

rac-2-Hydroxy-2,8-dimethyl-4-morpholinoethyl-1-thia-4-azaspiro[4.5]decan-3-one

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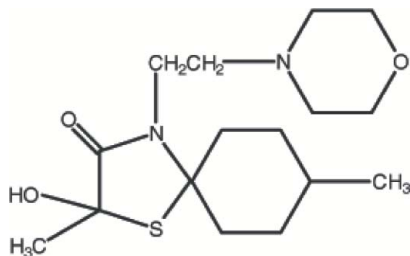
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.036; wR factor = 0.094; data-to-parameter ratio = 18.6.

The title compound, $\text{C}_{16}\text{H}_{28}\text{N}_2\text{O}_3\text{S}$, is dimerized by inversion symmetry-related intermolecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bonding, forming an $R_2^2(16)$ motif. The dimers are also linked through intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonding. The compound is chiral with a stereogenic centre located in the thiazole ring, but in the crystal structure it forms a racemate. The thiazole ring has an envelope conformation, while the cyclohexane and morpholine rings adopt chair conformations.

Related literature

For general background, see: Andres *et al.* (2000); Çapan *et al.* (1999); Srivastava *et al.* (2005). For related literature and bond-length data, see: Akkurt *et al.* (2007); Akkurt, Yalçın, Güzeldemirci *et al.* (2008); Akkurt, Yalçın, Klip *et al.* (2008). For ring conformation puckering parameters, see: Cremer & Pople (1975). For hydrogen-bond motifs, see: Bernstein *et al.* (1995).



Experimental

Crystal data

$\text{C}_{16}\text{H}_{28}\text{N}_2\text{O}_3\text{S}$
 $M_r = 328.47$

Triclinic, $P\bar{1}$
 $a = 8.0753$ (4) Å

$b = 10.2002$ (5) Å
 $c = 11.8734$ (6) Å
 $\alpha = 82.467$ (4)°
 $\beta = 71.487$ (4)°
 $\gamma = 68.965$ (4)°
 $V = 865.44$ (8) Å³

$Z = 2$
Mo $K\alpha$ radiation
 $\mu = 0.20$ mm⁻¹
 $T = 296$ K
 $0.73 \times 0.45 \times 0.29$ mm

Data collection

Stoe IPDS-2 diffractometer
Absorption correction: integration
(*X-RED32*; Stoe & Cie, 2002)
 $T_{\min} = 0.867$, $T_{\max} = 0.944$

18220 measured reflections
3693 independent reflections
3319 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.035$
 $wR(F^2) = 0.093$
 $S = 1.04$
3693 reflections

199 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.31$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.21$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|--|-------|-------------|-------------|---------------|
| $\text{O3}-\text{H3}\cdots\text{N2}^i$ | 0.82 | 2.00 | 2.8104 (14) | 169 |
| $\text{C7}-\text{H7B}\cdots\text{S1}$ | 0.97 | 2.82 | 3.2235 (18) | 106 |
| $\text{C14}-\text{H14B}\cdots\text{O1}^{ii}$ | 0.97 | 2.52 | 3.221 (2) | 129 |

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

Data collection: *X-AREA* (Stoe & Cie, 2002); cell refinement: *X-AREA*; data reduction: *X-RED32* (Stoe & Cie, 2002); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PLATON* (Spek, 2003).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: KP2184).

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supplementary materials

Acta Cryst. (2008). E64, o1574-o1575 [doi:10.1107/S1600536808022447]

***rac*-2-Hydroxy-2,8-dimethyl-4-morpholinoethyl-1-thia-4-azaspiro[4.5]decan-3-one**

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Comment

Thiazolidinones and their spiroheterocyclic analogues have been reported to exhibit antibacterial (Andres *et al.*, 2000), antifungal (Çapan *et al.*, 1999) and antimycobacterial (Srivastava *et al.*, 2005) activity. In view of these observations, we synthesized the title spiro[4.5]decan derivative as a racemate and report its crystal structure.

In the title molecule (Fig. 1), the values of the geometric parameters are normal and comparable with those in the similar compound, 8-methyl-4-morpholinoethyl-1-thia-4-azaspiro[4.5]decan-3-one (Akkurt, Yalçın, Klip *et al.*, 2008), which is a spiro[4.5]decan derivative.

