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2-Bromo-4-phenyl-1,3-thiazole

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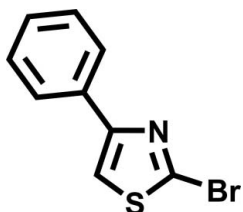
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Key indicators: single-crystal X-ray study; $T = 120$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.029; wR factor = 0.068; data-to-parameter ratio = 25.5.

In the title molecule, $\text{C}_9\text{H}_6\text{BrNS}$, the planes of the 2-bromo-1,3-thiazole and phenyl rings are inclined at 7.45 (10)° with respect to each other. In the crystal, molecules related by a centre of symmetry are held together *via* π - π interactions, with a short distance of 3.815 (2) Å between the centroids of the five- and six-membered rings. The crystal packing exhibits short intermolecular $\text{S} \cdots \text{Br}$ contacts of 3.5402 (6) Å.

Related literature

For syntheses and properties of compounds containing a thiazole fragment, see: Kelly & Lang (1995); Nicolaou *et al.* (1999); Cosford *et al.* (2003); Fyfe *et al.* (2004); Hamill *et al.* (2005). For the crystal structures of related compounds, see: Abbenante *et al.* (1996); Zhao *et al.* (2011); Ghabbour, Chia *et al.* (2012); Ghabbour, Kadi *et al.* (2012).



Experimental

Crystal data

 $\text{C}_9\text{H}_6\text{BrNS}$ $M_r = 240.12$

Monoclinic, $P2_1/n$
 $a = 5.8934$ (3) Å
 $b = 10.6591$ (6) Å
 $c = 13.8697$ (7) Å
 $\beta = 90.812$ (1)°
 $V = 871.18$ (8) Å³

$Z = 4$
Mo $K\alpha$ radiation
 $\mu = 4.89$ mm⁻¹
 $T = 120$ K
 $0.15 \times 0.12 \times 0.12$ mm

Data collection

Bruker APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2003)
 $T_{\min} = 0.527$, $T_{\max} = 0.591$

12144 measured reflections
2780 independent reflections
2258 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.045$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.029$
 $wR(F^2) = 0.068$
 $S = 1.03$
2780 reflections

109 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.40$ e Å⁻³
 $\Delta\rho_{\min} = -0.51$ e Å⁻³

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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Supporting information for this paper is available from the IUCr electronic archives (Reference: CV5440).

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2-Bromo-4-phenyl-1,3-thiazole

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S1. Comment

1,3-Thiazole rings appear in many compounds that exhibit important biological and pharmacological activities. For example, these rings feature in all the potent epothilones (Nicolaou *et al.*, 1999) used against multidrug-resistant tumor cell lines. They are also found among pharmaceuticals used for the treatment of type 2 diabetes (Fyfe *et al.*, 2004), antibiotic-like compounds (Kelly *et al.*, 1995), and metabotropic glutamate receptor subtype (mGluR5) antagonists (Cosford *et al.*, 2003; Hamill *et al.*, 2005). Herewith we present the title compound (**I**) prepared by the reaction of 2-amino-4-phenylthiazole with *n*-butyl nitrite and CuBr (Figure 1).

In **I** (Fig. 2), the bond lengths and angles are in a good agreement with those found in the related compounds (Abbenante *et al.*, 1996; Zhao *et al.*, 2011; Ghabbour, Chia *et al.*, 2012; Ghabbour, Kadi *et al.*, 2012). The 2-bromo-1,3-thiazole mean plane and phenyl ring are twisted by 7.45 (10)°.

