Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.006 Å R factor = 0.035 wR factor = 0.082 Data-to-parameter ratio = 14.8

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

3-(5-Bromo-2-hydroxybenzylammonio)propanoate

The title compound, $C_{10}H_{12}BrNO_3$, exists as a zwitterion in the solid state; the delocalized $-CO_2$ unit carries the negative charge and the secondary N atom carries the positive charge.

Received 18 October 2006 Accepted 31 October 2006

Comment

Schiff bases derived from 5-bromosalicylaldimine possess useful antifungal and anticancer properties (Liang *et al.*, 2006; Zhao & Liu, 2002). The C—N double bond in these systems is rigid, but can be reduced. The reduced compounds are of chemical interest as they are more flexible (Chen *et al.*, 2003; Hazell *et al.*, 1997); for example, N-(2-hydroxybenzyl)alanine functions as a chelate to Cu^{II} (Sureshan & Bhattacharya, 1998; Yang *et al.*, 2001, 2004).



The 4-bromo-substituted title compound, (I), exists as a zwitterion in the solid state (Fig. 1), with a delocalized carboxylate group at one end of the molecule. The secondary ammonium group interacts with the O atoms of other zwitterions, forming a linear chain structure. Interestingly, both H atoms are engaged in bifurcated hydrogen bonds (Table 1).

Experimental

5-Bromosalicylaldimine (0.20 g, 1 mmol), β -alanine (0.09 g, 1 mmol) and potassium hydroxide (0.06 g, 1 mmol) were dissolved in 80% aqueous methanol (20 ml). The mixture was kept cold while stirring, giving a clear yellow solution. An aqueous solution (5 ml) of sodium borohydride (0.50 mmol, 0.02 g) was added, followed by addition of 1 *M* hydrochloric acid until the the pH of the reaction mixture reached 5.5. The resulting white solid was collected and washed with ethanol. The compound was purified by recrystallization from 50% aqueous methanol (15 ml). Colourless prismatic crystals separated from the solution after 2 weeks in about 60% yield. Elemental analysis found: C 43.80, H 4.37, N 5.17%; calculated: C 43.78, H 4.38, N 5.15%.

Crystal data

 $C_{10}H_{12}BrNO_3$ $M_r = 274.12$ Orthorhombic, *Pna2*₁ a = 13.971 (4) Å b = 14.280 (4) Å c = 5.553 (2) Å V = 1107.9 (8) Å³

Z = 4 D_x = 1.643 Mg m⁻³ Mo K α radiation μ = 3.70 mm⁻¹ T = 293 (2) K Prism, colourless 0.30 × 0.20 × 0.20 mm

Acta Cryst. (2006). E62, o5409–o5410

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Data collection

Bruker APEX-II CCD diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\min} = 0.366, T_{\max} = 0.477$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.035$ $wR(F^2) = 0.082$ S = 0.922177 reflections 147 parameters H atoms treated by a mixture of independent and constrained refinement

Table 1Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
O1−H1o···O3 ⁱ	0.85 (1)	1.80 (2)	2.620 (4)	161 (5)
$N1 - H1n \cdot \cdot \cdot O2$	0.85(1)	2.22 (3)	2.836 (4)	129 (3)
$N1 - H1n \cdot \cdot \cdot O2^{ii}$	0.85(1)	2.38 (3)	2.929 (4)	122 (3)
$N1 - H2n \cdot \cdot \cdot O2^{iii}$	0.85 (1)	2.28 (3)	2.993 (5)	141 (4)
$N1 - H2n \cdot \cdot \cdot O3^{iii}$	0.85 (1)	2.36 (3)	3.106 (5)	147 (4)

5103 measured reflections

 $R_{\rm int}=0.042$

 $\theta_{\rm max} = 27.1^{\circ}$

 $(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta\rho_{\text{max}} = 0.54 \text{ e} \text{ Å}^{-3}$

 $\Delta \rho_{\rm min} = -0.40 \text{ e} \text{ Å}^{-3}$

825 Friedel pairs

Flack parameter: 0.04 (1)

2177 independent reflections

 $w = 1/[\sigma^2(F_o^2) + (0.0466P)^2]$

where $P = (F_0^2 + 2F_c^2)/3$

Absolute structure: Flack (1983),

1687 reflections with $I > 2\sigma(I)$

Symmetry codes: (i) $-x + \frac{3}{2}$, $y - \frac{1}{2}$, $z + \frac{1}{2}$; (ii) -x + 1, -y + 2, $z + \frac{1}{2}$; (iii) x, y, z + 1.

Carbon-bound H atoms were placed at calculated positions (C–H = 0.93–0.97 Å) and were included in the refinement in the ridingmodel approximation, with $U_{iso}(H) = 1.2U_{eq}(C)$. The ammonium and hydroxy H atoms were located in a difference Fourier map, and were refined with distance restraints of N–H = O–H = 0.85 (1) Å, and with isotropic displacement parameters.

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *SHELXL97*.



Figure 1

The molecular structure of (I), showing displacement ellipsoids at the 50% probability level and H atoms as spheres of arbitrary radii.

This work was supported by the Natural Science Foundation of the Guangxi Zhuang Autonomous Region (grant No. 0339034) and the Science Research Foundation of Guangxi Universities. We also thank Hechi University and the University of Malaya for supporting this study.

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