The title compound is dimerized by inversion-symmetry related intermolecular O—H \cdots N hydrogen bonds, forming an $R_2^2(16)$ motif (Bernstein *et al.*, 1995) (Fig. 2). The dimers are interlinked through intermolecular C—H \cdots O hydrogen bonds. The compound is chiral with a stereogenic centre C1, in the thiazole ring. As the structure is centrosymmetric, racemate occurs in the crystal. The thiazole ring (C1–C3/S1/N1) has an envelope conformation on S1 [puckering parameters (Cremer & Pople, 1975): $Q(2) = 0.1885(12)$ Å, $\varphi(2) = 5.7(4)^\circ$]. The cyclohexane and morpholine rings (C3–C8) and (C12–C15/N2/O2) adopt chair conformations [puckering parameters: $Q_T = 0.563(2)$ Å, $\theta = 177.6(2)^\circ$, $\varphi = 58(4)^\circ$, and $Q_T = 0.576(2)$ Å, $\theta = 0.28(15)^\circ$, $\varphi = 340(60)^\circ$, respectively].

The structure is stabilized by intramolecular C—H \cdots S and intermolecular O—H \cdots N and C—H \cdots O hydrogen bonds (Table 1, Fig. 2).

Experimental

A mixture of morpholinoethylamin (5 mmol), 4-methyl cyclohexanone (5 mmol) and α -mercaptopropionic acid (20 mmol) in dry benzene (20 ml) was refluxed for 18 h using a Dean-Stark water separator. Excess solvent was evaporated *in vacuo*. The residue was taken up in chloroform. The chloroform layer was triturated with saturated NaHCO₃ solution (2x) before drying over sodium sulfate and concentrated under reduced pressure to dryness. The crude product was triturated with diethyl ether several times and recrystallized from ethanol to yield racemic mixture as colourless prisms. IR (ν , cm⁻¹): 1678 (C=O). ¹H-NMR (δ , DMSO-d₆, 400 MHz): 0.84 (3H, d, $J=6.0$ Hz, 8-CH₃), 0.96–1.03 (1H, m, cycl. CH), 1.15–1.24 (1H, m, cycl.CH), 1.32–1.42 (3H, m, cycl.CH), 1.51–1.52 (1H, m, cycl.CH), 1.66 (2H, d, $J=12.8$ Hz, cycl.CH), 1.79 (1H, dd, $J=12.8, 2.8$ Hz, cycl.CH), 1.95–2.10 (3H, m, SCHCH₃), 2.33–2.44 (6H, m, morph.N—CH₂), 3.30–3.37 (3H, m, SCH and N—CH₂), 3.54 (4H, t, $J=4.4$ Hz, OCH₂). LC—MS (m/z): 313 ($M+1$).

Refinement

H atoms were positioned geometrically, with C—H = 0.98–0.96 Å, O—H = 0.82 Å and constrained to ride on their parent atoms. The thermal parameter of H-atoms of methyl and hydroxyl groups was taken 1.5 times of the parent atom, whereas for all other H-atoms it was taken 1.2 times of their parent atoms.

Figures

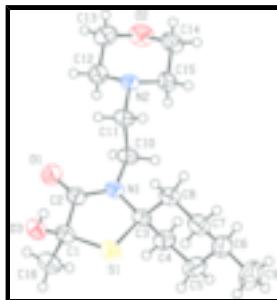


Fig. 1. The *ORTEP* diagram of the title compound with the atom numbering scheme. Displacement ellipsoids for non-H atoms are drawn at the 50% probability level.

