

Crystal structure of 1,5-diethyl-1*H*-1,5-benzodiazepine-2,4(3*H*,5*H*)-dithione

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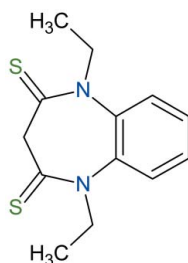
In the title compound, C₁₃H₁₆N₂S₂, the seven-membered ring adopts a boat conformation, with the two phenylene C atoms representing the stern and the methylene C atom as the prow. The thione S atoms and N-bound ethyl groups lie on the opposite side of the molecule to the phenylene ring so that the molecule approximates mirror symmetry. In the crystal, supramolecular layers in the *bc* plane are sustained by a pair of C—H...S interactions to the same S atom acceptor.

Keywords: crystal structure; benzodiazepine; boat conformation..

CCDC reference: 1040593

1. Related literature

For the biological activity of benzodiazepine derivatives, see: Kumar *et al.* (2006); Swamy *et al.* (2008). For a related structure, see: Ourahou *et al.* (2010).



2. Experimental

2.1. Crystal data

C ₁₃ H ₁₆ N ₂ S ₂	<i>V</i> = 2659.75 (5) Å ³
<i>M_r</i> = 264.40	<i>Z</i> = 8
Monoclinic, <i>C</i> 2/ <i>c</i>	Mo <i>K</i> α radiation
<i>a</i> = 19.8896 (2) Å	<i>μ</i> = 0.38 mm ⁻¹
<i>b</i> = 8.8743 (1) Å	<i>T</i> = 150 K
<i>c</i> = 15.5361 (2) Å	0.44 × 0.28 × 0.26 mm
<i>β</i> = 104.087 (1)°	

2.2. Data collection

Bruker APEXII CCD area-detector diffractometer	14634 measured reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2009)	3312 independent reflections
<i>T_{min}</i> = 0.880, <i>T_{max}</i> = 0.906	2963 reflections with <i>I</i> > 2σ(<i>I</i>)
	<i>R_{int}</i> = 0.022

2.3. Refinement

<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)] = 0.029	154 parameters
<i>wR</i> (<i>F</i> ²) = 0.083	H-atom parameters constrained
<i>S</i> = 1.05	Δ <i>ρ</i> _{max} = 0.39 e Å ⁻³
3312 reflections	Δ <i>ρ</i> _{min} = -0.21 e Å ⁻³

Table 1

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>
C3—H3...S1 ⁱ	0.95	2.86	3.6474 (13)	141
C12—H12A...S1 ⁱⁱ	0.99	2.87	3.4887 (13)	121

Symmetry codes: (i) *x*, -*y* + 2, *z* + ½; (ii) *x*, -*y* + 1, *z* + ½.

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT-Plus* (Bruker, 2009); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *pubCIF* (Westrip, 2010).

Supporting information for this paper is available from the IUCr electronic archives (Reference: TK5353).

References

- Bruker (2009). *APEX2*, *SADABS* and *SAINT-Plus*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Kumar, R., Chaudhary, P., Nimesh, S., Verma, A. K. & Chandra, R. (2006). *Green Chem.* **8**, 519–521.
- Ourahou, S., Zouihri, H., Essassi, E. M. & Ng, S. W. (2010). *Acta Cryst.* **E66**, o1653.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Spek, A. L. (2009). *Acta Cryst.* **D65**, 148–155.
- Swamy, G. Y. S. K., Sridhar, B., Ravikumar, K., Reddy, K. S. & Reddy, V. V. N. (2008). *J. Struct. Chem.* **49**, 775–779.
- Westrip, S. P. (2010). *J. Appl. Cryst.* **43**, 920–925.

supporting information

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Crystal structure of 1,5-diethyl-1*H*-1,5-benzodiazepine-2,4(3*H*,5*H*)-dithione

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S1. Introduction

Benzodiazepines and their derivatives are an important class of bioactive compound. They have attracted attention of chemists in the field of pharmaceuticals (Kumar *et al.*, 2006). Some benzodiazepine derivatives have been widely used as anti-bacterial, anti-fungal, analgesic and anti-convulsant agents (Swamy *et al.* 2008).

