

2-Amino-4-[4-(benzyloxy)phenyl]-5-methylthiazol-3-ium bromide monohydrate

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Key indicators

Single-crystal X-ray study
 $T = 294\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$
 R factor = 0.034
 wR factor = 0.086
Data-to-parameter ratio = 17.8

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

The crystal packing of the hydrated title salt, $\text{C}_{17}\text{H}_{17}\text{N}_2\text{OS}^+\cdot\text{Br}^-\cdot\text{H}_2\text{O}$, dominated by an extensive $\text{N}-\text{H}\cdots\text{O}_{\text{water}}$, $\text{N}-\text{H}\cdots\text{Br}$ and $\text{O}_{\text{water}}-\text{H}\cdots\text{Br}$ hydrogen-bonding network between organic molecules, bromine anions and water molecules. The anions are located on crystallographic twofold axes.

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Comment

2-Amino-4-arylthiazoles are an important class of heterocyclic compounds in the field of organic pharmaceutical chemistry (Marcantonio *et al.*, 2002). Heterocyclic compounds containing a thiazole ring generally exhibit broad-spectrum biological activity and some thiazole derivatives have been used as local anaesthetics, as anticonvulsant, antiviral, and antimicrobial agents, insecticides and so on (Narayana *et al.*, 2004). We report here the synthesis and structure of the title 2-amino-4-arylthiazole derivative, (I).

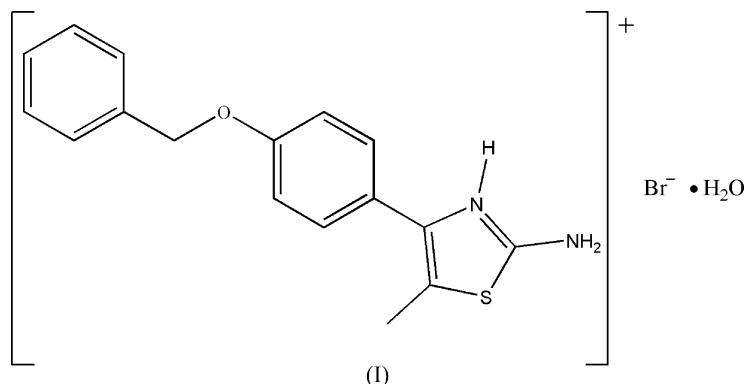


Fig. 1 shows the molecular structure of (I), with the atomic numbering scheme. The molecule of (I) contains a planar thiazole ring and two benzene rings, with dihedral angles of $42.7(2)^\circ$ and $10.6(2)^\circ$, respectively, between the least-squares planes of the two benzene rings (C5–C10 and C12–C17) and the thiazole ring. The dihedral angle between the two benzene rings is $51.3(2)^\circ$.

The anions lie on twofold axes. The cations, anions and water molecules in (I) associate *via* $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds between the amino H atoms and water O atoms, with the hydrogen-bonding array additionally involving $\text{O}-\text{H}\cdots\text{Br}$ and $\text{N}-\text{H}\cdots\text{Br}$ hydrogen bonds (Table 1). The crystal structure is further stabilized by van der Waals forces.

Experimental

1-[4-(Benzyloxy)phenyl]propan-1-one (0.005 mol) was dissolved in 100 ml ethanol and the mixture was stirred and heated to reflux.

Cupric bromide (0.01 mol) was added to the reaction mixture in batches and the course of the reaction was followed by thin-layer chromatography. After the reaction had finished, the mixture was filtered, 0.05 mol thiurea was added to the filtrate, and the mixture was heated at reflux for 3 h. The mixture was then cooled in a refrigerator, giving a yellow precipitate, which was filtered off and dried to obtain the title compound (I). MS (m/z): 296 (base, M^+), 205 ($M^+ - \text{PhCH}_2$), 91 (100%, PhCH_2^+), 80, 65, 51. Crystals suitable for X-ray crystallographic analysis were obtained by slow evaporation of an ethanol solution at room temperature.

