Acta Crystallographica Section E

Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 120 KMean $\sigma(C-C) = 0.005 \text{ Å}$ R factor = 0.061 wR factor = 0.126 Data-to-parameter ratio = 12.7

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

2-(Benzoylmethylsulfanyl)-6-methoxy-1*H*-benzimidazole

The structure of the title compound, C₁₆H₁₄N₂O₂S, has been determined and shows a twisted molecule with a dihedral angle of 66.31 (11) Å between the two aromatic ring systems. A single N-H···N association creates a slightly convoluted, linear hydrogen-bonded chain [graph set $C_2^2(4)$] in the b-axis direction.

Received 5 February 2004 Accepted 9 February 2004 Online 14 February 2004

Comment

A search of the April 2003 release of the Cambridge Structural Database (Allen, 2002) reveals that the 2-benzimidazoylthio moiety has been structurally characterized more often as a ligand in metal coordination and organometallic complexes than as a fragment of an organic compound; ratio 13:7, respectively, with four of the seven compounds being sulfinyls. As part of a series of synthetic and structural studies on S-substituted 2-benzimidazoylthio compounds we have characterized the structure of the title compound, (I), which shows a twisted molecule with a dihedral angle of 66.31 (11) Å between the two aromatic ring systems (Fig. 1). The assignment of N1 in the imidazoyl ring was confirmed by both the initial location of H1 in a difference syntheses and the notable differences between the N1–C2 and C2–N3 bond distances. A single N-H···N association creates a slightly convoluted, linear hydrogen-bonded chain [graph set $C_2^2(4)$] in the b-axis direction. Hydrogen-bonding geometry is given in Table 1.

Experimental

The title compound was obtained from Key Organics Ltd and was crystallized from ethanol.

Figure 1

The molecular configuration and atom-numbering scheme for (I). Displacement ellipsoids are drawn at the 50% probability level.

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Crystal data

$D_x = 1.432 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation
Cell parameters from 17502
reflections
$\theta = 2.9 - 27.5^{\circ}$
$\mu = 0.24 \text{ mm}^{-1}$
T = 120 (2) K
Plate, colourless
$0.26 \times 0.14 \times 0.02 \text{ mm}$

Data collection

Bruker–Nonius KappaCCD areadetector diffractometer φ and ω scans G areadesis and G scans G areadesis and G scans G and G scans G areadesis and G are areadesis and G are areadesis and G are areadesis and G are areadesis and G areadesis and G are areadesis and G areadesis and G areadesis and G are areadesis and G areadesis and G are areadesis and G areadesis and G are areadesis and G are areadesis and G are areadesis and G are areadesis and G areadesis and G are areadesis and G are areadesis and G areadesis and G areadesis

Refinement

 $\begin{array}{lll} \mbox{Refinement on } F^2 & w = 1/[\sigma^2(F_o^2) + (0.052P)^2 \\ R[F^2 > 2\sigma(F^2)] = 0.061 & + 0.5232P] \\ wR(F^2) = 0.126 & \mbox{where } P = (F_o^2 + 2F_c^2)/3 \\ S = 1.07 & (\Delta/\sigma)_{\rm max} < 0.001 \\ 2413 \ {\rm reflections} & \Delta\rho_{\rm max} = 0.28 \ {\rm e \ \mathring{A}}^{-3} \\ 190 \ {\rm parameters} & \Delta\rho_{\rm min} = -0.31 \ {\rm e \ \mathring{A}}^{-3} \end{array}$

Table 1 Hydrogen-bonding geometry (Å, °).

D $ H$ $\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	D $ H$ $\cdot \cdot \cdot A$
$N1-H1\cdots N3^{i}$	0.88	2.14	3.017 (4)	176

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

All H atoms were included in the refinement at calculated positions, in the riding-model approximation, with an N-H distance of 0.88 Å and C-H distances of 0.95 (aromatic H atoms), 0.99 (CH₂ H atoms) and 0.98 Å (CH₃ H atoms). The isotropic displacement parameters were set equal to $1.25U_{\rm eq}$ of the carrier atom. The high $R_{\rm int}$ value for the title compound was the result of weak high-angle data.

Data collection: *DENZO* (Otwinowski & Minor, 1997) and *COLLECT* (Hooft, 1998); cell refinement: *DENZO* and *COLLECT*; data reduction: *DENZO*, *SCALEPACK* (Otwinowski & Minor, 1997) and *COLLECT*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *PLATON*97 (Spek, 1997); software used to prepare material for publication: *SHELXL*97.

The authors thank the EPSRC National Crystallography Service (Southampton, England) and the EPSRC Chemical Database Service at Daresbury Laboratory, England (Fletcher *et al.*, 1996).

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