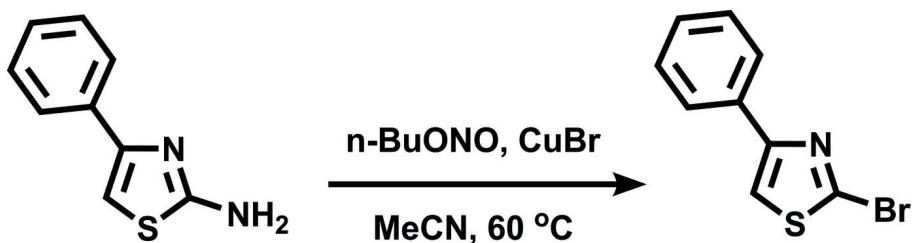
In the crystal, the molecules related by center of symmetry held together *via* $\pi\cdots\pi$ interactions proved by short $Cg^5\cdots Cg^{6i}$ distance of 3.815 (2) Å between the centroids of five-membered (Cg^5) and six-membered (Cg^6) rings [symmetry code: (i) $-x, 1-y, 1-z$]. The crystal packing exhibits short intermolecular $S\cdots Br^{ii}$ contacts of 3.5402 (6) Å (Figure 3) [symmetry code: (ii) $-1+x, y, z$].

S2. Experimental

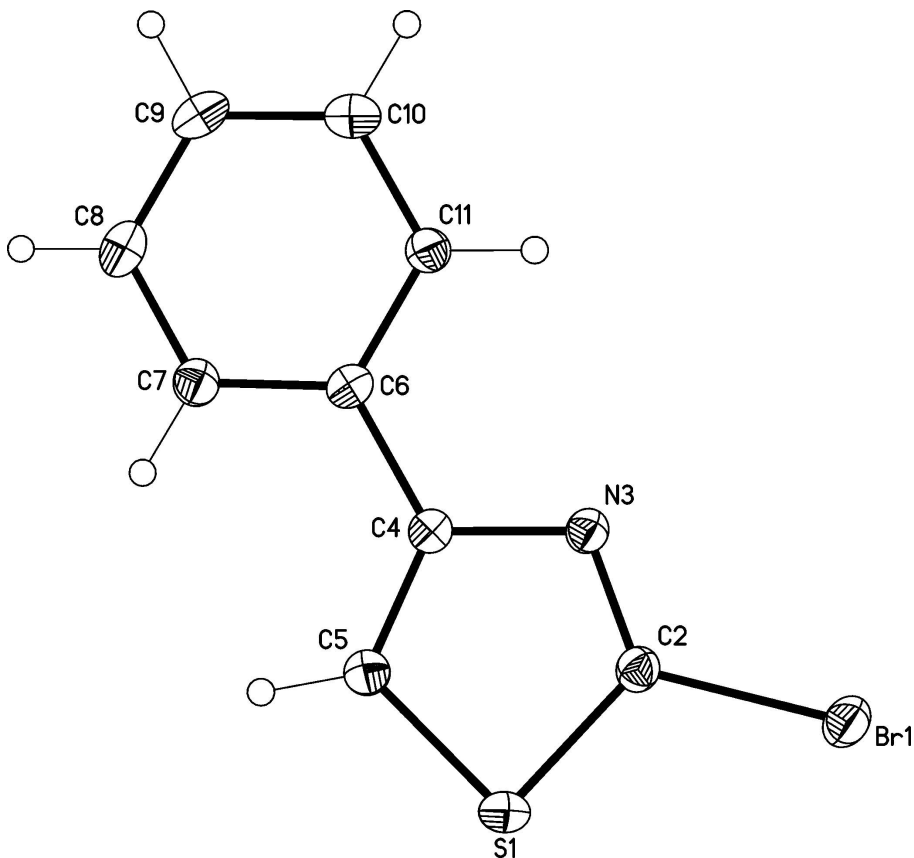
The 4-phenyl-2-aminothiazole (8.1 g, 46.9 mmol) and CuBr (10.7 g, 74.6 mmol) were dissolved in acetonitrile at room temperature. *n*-Butyl nitrite (8.7 ml, 7.69 g, 74.6 mmol) was added with stirring, and the solution was heated to 333 K. The reaction completed after 15 min. The reaction mixture was then evaporated to dryness in *vacuo*. The residue was dissolved in ethyl acetate (50 ml) and washed with ammonia solution (0.1 M, 2 × 50). The organic layer was dried over MgSO₄ and evaporated to dryness in *vacuo*. The residue was purified by chromatography on silica gel (heptane-ethyl acetate; 70:3, *v/v*). The residue crystallized from 5% solution in heptane. Yield is 53%. The single-crystal of the product **I** was obtained by slow crystallization from hexane. *M.p.* = 327–328 K. IR (KBr), ν/cm^{-1} : 3098, 3063, 1476, 1420, 1263, 1070, 1010, 836, 730, 689. ¹H NMR (500 MHz, DMSO-*d*₆, 304 K): 7.40–6.37 (m, 1H, Ph), 7.46 (t, 2H, *J* = 7.63, Ph), 7.92 (d, 2H, *J* = 7.32, Ph), 8.16 (s, 1H, thiazole). Anal. Calcd for C₉H₆BrNS: C, 45.02; H, 2.52. Found: C, 45.09; H, 2.57.