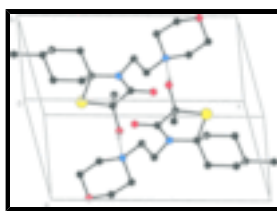


Fig. 2. View of the dimer generated by inversion symmetry related O—H...N hydrogen bonds, forming a $R_2^2(16)$ motif. For clarity, H atoms not involved in hydrogen bonds have been omitted.

rac-2-Hydroxy-2,8-dimethyl-4-morpholinoethyl-1-thia-4- azaspiro[4.5]decan-3-one

Crystal data

$C_{16}H_{28}N_2O_3S$

$M_r = 328.47$

Triclinic, $P\bar{1}$

Hall symbol: $-P\ 1$

$a = 8.0753\ (4)\ \text{\AA}$

$b = 10.2002\ (5)\ \text{\AA}$

$c = 11.8734\ (6)\ \text{\AA}$

$\alpha = 82.467\ (4)^\circ$

$\beta = 71.487\ (4)^\circ$

$\gamma = 68.965\ (4)^\circ$

$V = 865.44\ (8)\ \text{\AA}^3$

$Z = 2$

$F_{000} = 356$

$D_x = 1.260\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation

$\lambda = 0.71073\ \text{\AA}$

Cell parameters from 29023 reflections

$\theta = 2.1\text{--}27.4^\circ$

$\mu = 0.20\ \text{mm}^{-1}$

$T = 296\ \text{K}$

Block, colourless

$0.73 \times 0.45 \times 0.29\ \text{mm}$

Data collection

Stoe IPDS-2
diffractometer

Monochromator: plane graphite

3693 independent reflections

3319 reflections with $I > 2\sigma(I)$

Detector resolution: 6.67 pixels mm⁻¹
 $T = 296$ K
 ω scans
 Absorption correction: integration
 (X-RED32; Stoe & Cie, 2002)
 $T_{\min} = 0.867$, $T_{\max} = 0.944$
 18220 measured reflections

$R_{\text{int}} = 0.038$
 $\theta_{\max} = 26.9^\circ$
 $\theta_{\min} = 2.1^\circ$
 $h = -10 \rightarrow 10$
 $k = -12 \rightarrow 12$
 $l = -15 \rightarrow 15$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.093$
 $S = 1.04$
 3693 reflections
 199 parameters
 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
 $w = 1/[\sigma^2(F_o^2) + (0.0478P)^2 + 0.1524P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.31 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$
 Extinction correction: none

Special details

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

Refinement. Refinement on F^2 for ALL reflections except those flagged by the user for potential systematic errors. Weighted R -factors wR and all goodnesses 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 observed criterion of $F^2 > \sigma(F^2)$ is used only for calculating $-R$ -factor-obs *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 | 0.33771 (4) | 0.24227 (4) | 0.79561 (3) | 0.0486 (1) |
| O1 | 0.45350 (15) | 0.25388 (12) | 0.45525 (9) | 0.0555 (3) |
| O2 | 1.23062 (15) | 0.29531 (12) | 0.15285 (9) | 0.0597 (3) |
| O3 | 0.11938 (12) | 0.32395 (10) | 0.65362 (9) | 0.0480 (3) |
| N1 | 0.59680 (14) | 0.23670 (11) | 0.59509 (9) | 0.0389 (3) |
| N2 | 0.90885 (14) | 0.39307 (11) | 0.34820 (9) | 0.0372 (3) |
| C1 | 0.28811 (17) | 0.22315 (13) | 0.65981 (12) | 0.0417 (4) |
| C2 | 0.45358 (17) | 0.24085 (13) | 0.55883 (11) | 0.0408 (3) |
| C3 | 0.58346 (16) | 0.22120 (13) | 0.72210 (10) | 0.0369 (3) |
| C4 | 0.7085 (2) | 0.07643 (14) | 0.75151 (12) | 0.0482 (4) |

