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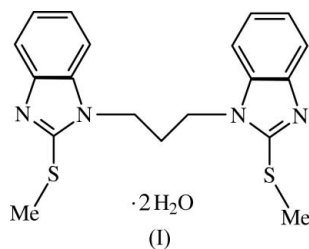
Key indicators

Single-crystal X-ray study
 $T = 296$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.042
 wR factor = 0.119
Data-to-parameter ratio = 17.4For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.1,1'-(Propane-1,3-diyl)bis[2-(methylsulfanyl)-1*H*-benzimidazole] dihydrate

In the title compound, $\text{C}_{19}\text{H}_{20}\text{N}_4\text{S}_2 \cdot 2\text{H}_2\text{O}$, the two 1*H*-benzimidazole groups are bridged by a propane chain; the dihedral angle between the benzimidazole ring systems is $74.61(3)^\circ$. The organic molecules and solvent water molecules form $\text{O}-\text{H} \cdots \text{N}$ hydrogen bonds. There are also $\text{O}-\text{H} \cdots \text{O}$, $\text{C}-\text{H} \cdots \text{S}$, $\text{C}-\text{H} \cdots \pi$ and $\pi-\pi$ interactions in the crystal structure.

Comment

The benzimidazole ring system has an interesting chemistry, and it is an effective pharmacophore in medicinal chemistry. Bis-benzimidazoles have potent activity against a number of microorganisms, including those that lead to AIDS-related infections (Bell *et al.*, 1993). These compounds bind to DNA in AT-rich sequences. Recently, drugs derived from benzimidazole have received much attention owing to the fact that the benzimidazole residue is a constituent of vitamin B_{12} (Skalitzky *et al.*, 2003), which supports their potential use as therapeutics (Hauel *et al.*, 2002; Valdez *et al.*, 2002). Benzimidazole, sometimes called 1,3-dideazapurine, and its derivatives can serve as model compounds for purine due to the structural similarity (Seela & Wenzel, 1995; Moreno *et al.*, 2004). In this paper, we report the crystal structure of the title compound, (I).



The asymmetric unit of (I) is shown Fig. 1. The organic molecule has two 1*H*-benzimidazole groups which are linked by a propane chain. The dihedral angles between benzimidazole plane *A* (C2–C7/N1/C1/N2), propane plane *B* (C8–C10) and the other benzimidazole plane *C* (C12–C17/N3/C11/N4) are $A/B = 85.66(12)$, $A/C = 74.61(3)$ and $B/C = 67.80(17)^\circ$. The geometric parameters for (I) agree well with those reported for other structures of bis(benzimidazole)s with rigid two-carbon bridges (Stibrany *et al.*, 2005). In particular, in the $\text{N}-\text{C}-\text{N}$ fragments, the $\text{C}1-\text{N}1$ and $\text{C}11-\text{N}4$ distances are *ca* 0.05 Å shorter than the $\text{C}1-\text{N}2$ and $\text{C}11-\text{N}3$ distances (Table 1), consistent with the partial double-bond character of the former bonds.

In (I), the organic molecules are linked by solvent water molecules, forming $\text{O}-\text{H} \cdots \text{N}$ hydrogen bonds (Table 2) in

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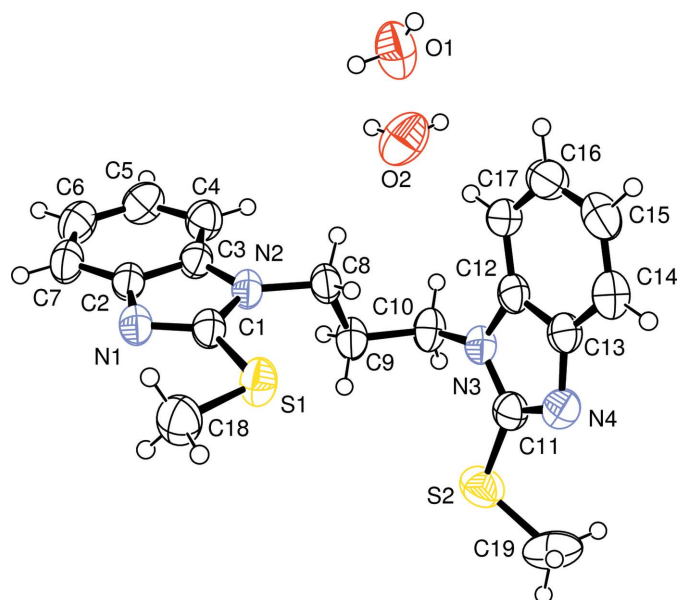