S3. Refinement

All hydrogen atoms were placed in the calculated positions [C—H = 0.95 Å] and refined in the riding model, with $U_{iso}(H) = 1.2U_{eq}(C)$.

**Figure 1**

Synthesis of 2-bromo-4-phenylthiazole.

**Figure 2**

Molecular structure of **I**. Displacement ellipsoids are presented at the 50% probability level. H atoms are depicted as small spheres of arbitrary radius.

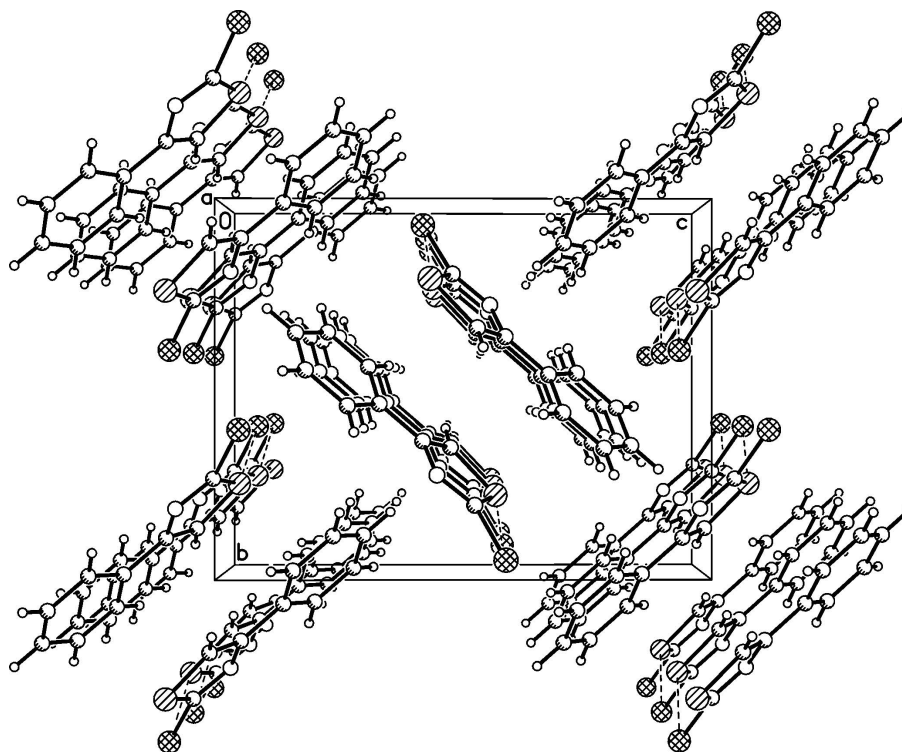


Figure 3

The crystal packing of **I** viewed along the *a* axis. Dashed lines indicate the intermolecular secondary S...Br interactions.