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| | | | | |
|------|--------------|--------------|--------------|------------|
| C5 | 0.7034 (2) | 0.06187 (17) | 0.88162 (14) | 0.0583 (5) |
| C6 | 0.7495 (2) | 0.17691 (19) | 0.92078 (13) | 0.0587 (5) |
| C7 | 0.6200 (2) | 0.31982 (17) | 0.89350 (13) | 0.0541 (5) |
| C8 | 0.62959 (19) | 0.33651 (14) | 0.76215 (12) | 0.0445 (4) |
| C9 | 0.7389 (3) | 0.1587 (3) | 1.05180 (17) | 0.0881 (8) |
| C10 | 0.76951 (17) | 0.23709 (13) | 0.50474 (11) | 0.0413 (3) |
| C11 | 0.75792 (17) | 0.38482 (13) | 0.45390 (11) | 0.0408 (3) |
| C12 | 0.90146 (19) | 0.34106 (16) | 0.24137 (11) | 0.0471 (4) |
| C13 | 1.0540 (2) | 0.36400 (19) | 0.13593 (12) | 0.0584 (5) |
| C14 | 1.2384 (2) | 0.34737 (18) | 0.25562 (13) | 0.0554 (5) |
| C15 | 1.09318 (17) | 0.32359 (15) | 0.36461 (11) | 0.0432 (4) |
| C16 | 0.2750 (2) | 0.08031 (15) | 0.65024 (15) | 0.0561 (5) |
| H3 | 0.12210 | 0.40260 | 0.65820 | 0.0720* |
| H4A | 0.66870 | 0.00470 | 0.73310 | 0.0580* |
| H4B | 0.83500 | 0.06140 | 0.70220 | 0.0580* |
| H5A | 0.58070 | 0.06390 | 0.93020 | 0.0700* |
| H5B | 0.79130 | -0.02850 | 0.89480 | 0.0700* |
| H6 | 0.87690 | 0.16960 | 0.87490 | 0.0700* |
| H7A | 0.65370 | 0.39320 | 0.91510 | 0.0650* |
| H7B | 0.49370 | 0.33050 | 0.94100 | 0.0650* |
| H8A | 0.54300 | 0.42730 | 0.74850 | 0.0530* |
| H8B | 0.75330 | 0.33370 | 0.71510 | 0.0530* |
| H9A | 0.82270 | 0.06800 | 1.06540 | 0.1320* |
| H9B | 0.61470 | 0.16630 | 1.09830 | 0.1320* |
| H9C | 0.77270 | 0.23040 | 1.07430 | 0.1320* |
| H10A | 0.79540 | 0.17570 | 0.44090 | 0.0500* |
| H10B | 0.87110 | 0.20060 | 0.53970 | 0.0500* |
| H11A | 0.64170 | 0.42750 | 0.43460 | 0.0490* |
| H11B | 0.75450 | 0.44050 | 0.51540 | 0.0490* |
| H12A | 0.78170 | 0.39070 | 0.22810 | 0.0570* |
| H12B | 0.91750 | 0.24180 | 0.25140 | 0.0570* |
| H13A | 1.05010 | 0.32880 | 0.06510 | 0.0700* |
| H13B | 1.03320 | 0.46390 | 0.12390 | 0.0700* |
| H14A | 1.21920 | 0.44710 | 0.24480 | 0.0670* |
| H14B | 1.36030 | 0.30050 | 0.26640 | 0.0670* |
| H15A | 1.11570 | 0.22360 | 0.37830 | 0.0520* |
| H15B | 1.09970 | 0.36120 | 0.43340 | 0.0520* |
| H16A | 0.17160 | 0.06840 | 0.71300 | 0.0840* |
| H16B | 0.38720 | 0.00790 | 0.65670 | 0.0840* |
| H16C | 0.25790 | 0.07430 | 0.57490 | 0.0840* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|----|------------|------------|------------|-------------|-------------|-------------|
| S1 | 0.0372 (2) | 0.0676 (2) | 0.0380 (2) | -0.0232 (2) | -0.0003 (1) | -0.0024 (1) |
| O1 | 0.0540 (6) | 0.0738 (7) | 0.0406 (5) | -0.0258 (5) | -0.0106 (4) | -0.0033 (4) |
| O2 | 0.0495 (6) | 0.0739 (7) | 0.0429 (5) | -0.0201 (5) | 0.0074 (4) | -0.0128 (5) |
| O3 | 0.0364 (5) | 0.0458 (5) | 0.0612 (6) | -0.0141 (4) | -0.0110 (4) | -0.0065 (4) |