Crystal data

$\text{C}_{17}\text{H}_{17}\text{N}_2\text{OS}^+ \cdot \text{Br}^- \cdot \text{H}_2\text{O}$	$Z = 8$
$M_r = 395.31$	$D_x = 1.483 \text{ Mg m}^{-3}$
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
$a = 10.0462 (7) \text{ \AA}$	$\mu = 2.45 \text{ mm}^{-1}$
$b = 16.2513 (11) \text{ \AA}$	$T = 294 (2) \text{ K}$
$c = 21.7802 (15) \text{ \AA}$	Block, yellow
$\beta = 95.1720 (10)^\circ$	$0.48 \times 0.45 \times 0.40 \text{ mm}$
$V = 3541.4 (4) \text{ \AA}^3$	

Data collection

Bruker AXS SMART 1000 CCD diffractometer	8967 measured reflections
ω scans	3841 independent reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	2494 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.328$, $T_{\max} = 0.375$	$R_{\text{int}} = 0.029$
	$\theta_{\max} = 27.0^\circ$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0329P)^2 + 2.5508P]$
$R[F^2 > 2\sigma(F^2)] = 0.035$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.086$	$(\Delta/\sigma)_{\max} = 0.001$
$S = 1.02$	$\Delta\rho_{\max} = 0.30 \text{ e \AA}^{-3}$
3841 reflections	$\Delta\rho_{\min} = -0.31 \text{ e \AA}^{-3}$
216 parameters	
H atoms treated by a mixture of independent and constrained refinement	

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1A}\cdots\text{O2W}$	0.86	1.88	2.724 (3)	166
$\text{N2}-\text{H2A}\cdots\text{Br2}$	0.86	2.68	3.483 (2)	155
$\text{N2}-\text{H2B}\cdots\text{Br1}^\dagger$	0.86	2.59	3.362 (2)	149
$\text{O2W}-\text{H2C}\cdots\text{Br1}$	0.78 (3)	2.52 (4)	3.290 (2)	171 (4)
$\text{O2W}-\text{H2D}\cdots\text{Br2}$	0.79 (3)	2.50 (4)	3.259 (2)	161 (4)

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

All H atoms except for the water H atoms were refined in the riding-model approximation, with N—H distances of 0.86 \AA and C—H distances of 0.93 (aromatic), 0.96 (methyl) and 0.97 \AA (methylene), and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$ or $1.5U_{\text{eq}}(\text{methyl C})$. The water H atoms were located in Fourier syntheses and $U_{\text{iso}}(\text{H})$ values were set equal to $1.2U_{\text{eq}}(\text{O})$.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT-Plus (Bruker, 2003); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics:

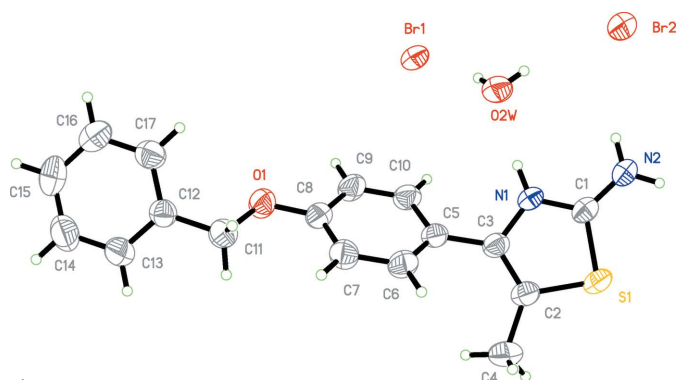


Figure 1

The molecular structure of (I), showing the atom-labelling scheme and 50% probability displacement ellipsoids (arbitrary spheres for H atoms).

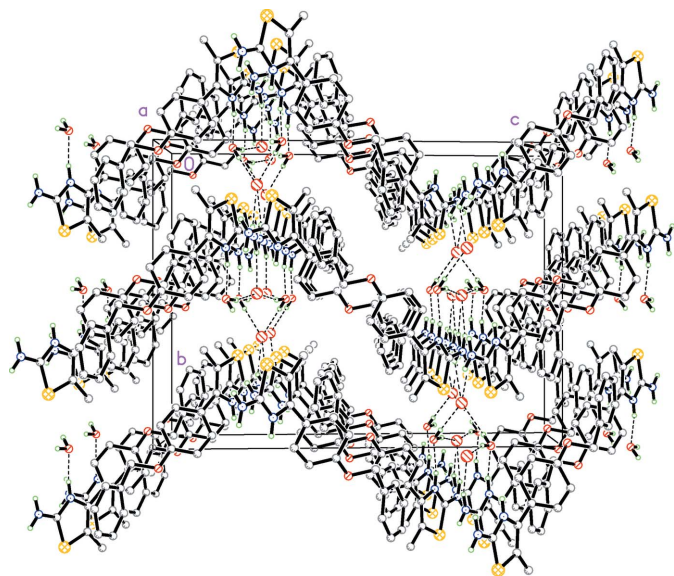


Figure 2

A packing diagram for (I). Dashed lines indicate hydrogen bonds.

SHELXTL (Bruker, 1997); software used to prepare material for publication: SHELXTL.

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