Figure 1
The asymmetric unit of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level.

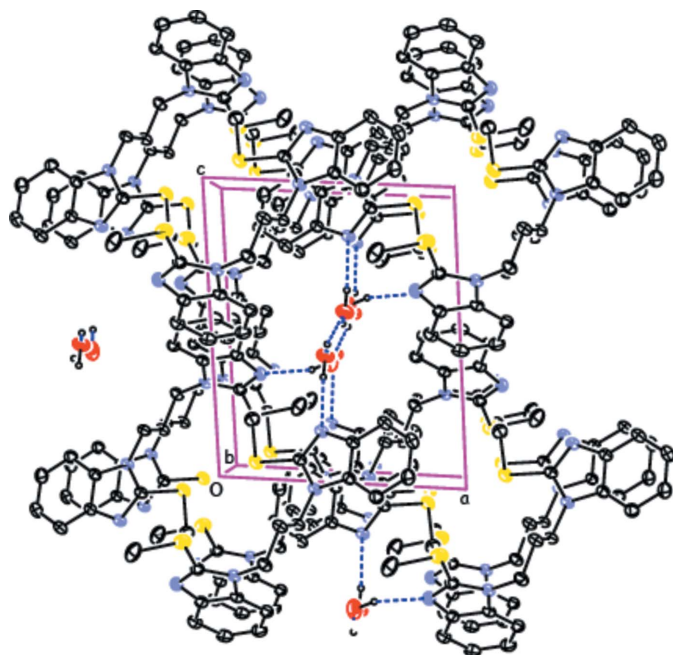


Figure 2
The crystal structure of (I). H atoms have been omitted unless they are involved in hydrogen bonds (dashed lines).

the crystal structure. As a result of these interactions, the structure exhibits channels of cavities passing through the centre of the unit cell and parallel to the *b* axis, with a mean diameter of 7.1807 (6) Å (Fig. 2). In addition, the other columns of cavity lie at the edges of the unit cell parallel to the *b* axis. There are also O—H···O hydrogen bonds between water molecules, intramolecular C—H···S, intermolecular C—H··· π and π – π interactions in the crystal structure (Fig. 3). The imidazole rings are oriented in such a way that the perpendicular distance between the the $Cg1$ (N1/C1/N2/C3/

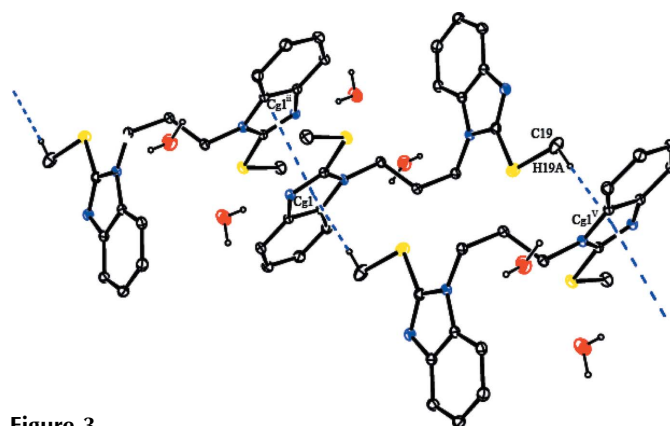


Figure 3
Intermolecular π – π and C—H··· π interactions (dashed lines). Other H atoms have been omitted. [Symmetry codes: (ii) $1 - x, -y, -z$; (v) $2 - x, 1 - y, -z$.]

