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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.003 Å R factor = 0.035 wR factor = 0.096 Data-to-parameter ratio = 14.5

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

1-Butyl-3-(4-chloro-1,3-benzothiazol-2-yl)urea monohydrate

In the crystal structure of the title compound, $C_{12}H_{14}ClN_3OS$ - H_2O , the butyl(chlorobenzo[*d*]thiazolyl)urea molecule, excluding butyl H atoms, displays a roughly planar structure. The water molecules link with the urea derivative molecules *via* $O-H\cdots O$ and $N-H\cdots O$ hydrogen bonding.

Comment

1-Butyl-3-(4-chlorobenzo[d]thiazol-2-yl)urea is useful as an inhibitor of serine/threonine and tyrosine kinases (Cusack *et al.*, 2003). Recently we obtained single crystals of its monohydrate, (I), and determined the crystal structure of (I).



The molecular structure of (I) is shown in Fig. 1. The butyl(chlorobenzo[d]thiazolyl)urea molecule, excluding butyl H atoms, displays a roughly planar structure. The C9–C10– C11–C12 torsion angle of -178.95 (19)° shows the extended conformation of the butyl group. The N3–C9 bond distance indicates a typical single bond, but the N2–C7 bond is much shorter than the N3–C9 bond (Table 1) and shows somewhat the character of a C=N double bond. This is presumably due to the overlap of the non-bonding orbital of the imino N2 atom with the π orbital of the thiazole ring.

The water molecules link with the urea derivative molecules *via* $O-H\cdots O$ and $N-H\cdots O$ hydrogen bonding (Table 2), which helps to stabilize the crystal structure.

Experimental

Butan-1-amine (9 mmol) was mixed with a dry tetrahydrofuran solution (12 ml) of phenyl 4-chlorobenzo[d]thiazol-2-ylcarbamate (3 mmol). The mixture was heated for 20 min at 423 K in a microwave synthetic reactor. The solvent was then removed under reduced pressure. Colourless prismatic single crystals of (I) were obtained from an acetone–water (10:1)solution.

Crystal data

$C_{12}H_{14}CIN_3OS \cdot H_2O$	V = 712.9 (3) Å ³
$M_r = 301.8$	Z = 2
Triclinic, P1	$D_x = 1.406 \text{ Mg m}^{-3}$
a = 7.0075 (16) Å	Mo $K\alpha$ radiation
$b = 7.6324 (18) \text{\AA}$	$\mu = 0.42 \text{ mm}^{-1}$
c = 14.826 (3) Å	T = 293 (2) K
$\alpha = 96.568 \ (4)^{\circ}$	Prism, colourless
$\beta = 91.451 \ (4)^{\circ}$	$0.30 \times 0.26 \times 0.20 \text{ mm}$
$\gamma = 114.775 \ (4)^{\circ}$	

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Data collection

Bruker SMART 1000 CCD areadetector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 2002) $T_{\min} = 0.885$, $T_{\max} = 0.921$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.035$ $wR(F^2) = 0.096$ S = 1.022507 reflections 173 parameters H-atom parameters constrained 3635 measured reflections 2507 independent reflections 1939 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.027$ $\theta_{\text{max}} = 25.0^{\circ}$

$$\begin{split} w &= 1/[\sigma^2(F_o^2) + (0.0458P)^2 \\ &+ 0.1733P] \\ \text{where } P &= (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\text{max}} &= 0.001 \\ \Delta\rho_{\text{max}} &= 0.34 \text{ e } \text{\AA}^{-3} \\ \Delta\rho_{\text{min}} &= -0.21 \text{ e } \text{\AA}^{-3} \end{split}$$

Table 1

Selected bond lengths (Å).

S1-C5	1.741 (2)	N2-C7	1.365 (3)
S1-C7	1.753 (2)	N2-C8	1.391 (3)
O1-C8	1.223 (2)	N3-C8	1.333 (3)
N1-C6	1.386 (2)	N3-C9	1.453 (3)
N1-C7	1.300 (3)		

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots \mathbf{A}$
$N2-H2\cdots O2^{i}$	0.86	2.05	2.826 (2)	149
$N3-H3\cdots O2^{i}$	0.86	2.21	2.962 (2)	147
$O2-H2B\cdots N1^{ii}$	0.85	2.13	2.939 (2)	158
$O2-H2C\cdots O1^{iii}$	0.85	2.19	2.918 (2)	144

Symmetry codes: (i) -x + 2, -y + 1, -z; (ii) x - 1, y - 1, z; (iii) -x + 1, -y + 1, -z.

Water H atoms were located in a difference Fourier map and refined as riding in their as-found relative positions, with $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm O})$. Methyl H atoms were placed in calculated positions with C-H = 0.96 Å and the torsion angle was refined to fit the electron density; $U_{\rm iso}({\rm H}) = 1.5U_{\rm eq}({\rm C})$. Other H atoms were placed in calculated positions with N-H = 0.86 Å, and C-H = 0.93 (aromatic) or 0.97 Å (methylene), and refined in riding mode, with $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$.



Figure 1

The molecular structure of (I), with 30% probability displacement ellipsoids (arbitrary spheres for H atoms).

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1999); software used to prepare material for publication: *SHELXTL*.

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