scheme). We present here the crystal structure of (3).



A perspective view of compound (3) is shown in Fig. 1. Bond lengths and angles can be regarded as normal (Cambridge Structural Database, Version 5.27, November 2005, updated August 2006; MOGUL, Version 1.1; Allen, 2002). The molecule is essentially planar (r.m.s. deviation for all non-H atoms is 0.059 Å).

The molecular conformation of (3) is stabilized by two intramolecular  $C-H\cdots N$  hydrogen bonds (Table 1).

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# **Experimental**

For the synthesis of compound (3), 5-(2-bromo-5-methoxyphenyl)-4*H*-1,2,4-triazole-3-thiol (2.86 g, 0.01 mol) and the appropriate 2bromo-1-phenylethanone (2 g, 0.01 mol) were refluxed in ethanol for 4 h. The progress of the reaction was monitored by thin-layer chromatography. After completion of the reaction, the reaction mixture was cooled and the precipitated solid was filtered off. The solid obtained was recrystallized from a methanol–acetone solvent mixture (1:1). The compound was obtained as creamish crystals in 48% yield (m.p. 483 K). Analysis for  $C_{17}H_{12}BrN_3OS$ , found (calculated): C 52.80 (52.86), H 3.07 (3.13), N 10.72 (10.88)%. Spectroscopic data: IR (KBr,  $\nu$ , cm-1): 3118 and 3070 (–CH), 1475 (–C=N–), 734 (C–Br).

## Crystal data

 $C_{17}H_{12}BrN_3OS$   $M_r = 386.27$ Orthorhombic, *Pbca*  a = 14.9271 (13) Å b = 10.9247 (13) Å c = 18.8582 (16) Å V = 3075.3 (5) Å<sup>3</sup>

#### Z = 8 $D_x$ = 1.669 Mg m<sup>-3</sup> Mo K $\alpha$ radiation $\mu$ = 2.82 mm<sup>-1</sup> T = 173 (2) K Plate, colourless 0.32 × 0.16 × 0.08 mm

10330 measured reflections

 $R_{\rm int}=0.064$ 

 $\theta_{\rm max} = 25.6^\circ$ 

2883 independent reflections

2189 reflections with  $I > 2\sigma(I)$ 

#### Data collection

Stoe IPDS II two-circle diffractometer  $\omega$  scans Absorption correction: multi-scan (*MULABS*; Spek, 2003; Blessing, 1995)  $T_{\min} = 0.466, T_{\max} = 0.806$ 

# Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.037$   $wR(F^2) = 0.087$  S = 0.982883 reflections 210 parameters H-atom parameters constrained  $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0505P)^{2}]$ where  $P = (F_{o}^{2} + 2F_{c}^{2})/3$  $(\Delta/\sigma)_{max} = 0.001$  $\Delta\rho_{max} = 0.55 \text{ e } \text{Å}^{-3}$  $\Delta\rho_{min} = -0.80 \text{ e } \text{Å}^{-3}$ Extinction correction: *SHELXL97* (Sheldrick, 1997) Extinction coefficient: 0.0046 (4)

#### Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - \mathbf{H} \cdots A$
C26—H26···N8	0.95	2.40	2.788 (4)	104
C12—H12···N6	0.95	2.35	3.040 (4)	129



## Figure 1

The molecular structure of compound (3), with the atom numbering; displacement ellipsoids are drawn at the 50% probability level.

H atoms were found in a difference map but they were subsequently refined using a riding model, with C-H = 0.95 Å and  $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm C})$  or C-H = 0.98 Å and  $U_{\rm iso}({\rm H}) = 1.5 U_{\rm eq}({\rm C}_{\rm methyl})$ . The methyl group was allowed to rotate but not to tip.

Data collection: X-AREA (Stoe & Cie, 2001); cell refinement: X-AREA; data reduction: X-AREA; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXL97.

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