$C2$) and $Cg1^{ii}$ rings is 3.478 Å, with the ring centroids separated by 3.5346 (11) Å [symmetry code: (ii) $1 - x, -y, -z$]. A C—H··· π interaction is observed between $Cg1$ and the C19 methyl group: H19A··· $Cg1^v = 2.5833$ Å, C19—H19A··· $Cg1^v = 175.1^\circ$ [symmetry code: (v) $2 - x, 1 - y, -z$].

Experimental

To a mixture of 2-methylthio-1*H*-benzimidazole (0.91 g, 3.14 mmol) and finely powdered NaOH (0.5 g, 12.5 mmol) in DMSO (7 ml) was added dropwise 1,3-dibromopropane (0.35 ml, 3.45 mmol). The resulting solution was stirred at 308–313 K for 1 h. Water was then added to the reaction mixture, and the solid that precipitated was collected and recrystallized from ethanol (yield 85%, m.p. 391–393 K).

Crystal data

$C_{19}H_{20}N_4S_2 \cdot 2H_2O$
 $M_r = 404.56$
Triclinic, $P\bar{1}$
 $a = 9.2655$ (7) Å
 $b = 9.9754$ (7) Å
 $c = 11.2980$ (8) Å
 $\alpha = 102.125$ (6) $^\circ$
 $\beta = 93.415$ (6) $^\circ$
 $\gamma = 96.443$ (6) $^\circ$

$V = 1010.78$ (13) Å³
 $Z = 2$
 $D_x = 1.329$ Mg m⁻³
Mo $K\alpha$ radiation
 $\mu = 0.28$ mm⁻¹
 $T = 296$ K
Prism, colorless
 $0.57 \times 0.36 \times 0.18$ mm

Data collection

Stoe IPDS-2 area-detector diffractometer
 ω scans
Absorption correction: integration (*X-RED32*; Stoe & Cie, 2002)
 $T_{min} = 0.880$, $T_{max} = 0.958$

19069 measured reflections
4506 independent reflections
3322 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.062$
 $\theta_{max} = 27.3^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.119$
 $S = 1.04$
4506 reflections
259 parameters
H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.065P)^2 + 0.0747P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} = 0.002$
 $\Delta\rho_{max} = 0.19$ e Å⁻³
 $\Delta\rho_{min} = -0.55$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

S1—C1	1.7319 (19)	N2—C8	1.457 (2)
S2—C11	1.738 (2)	N3—C11	1.363 (2)
N1—C1	1.318 (2)	N3—C10	1.457 (2)
N2—C1	1.365 (2)	N4—C11	1.313 (3)
C18—S1—C1—N1	−6.5 (2)	C10—N3—C11—N4	176.71 (16)
N2—C8—C9—C10	174.45 (16)	C19—S2—C11—N4	−14.5 (2)
C8—C9—C10—N3	66.1 (2)		

Table 2

Hydrogen-bond geometry (Å, °).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
O2—H4W...N4 ⁱ	0.83 (3)	2.05 (3)	2.851 (3)	162 (5)
O1—H1W...N1 ⁱⁱ	0.820 (16)	2.088 (18)	2.905 (2)	175 (3)
O1—H2W...O1 ⁱⁱⁱ	0.82 (4)	2.10 (5)	2.905 (3)	167 (7)
O2—H3W...O2 ^{iv}	0.83 (2)	2.25 (4)	2.922 (4)	139 (4)
C8—H8A...S1	0.97	2.72	3.1284 (19)	106
C10—H10A...S2	0.97	2.79	3.138 (2)	102

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

Water H atoms were located in difference maps and refined isotropically with bond restraints O—H = 0.83 (2) Å. C-bound H atoms were positioned geometrically and treated as riding, with C—H = 0.93–0.97 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}(\text{methyl C})$.

Data collection: *X-AREA* (Stoe & Cie, 2002); cell refinement: *X-AREA*; data reduction: *X-RED32* (Stoe & Cie, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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