2-Bromo-4-phenyl-1,3-thiazole

Crystal data

C_9H_6BrNS

$M_r = 240.12$

Monoclinic, $P2_1/n$

Hall symbol: $-P\ 2_1n$

$a = 5.8934$ (3) Å

$b = 10.6591$ (6) Å

$c = 13.8697$ (7) Å

$\beta = 90.812$ (1)°

$V = 871.18$ (8) Å³

$Z = 4$

$F(000) = 472$

$D_x = 1.831$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 3185 reflections

$\theta = 2.4\text{--}29.5^\circ$

$\mu = 4.89$ mm⁻¹

$T = 120$ K

Prism, yellow

$0.15 \times 0.12 \times 0.12$ mm

Data collection

Bruker APEXII CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2003)

$T_{\min} = 0.527$, $T_{\max} = 0.591$

12144 measured reflections

2780 independent reflections

2258 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.045$

$\theta_{\max} = 31.0^\circ$, $\theta_{\min} = 2.4^\circ$

$h = -8 \rightarrow 8$

$k = -15 \rightarrow 15$

$l = -20 \rightarrow 19$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.029$	H-atom parameters constrained
$wR(F^2) = 0.068$	$w = 1/[\sigma^2(F_o^2) + (0.0323P)^2 + 0.1245P]$
$S = 1.03$	where $P = (F_o^2 + 2F_c^2)/3$
2780 reflections	$(\Delta/\sigma)_{\max} = 0.002$
109 parameters	$\Delta\rho_{\max} = 0.40 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta\rho_{\min} = -0.51 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.21824 (3)	0.90020 (2)	0.576233 (15)	0.02238 (7)
S1	-0.24024 (8)	0.75866 (5)	0.55984 (4)	0.01975 (11)
C2	0.0322 (3)	0.77659 (19)	0.52049 (14)	0.0165 (4)
N3	0.0960 (3)	0.70330 (16)	0.45150 (11)	0.0161 (3)
C4	-0.0825 (3)	0.62475 (17)	0.42443 (14)	0.0144 (4)
C5	-0.2768 (3)	0.64278 (19)	0.47549 (14)	0.0178 (4)
H5	-0.4135	0.5972	0.4655	0.021*
C6	-0.0500 (3)	0.53485 (18)	0.34463 (14)	0.0151 (4)
C7	-0.2155 (3)	0.4453 (2)	0.32167 (14)	0.0184 (4)
H7	-0.3508	0.4419	0.3580	0.022*
C8	-0.1849 (4)	0.3613 (2)	0.24652 (14)	0.0214 (4)
H8	-0.2988	0.3009	0.2318	0.026*
C9	0.0134 (4)	0.3655 (2)	0.19254 (15)	0.0213 (4)
H9	0.0354	0.3080	0.1411	0.026*
C10	0.1778 (4)	0.4544 (2)	0.21484 (15)	0.0208 (4)
H10	0.3130	0.4576	0.1784	0.025*
C11	0.1471 (3)	0.53900 (19)	0.28978 (14)	0.0174 (4)
H11	0.2605	0.5999	0.3038	0.021*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.02065 (11)	0.02080 (11)	0.02561 (12)	-0.00021 (8)	-0.00253 (8)	-0.00725 (8)
S1	0.0181 (2)	0.0210 (3)	0.0202 (2)	0.00233 (19)	0.00358 (19)	-0.00361 (19)
C2	0.0156 (9)	0.0158 (9)	0.0181 (9)	0.0006 (7)	-0.0012 (7)	-0.0013 (7)