| | | | | | | |
|-----|-------------|-------------|-------------|--------------|-------------|--------------|
| N1 | 0.0332 (5) | 0.0452 (6) | 0.0346 (5) | -0.0165 (4) | -0.0011 (4) | -0.0009 (4) |
| N2 | 0.0344 (5) | 0.0407 (5) | 0.0322 (5) | -0.0144 (4) | -0.0012 (4) | -0.0031 (4) |
| C1 | 0.0360 (6) | 0.0429 (6) | 0.0448 (7) | -0.0163 (5) | -0.0054 (5) | -0.0035 (5) |
| C2 | 0.0389 (6) | 0.0406 (6) | 0.0407 (6) | -0.0150 (5) | -0.0059 (5) | -0.0034 (5) |
| C3 | 0.0334 (5) | 0.0390 (6) | 0.0353 (6) | -0.0152 (5) | -0.0026 (4) | -0.0006 (4) |
| C4 | 0.0480 (7) | 0.0408 (7) | 0.0478 (7) | -0.0115 (6) | -0.0081 (6) | 0.0007 (5) |
| C5 | 0.0593 (9) | 0.0556 (8) | 0.0498 (8) | -0.0133 (7) | -0.0140 (7) | 0.0106 (6) |
| C6 | 0.0453 (7) | 0.0845 (11) | 0.0437 (7) | -0.0222 (7) | -0.0098 (6) | -0.0002 (7) |
| C7 | 0.0564 (8) | 0.0637 (9) | 0.0458 (7) | -0.0296 (7) | -0.0056 (6) | -0.0099 (6) |
| C8 | 0.0460 (7) | 0.0430 (7) | 0.0447 (7) | -0.0210 (5) | -0.0064 (5) | -0.0018 (5) |
| C9 | 0.0772 (13) | 0.1301 (19) | 0.0518 (10) | -0.0263 (12) | -0.0219 (9) | -0.0022 (10) |
| C10 | 0.0330 (6) | 0.0421 (6) | 0.0391 (6) | -0.0125 (5) | 0.0022 (5) | -0.0012 (5) |
| C11 | 0.0355 (6) | 0.0396 (6) | 0.0387 (6) | -0.0126 (5) | 0.0018 (5) | -0.0038 (5) |
| C12 | 0.0457 (7) | 0.0579 (8) | 0.0387 (6) | -0.0195 (6) | -0.0102 (5) | -0.0038 (5) |
| C13 | 0.0639 (9) | 0.0734 (10) | 0.0334 (7) | -0.0260 (8) | -0.0045 (6) | -0.0019 (6) |
| C14 | 0.0409 (7) | 0.0703 (9) | 0.0520 (8) | -0.0259 (7) | 0.0014 (6) | -0.0080 (7) |
| C15 | 0.0355 (6) | 0.0517 (7) | 0.0395 (6) | -0.0150 (5) | -0.0048 (5) | -0.0055 (5) |
| C16 | 0.0533 (8) | 0.0466 (8) | 0.0683 (9) | -0.0239 (6) | -0.0077 (7) | -0.0058 (6) |

Geometric parameters (Å, °)

| | | | |
|----------|-------------|-----------|--------|
| S1—C1 | 1.8295 (14) | C4—H4B | 0.9700 |
| S1—C3 | 1.8410 (14) | C5—H5A | 0.9700 |
| O1—C2 | 1.2197 (16) | C5—H5B | 0.9700 |
| O2—C13 | 1.412 (2) | C6—H6 | 0.9800 |
| O2—C14 | 1.4204 (19) | C7—H7A | 0.9700 |
| O3—C1 | 1.3990 (17) | C7—H7B | 0.9700 |
| O3—H3 | 0.8200 | C8—H8A | 0.9700 |
| N1—C3 | 1.4688 (15) | C8—H8B | 0.9700 |
| N1—C10 | 1.4659 (18) | C9—H9A | 0.9600 |
| N1—C2 | 1.3420 (19) | C9—H9B | 0.9600 |
| N2—C12 | 1.4633 (17) | C9—H9C | 0.9600 |
| N2—C15 | 1.4651 (19) | C10—H10A | 0.9700 |
| N2—C11 | 1.4636 (17) | C10—H10B | 0.9700 |
| C1—C2 | 1.535 (2) | C11—H11A | 0.9700 |
| C1—C16 | 1.520 (2) | C11—H11B | 0.9700 |
| C3—C8 | 1.524 (2) | C12—H12A | 0.9700 |
| C3—C4 | 1.5298 (19) | C12—H12B | 0.9700 |
| C4—C5 | 1.521 (2) | C13—H13A | 0.9700 |
| C5—C6 | 1.516 (2) | C13—H13B | 0.9700 |
| C6—C9 | 1.521 (2) | C14—H14A | 0.9700 |
| C6—C7 | 1.523 (2) | C14—H14B | 0.9700 |
| C7—C8 | 1.527 (2) | C15—H15A | 0.9700 |
| C10—C11 | 1.5297 (18) | C15—H15B | 0.9700 |
| C12—C13 | 1.513 (2) | C16—H16A | 0.9600 |
| C14—C15 | 1.506 (2) | C16—H16B | 0.9600 |
| C4—H4A | 0.9700 | C16—H16C | 0.9600 |
| S1...N1 | 2.6146 (11) | H5B...H9A | 2.4800 |
| S1...H5A | 2.8900 | H6...H8B | 2.5600 |