S2. Synthesis and crystallization

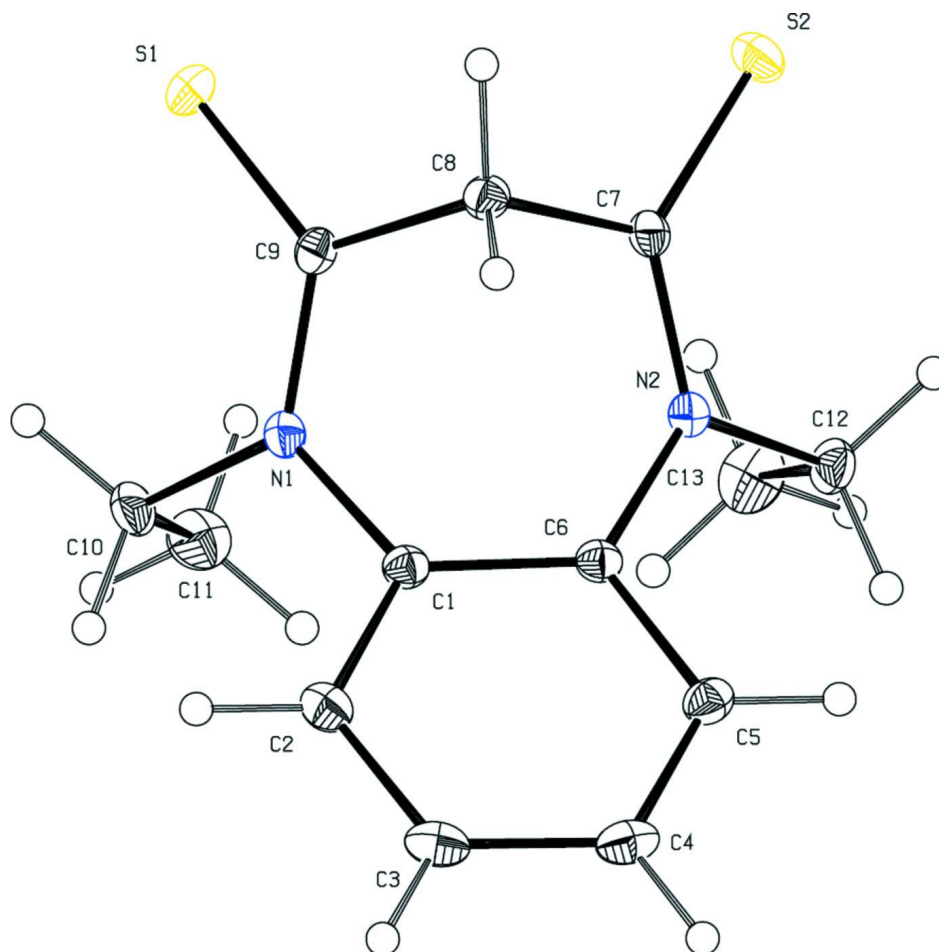
In a round flask, the 1,5 diethyl benzodiazepine-2,4-diones (2,22 g, 10 ml) and P₂S₅ (4.44 g, 20 ml) were mixed in acetonitrile (50 ml). The mixture was refluxed for 4 h. After this time, the solvent was evaporated, and the residue formed was washed with HCl (2 N) solution and distilled water, dried and recrystallised from toluene-chloroform (90/10). After some days, pale-yellow crystals were isolated (yield: 92.1 %, 2.34 g).

S3. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 1. H-atoms were placed in calculated positions (C—H 0.95–0.99 Å) and were included in the refinement in the riding model approximation, with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl H atoms and $1.2U_{\text{eq}}(\text{C})$ for all other H atoms.

S4. Results and discussion

The title compound crystallizes in the space group C2/c with one independent molecule in the asymmetric unit (Fig. 1). In the molecule, the diazepine ring system adopts a boat conformation with the two C1 and C6 atoms representing the stern and the C8 atom the prow with maximum deviation of 0.6626 (12) Å. The puckering parameters are: $q_2 = 0.9557$ (11) Å, $q_3 = 0.2328$ (11) Å, $\varphi_2 = 29.60$ (7)° and $\varphi_3 = 128.4$ (3)°. The mean plane of the diazepine ring is twisted with respect to that of the benzene ring by 32.27 (5)°. The geometric parameters of the title compound are comparable to those reported for similar structures (Ourahou *et al.*, 2010).