N3	0.0167 (8)	0.0160 (8)	0.0155 (8)	-0.0008 (6)	-0.0005 (6)	-0.0006 (6)
C4	0.0162 (9)	0.0129 (9)	0.0142 (8)	0.0009 (7)	-0.0008 (7)	0.0015 (7)
C5	0.0173 (9)	0.0173 (9)	0.0190 (10)	0.0002 (7)	0.0009 (7)	-0.0011 (8)
C6	0.0187 (9)	0.0137 (9)	0.0129 (8)	0.0020 (7)	-0.0013 (7)	0.0015 (7)
C7	0.0194 (9)	0.0182 (9)	0.0176 (9)	-0.0013 (8)	0.0002 (7)	0.0010 (8)
C8	0.0264 (10)	0.0180 (10)	0.0198 (10)	-0.0019 (8)	-0.0054 (8)	-0.0004 (8)
C9	0.0307 (11)	0.0180 (10)	0.0153 (9)	0.0045 (8)	-0.0019 (8)	-0.0026 (7)
C10	0.0224 (10)	0.0225 (10)	0.0175 (9)	0.0041 (8)	0.0028 (8)	0.0006 (8)
C11	0.0184 (9)	0.0165 (9)	0.0172 (9)	-0.0013 (7)	0.0019 (7)	-0.0004 (7)

Geometric parameters (Å, °)

Br1—C2	1.874 (2)	C7—C8	1.388 (3)
S1—C5	1.713 (2)	C7—H7	0.9500
S1—C2	1.714 (2)	C8—C9	1.398 (3)
C2—N3	1.295 (2)	C8—H8	0.9500
N3—C4	1.392 (2)	C9—C10	1.388 (3)
C4—C5	1.368 (3)	C9—H9	0.9500
C4—C6	1.478 (3)	C10—C11	1.390 (3)
C5—H5	0.9500	C10—H10	0.9500
C6—C11	1.398 (3)	C11—H11	0.9500
C6—C7	1.398 (3)		
C5—S1—C2	88.40 (10)	C8—C7—H7	119.5
N3—C2—S1	116.81 (15)	C6—C7—H7	119.5
N3—C2—Br1	123.68 (15)	C7—C8—C9	120.0 (2)
S1—C2—Br1	119.49 (11)	C7—C8—H8	120.0
C2—N3—C4	109.64 (17)	C9—C8—H8	120.0
C5—C4—N3	114.24 (17)	C10—C9—C8	119.28 (19)
C5—C4—C6	126.63 (18)	C10—C9—H9	120.4
N3—C4—C6	119.11 (17)	C8—C9—H9	120.4
C4—C5—S1	110.91 (15)	C9—C10—C11	120.80 (19)
C4—C5—H5	124.5	C9—C10—H10	119.6
S1—C5—H5	124.5	C11—C10—H10	119.6
C11—C6—C7	118.68 (18)	C10—C11—C6	120.30 (19)
C11—C6—C4	120.27 (17)	C10—C11—H11	119.9
C7—C6—C4	121.05 (17)	C6—C11—H11	119.9
C8—C7—C6	120.92 (19)		
C5—S1—C2—N3	-0.65 (17)	C5—C4—C6—C7	-8.5 (3)
C5—S1—C2—Br1	178.28 (13)	N3—C4—C6—C7	173.20 (18)
S1—C2—N3—C4	0.5 (2)	C11—C6—C7—C8	0.5 (3)
Br1—C2—N3—C4	-178.35 (13)	C4—C6—C7—C8	179.78 (19)
C2—N3—C4—C5	-0.1 (2)	C6—C7—C8—C9	0.0 (3)
C2—N3—C4—C6	178.41 (17)	C7—C8—C9—C10	-0.2 (3)
N3—C4—C5—S1	-0.4 (2)	C8—C9—C10—C11	-0.1 (3)
C6—C4—C5—S1	-178.74 (16)	C9—C10—C11—C6	0.6 (3)
C2—S1—C5—C4	0.56 (16)	C7—C6—C11—C10	-0.8 (3)

C5—C4—C6—C11	170.74 (19)	C4—C6—C11—C10	179.93 (18)
N3—C4—C6—C11	-7.5 (3)		