supplementary materials

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|--------------------------|-------------|---------------------------|--------|
| S1...H7B | 2.8200 | H6...H16A ^v | 2.5100 |
| O1...O3 | 2.8913 (15) | H7A...H9C | 2.5400 |
| O1...C11 | 3.187 (2) | H7A...H14A ^{vi} | 2.5400 |
| O1...C14 ⁱ | 3.221 (2) | H7B...S1 | 2.8200 |
| O1...C15 ⁱ | 3.230 (2) | H7B...O2 ^x | 2.8100 |
| O2...N2 | 2.8364 (16) | H7B...H5A | 2.5600 |
| O3...C15 ⁱⁱ | 3.3942 (17) | H7B...H9B | 2.5300 |
| O3...O1 | 2.8913 (15) | H8A...H12A ⁱⁱ | 2.5600 |
| O3...N2 ⁱⁱ | 2.8104 (14) | H8B...O3 ^v | 2.7800 |
| O1...H10A | 2.5400 | H8B...C10 | 2.7500 |
| O1...H11A | 2.6600 | H8B...C11 | 3.0700 |
| O1...H15B ⁱ | 2.7500 | H8B...H6 | 2.5600 |
| O1...H16C | 2.8000 | H8B...H10B | 2.3800 |
| O1...H14B ⁱ | 2.5200 | H8B...H11B | 2.4700 |
| O1...H16B ⁱⁱⁱ | 2.8200 | H8B...H14A ^{vi} | 2.4400 |
| O2...H9B ^{iv} | 2.7900 | H9A...H5B | 2.4800 |
| O2...H7B ^{iv} | 2.8100 | H9B...O2 ^x | 2.7900 |
| O3...H15B ⁱ | 2.6400 | H9B...C14 ^x | 3.0700 |
| O3...H8B ⁱ | 2.7800 | H9B...H5A | 2.5000 |
| O3...H14A ⁱⁱ | 2.9100 | H9B...H7B | 2.5300 |
| N1...S1 | 2.6146 (11) | H9B...H14B ^x | 2.4800 |
| N2...O2 | 2.8364 (16) | H9C...C12 ^{xi} | 3.0000 |
| N2...O3 ⁱⁱ | 2.8104 (14) | H9C...H7A | 2.5400 |
| N2...H3 ⁱⁱ | 2.0000 | H10A...O1 | 2.5400 |
| C2...C11 ⁱⁱ | 3.5847 (18) | H10A...C12 | 2.8100 |
| C8...C11 | 3.4925 (18) | H10A...H12B | 2.2700 |
| C11...C8 | 3.4925 (18) | H10B...C4 | 2.8100 |
| C11...C2 ⁱⁱ | 3.5847 (18) | H10B...C8 | 2.9100 |
| C11...O1 | 3.187 (2) | H10B...C15 | 2.7700 |
| C14...O1 ^v | 3.221 (2) | H10B...H4B | 2.2500 |
| C15...O1 ^v | 3.230 (2) | H10B...H8B | 2.3800 |
| C15...O3 ⁱⁱ | 3.3942 (17) | H10B...H15A | 2.3300 |
| C2...H11A | 2.8300 | H11A...O1 | 2.6600 |
| C4...H10B | 2.8100 | H11A...C2 | 2.8300 |
| C8...H14A ^{vi} | 2.8600 | H11A...H12A | 2.3700 |
| C8...H10B | 2.9100 | H11A...C11 ⁱⁱ | 3.0600 |
| C8...H11B | 2.9700 | H11A...H11A ⁱⁱ | 2.4000 |
| C10...H12B | 2.8600 | H11B...C8 | 2.9700 |
| C10...H4B | 2.8300 | H11B...H8B | 2.4700 |
| C10...H15A | 2.6800 | H11B...H15B | 2.5000 |
| C10...H8B | 2.7500 | H12A...H11A | 2.3700 |
| C11...H8B | 3.0700 | H12A...H8A ⁱⁱ | 2.5600 |
| C11...H3 ⁱⁱ | 2.7100 | H12B...C10 | 2.8600 |