**Figure 1**

The structure of the title compound, showing atom labelling and 30% probability displacement ellipsoids.

1,5-Diethyl-1H-1,5-benzodiazepine-2,4(3H,5H)-dithione

Crystal data

$C_{13}H_{16}N_2S_2$

$M_r = 264.40$

Monoclinic, $C2/c$

Hall symbol: $-C\ 2yc$

$a = 19.8896\ (2)\ \text{\AA}$

$b = 8.8743\ (1)\ \text{\AA}$

$c = 15.5361\ (2)\ \text{\AA}$

$\beta = 104.087\ (1)^\circ$

$V = 2659.75\ (5)\ \text{\AA}^3$

$Z = 8$

$F(000) = 1120$

$D_x = 1.321\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 7669 reflections

$\theta = 2.5\text{--}28.3^\circ$

$\mu = 0.38\ \text{mm}^{-1}$

$T = 150\ \text{K}$

Block, pale-yellow

$0.44 \times 0.28 \times 0.26\ \text{mm}$

Data collection

Bruker APEXII CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2009)

$T_{\min} = 0.880$, $T_{\max} = 0.906$

14634 measured reflections

3312 independent reflections

2963 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.022$
 $\theta_{\text{max}} = 28.3^\circ$, $\theta_{\text{min}} = 2.1^\circ$

$h = -26 \rightarrow 26$
 $k = -11 \rightarrow 9$
 $l = -20 \rightarrow 20$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.029$
 $wR(F^2) = 0.083$
 $S = 1.05$
 3312 reflections
 154 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0451P)^2 + 1.6941P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.39 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.21 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.389586 (17)	0.69948 (4)	0.221477 (18)	0.02667 (9)
S2	0.401294 (17)	0.32244 (3)	0.37615 (2)	0.02698 (9)
N1	0.38266 (5)	0.81930 (10)	0.37619 (6)	0.01868 (19)
N2	0.38030 (5)	0.54640 (10)	0.48109 (6)	0.01790 (19)
C1	0.39904 (5)	0.82159 (12)	0.47124 (7)	0.0180 (2)
C2	0.41810 (6)	0.95882 (13)	0.51444 (8)	0.0236 (2)
H2	0.4222	1.0462	0.4808	0.028*
C3	0.43116 (6)	0.96826 (15)	0.60624 (8)	0.0270 (3)
H3	0.4443	1.0618	0.6353	0.032*
C4	0.42499 (6)	0.84098 (15)	0.65542 (8)	0.0258 (3)
H4	0.4323	0.8482	0.7180	0.031*
C5	0.40829 (6)	0.70361 (14)	0.61388 (7)	0.0217 (2)
H5	0.4055	0.6164	0.6483	0.026*
C6	0.39547 (5)	0.69179 (12)	0.52125 (7)	0.0176 (2)
C7	0.41267 (5)	0.49358 (12)	0.42024 (7)	0.0183 (2)
C8	0.45893 (6)	0.60552 (13)	0.38859 (7)	0.0196 (2)
H8A	0.4898	0.6575	0.4396	0.024*
H8B	0.4878	0.5541	0.3539	0.024*
C9	0.41007 (5)	0.71650 (12)	0.33112 (7)	0.0185 (2)
C10	0.32917 (6)	0.92648 (13)	0.32921 (8)	0.0255 (2)
H10A	0.3380	0.9521	0.2709	0.031*
H10B	0.3320	1.0205	0.3642	0.031*

C11	0.25747 (7)	0.85946 (17)	0.31559 (9)	0.0341 (3)
H11A	0.2230	0.9323	0.2846	0.051*
H11B	0.2485	0.8353	0.3733	0.051*
H11C	0.2544	0.7673	0.2801	0.051*
C12	0.32830 (6)	0.45201 (14)	0.50965 (8)	0.0245 (2)
H12A	0.3262	0.4814	0.5704	0.029*
H12B	0.3427	0.3450	0.5112	0.029*
C13	0.25706 (7)	0.46914 (18)	0.44750 (10)	0.0374 (3)
H13A	0.2239	0.4054	0.4681	0.056*
H13B	0.2588	0.4385	0.3875	0.056*
H13C	0.2424	0.5746	0.4467	0.056*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.03626 (18)	0.02858 (17)	0.01620 (14)	0.00172 (12)	0.00835 (11)	0.00226 (10)
S2	0.03522 (18)	0.01756 (15)	0.02775 (16)	0.00255 (11)	0.00685 (12)	-0.00303 (10)
N1	0.0206 (4)	0.0171 (5)	0.0179 (4)	0.0015 (3)	0.0039 (3)	0.0023 (3)
N2	0.0195 (4)	0.0165 (4)	0.0182 (4)	0.0003 (3)	0.0055 (3)	0.0014 (3)
C1	0.0159 (5)	0.0196 (5)	0.0186 (5)	0.0016 (4)	0.0044 (4)	-0.0005 (4)
C2	0.0233 (5)	0.0197 (5)	0.0279 (6)	-0.0003 (4)	0.0064 (4)	-0.0024 (4)
C3	0.0249 (6)	0.0270 (6)	0.0289 (6)	-0.0001 (5)	0.0062 (4)	-0.0108 (5)
C4	0.0221 (5)	0.0356 (7)	0.0196 (5)	0.0033 (5)	0.0047 (4)	-0.0066 (5)
C5	0.0193 (5)	0.0274 (6)	0.0191 (5)	0.0030 (4)	0.0059 (4)	0.0009 (4)
C6	0.0145 (5)	0.0192 (5)	0.0191 (5)	0.0019 (4)	0.0044 (4)	-0.0006 (4)
C7	0.0188 (5)	0.0180 (5)	0.0168 (4)	0.0043 (4)	0.0017 (4)	0.0031 (4)
C8	0.0181 (5)	0.0218 (5)	0.0198 (5)	0.0028 (4)	0.0063 (4)	0.0013 (4)
C9	0.0190 (5)	0.0185 (5)	0.0188 (5)	-0.0020 (4)	0.0065 (4)	0.0020 (4)
C10	0.0309 (6)	0.0198 (5)	0.0237 (5)	0.0078 (5)	0.0027 (4)	0.0046 (4)
C11	0.0255 (6)	0.0398 (8)	0.0349 (7)	0.0099 (6)	0.0035 (5)	0.0022 (6)
C12	0.0267 (6)	0.0230 (6)	0.0254 (5)	-0.0043 (5)	0.0097 (4)	0.0039 (4)
C13	0.0243 (6)	0.0473 (8)	0.0403 (7)	-0.0084 (6)	0.0071 (5)	0.0031 (6)

Geometric parameters (Å, °)

S1—C9	1.6591 (11)	C5—H5	0.9500
S2—C7	1.6587 (11)	C7—C8	1.5150 (15)
N1—C9	1.3437 (15)	C8—C9	1.5131 (15)
N1—C1	1.4329 (13)	C8—H8A	0.9900
N1—C10	1.4803 (14)	C8—H8B	0.9900
N2—C7	1.3506 (14)	C10—C11	1.5118 (18)
N2—C6	1.4329 (14)	C10—H10A	0.9900
N2—C12	1.4805 (14)	C10—H10B	0.9900
C1—C2	1.3976 (15)	C11—H11A	0.9800
C1—C6	1.4008 (15)	C11—H11B	0.9800
C2—C3	1.3886 (17)	C11—H11C	0.9800
C2—H2	0.9500	C12—C13	1.5157 (17)
C3—C4	1.3858 (19)	C12—H12A	0.9900