| | | | |
|----------------------------|-------------|----------------------------|--------|
| C11···H11A ⁱⁱ | 3.0600 | H12B···H10A | 2.2700 |
| C12···H3 ⁱⁱ | 2.9300 | H12B···H15A | 2.4700 |
| C12···H10A | 2.8100 | H13B···H14A | 2.3400 |
| C12···H9C ^{vii} | 3.0000 | H14A···H13B | 2.3400 |
| C14···H9B ^{iv} | 3.0700 | H14A···O3 ⁱⁱ | 2.9100 |
| C14···H3 ⁱⁱ | 3.0800 | H14A···C8 ^{vi} | 2.8600 |
| C15···H10B | 2.7700 | H14A···H3 ⁱⁱ | 2.5900 |
| C15···H3 ⁱⁱ | 2.7400 | H14A···H7A ^{vi} | 2.5400 |
| H3···N2 ⁱⁱ | 2.0000 | H14A···H8B ^{vi} | 2.4400 |
| H3···C11 ⁱⁱ | 2.7100 | H14B···O1 ^v | 2.5200 |
| H3···C12 ⁱⁱ | 2.9300 | H14B···H9B ^{iv} | 2.4800 |
| H3···C14 ⁱⁱ | 3.0800 | H15A···C10 | 2.6800 |
| H3···C15 ⁱⁱ | 2.7400 | H15A···H10B | 2.3300 |
| H3···H14A ⁱⁱ | 2.5900 | H15A···H12B | 2.4700 |
| H4A···H15A ^{viii} | 2.5800 | H15A···H4A ^{viii} | 2.5800 |
| H4B···C10 | 2.8300 | H15B···O1 ^v | 2.7500 |
| H4B···H10B | 2.2500 | H15B···O3 ^v | 2.6400 |
| H5A···S1 | 2.8900 | H15B···H11B | 2.5000 |
| H5A···H7B | 2.5600 | H16A···H6 ⁱ | 2.5100 |
| H5A···H9B | 2.5000 | H16B···O1 ⁱⁱⁱ | 2.8200 |
| H5A···H5A ^{ix} | 2.3300 | H16C···O1 | 2.8000 |
| C1—S1—C3 | 94.96 (6) | C6—C7—H7B | 109.00 |
| C13—O2—C14 | 109.74 (12) | C8—C7—H7A | 109.00 |
| C1—O3—H3 | 109.00 | C8—C7—H7B | 109.00 |
| C2—N1—C10 | 118.36 (10) | H7A—C7—H7B | 108.00 |
| C3—N1—C10 | 121.36 (11) | C3—C8—H8A | 109.00 |
| C2—N1—C3 | 120.12 (11) | C3—C8—H8B | 109.00 |
| C11—N2—C15 | 113.24 (10) | C7—C8—H8A | 109.00 |
| C12—N2—C15 | 109.83 (11) | C7—C8—H8B | 109.00 |
| C11—N2—C12 | 113.29 (11) | H8A—C8—H8B | 108.00 |
| S1—C1—C2 | 104.53 (10) | C6—C9—H9A | 109.00 |
| S1—C1—C16 | 113.45 (10) | C6—C9—H9B | 109.00 |
| S1—C1—O3 | 110.65 (9) | C6—C9—H9C | 109.00 |
| O3—C1—C16 | 107.00 (12) | H9A—C9—H9B | 110.00 |
| C2—C1—C16 | 108.91 (11) | H9A—C9—H9C | 109.00 |
| O3—C1—C2 | 112.40 (11) | H9B—C9—H9C | 109.00 |
| O1—C2—C1 | 121.93 (14) | N1—C10—H10A | 109.00 |
| N1—C2—C1 | 113.93 (11) | N1—C10—H10B | 109.00 |
| O1—C2—N1 | 124.12 (13) | C11—C10—H10A | 109.00 |
| S1—C3—N1 | 103.79 (9) | C11—C10—H10B | 109.00 |
| S1—C3—C8 | 109.34 (9) | H10A—C10—H10B | 108.00 |
| N1—C3—C4 | 111.50 (10) | N2—C11—H11A | 108.00 |
| S1—C3—C4 | 110.49 (9) | N2—C11—H11B | 108.00 |
| C4—C3—C8 | 110.34 (12) | C10—C11—H11A | 108.00 |
| N1—C3—C8 | 111.20 (10) | C10—C11—H11B | 108.00 |