C3—H3	0.9500	C12—H12B	0.9900
C4—C5	1.3815 (17)	C13—H13A	0.9800
C4—H4	0.9500	C13—H13B	0.9800
C5—C6	1.4030 (15)	C13—H13C	0.9800
C9—N1—C1	121.84 (9)	C9—C8—H8B	110.7
C9—N1—C10	120.88 (9)	C7—C8—H8B	110.7
C1—N1—C10	117.06 (9)	H8A—C8—H8B	108.8
C7—N2—C6	122.11 (9)	N1—C9—C8	114.72 (9)
C7—N2—C12	120.02 (10)	N1—C9—S1	124.58 (8)
C6—N2—C12	117.86 (9)	C8—C9—S1	120.56 (8)
C2—C1—C6	119.65 (10)	N1—C10—C11	110.88 (10)
C2—C1—N1	118.31 (10)	N1—C10—H10A	109.5
C6—C1—N1	122.03 (9)	C11—C10—H10A	109.5
C3—C2—C1	120.45 (11)	N1—C10—H10B	109.5
C3—C2—H2	119.8	C11—C10—H10B	109.5
C1—C2—H2	119.8	H10A—C10—H10B	108.1
C4—C3—C2	119.84 (11)	C10—C11—H11A	109.5
C4—C3—H3	120.1	C10—C11—H11B	109.5
C2—C3—H3	120.1	H11A—C11—H11B	109.5
C5—C4—C3	120.34 (11)	C10—C11—H11C	109.5
C5—C4—H4	119.8	H11A—C11—H11C	109.5
C3—C4—H4	119.8	H11B—C11—H11C	109.5
C4—C5—C6	120.55 (11)	N2—C12—C13	111.43 (10)
C4—C5—H5	119.7	N2—C12—H12A	109.3
C6—C5—H5	119.7	C13—C12—H12A	109.3
C1—C6—C5	119.09 (10)	N2—C12—H12B	109.3
C1—C6—N2	122.18 (9)	C13—C12—H12B	109.3
C5—C6—N2	118.72 (10)	H12A—C12—H12B	108.0
N2—C7—C8	115.48 (9)	C12—C13—H13A	109.5
N2—C7—S2	124.50 (9)	C12—C13—H13B	109.5
C8—C7—S2	119.93 (8)	H13A—C13—H13B	109.5
C9—C8—C7	105.35 (9)	C12—C13—H13C	109.5
C9—C8—H8A	110.7	H13A—C13—H13C	109.5
C7—C8—H8A	110.7	H13B—C13—H13C	109.5
C9—N1—C1—C2	131.40 (11)	C12—N2—C6—C5	47.51 (13)
C10—N1—C1—C2	-54.03 (14)	C6—N2—C7—C8	-7.63 (14)
C9—N1—C1—C6	-49.55 (15)	C12—N2—C7—C8	173.63 (9)
C10—N1—C1—C6	125.02 (11)	C6—N2—C7—S2	175.84 (8)
C6—C1—C2—C3	-2.23 (17)	C12—N2—C7—S2	-2.90 (14)
N1—C1—C2—C3	176.84 (10)	N2—C7—C8—C9	-71.66 (11)
C1—C2—C3—C4	-0.26 (18)	S2—C7—C8—C9	105.04 (9)
C2—C3—C4—C5	2.27 (18)	C1—N1—C9—C8	-0.07 (15)
C3—C4—C5—C6	-1.77 (17)	C10—N1—C9—C8	-174.43 (10)
C2—C1—C6—C5	2.70 (16)	C1—N1—C9—S1	175.67 (8)
N1—C1—C6—C5	-176.34 (10)	C10—N1—C9—S1	1.30 (15)
C2—C1—C6—N2	-176.18 (10)	C7—C8—C9—N1	77.46 (11)

N1—C1—C6—N2	4.78 (16)	C7—C8—C9—S1	-98.46 (10)
C4—C5—C6—C1	-0.73 (16)	C9—N1—C10—C11	87.54 (13)
C4—C5—C6—N2	178.20 (10)	C1—N1—C10—C11	-87.08 (12)
C7—N2—C6—C1	47.63 (15)	C7—N2—C12—C13	-85.53 (13)
C12—N2—C6—C1	-133.60 (11)	C6—N2—C12—C13	95.68 (12)
C7—N2—C6—C5	-131.26 (11)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C3—H3 \cdots S1 ⁱ	0.95	2.86	3.6474 (13)	141
C12—H12 <i>A</i> \cdots S1 ⁱⁱ	0.99	2.87	3.4887 (13)	121

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