supplementary materials

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| C3—C4—C5 | 111.95 (11) | H11A—C11—H11B | 107.00 |
| C4—C5—C6 | 112.88 (13) | N2—C12—H12A | 110.00 |
| C5—C6—C7 | 109.51 (14) | N2—C12—H12B | 110.00 |
| C7—C6—C9 | 111.87 (16) | C13—C12—H12A | 110.00 |
| C5—C6—C9 | 110.62 (16) | C13—C12—H12B | 110.00 |
| C6—C7—C8 | 111.54 (12) | H12A—C12—H12B | 108.00 |
| C3—C8—C7 | 111.69 (12) | O2—C13—H13A | 109.00 |
| N1—C10—C11 | 111.71 (11) | O2—C13—H13B | 109.00 |
| N2—C11—C10 | 115.84 (11) | C12—C13—H13A | 109.00 |
| N2—C12—C13 | 109.15 (13) | C12—C13—H13B | 109.00 |
| O2—C13—C12 | 111.53 (12) | H13A—C13—H13B | 108.00 |
| O2—C14—C15 | 111.11 (14) | O2—C14—H14A | 109.00 |
| N2—C15—C14 | 109.61 (11) | O2—C14—H14B | 109.00 |
| C3—C4—H4A | 109.00 | C15—C14—H14A | 109.00 |
| C3—C4—H4B | 109.00 | C15—C14—H14B | 109.00 |
| C5—C4—H4A | 109.00 | H14A—C14—H14B | 108.00 |
| C5—C4—H4B | 109.00 | N2—C15—H15A | 110.00 |
| H4A—C4—H4B | 108.00 | N2—C15—H15B | 110.00 |
| C4—C5—H5A | 109.00 | C14—C15—H15A | 110.00 |
| C4—C5—H5B | 109.00 | C14—C15—H15B | 110.00 |
| C6—C5—H5A | 109.00 | H15A—C15—H15B | 108.00 |
| C6—C5—H5B | 109.00 | C1—C16—H16A | 109.00 |
| H5A—C5—H5B | 108.00 | C1—C16—H16B | 109.00 |
| C5—C6—H6 | 108.00 | C1—C16—H16C | 109.00 |
| C7—C6—H6 | 108.00 | H16A—C16—H16B | 109.00 |
| C9—C6—H6 | 108.00 | H16A—C16—H16C | 109.00 |
| C6—C7—H7A | 109.00 | H16B—C16—H16C | 109.00 |
| C3—S1—C1—O3 | -136.07 (10) | C12—N2—C15—C14 | 56.88 (15) |
| C3—S1—C1—C2 | -14.85 (9) | C12—N2—C11—C10 | 71.48 (15) |
| C3—S1—C1—C16 | 103.66 (11) | S1—C1—C2—N1 | 12.26 (13) |
| C1—S1—C3—N1 | 13.84 (9) | O3—C1—C2—O1 | -49.39 (17) |
| C1—S1—C3—C4 | -105.81 (9) | O3—C1—C2—N1 | 132.32 (12) |
| C1—S1—C3—C8 | 132.57 (9) | C16—C1—C2—O1 | 69.00 (17) |
| C14—O2—C13—C12 | -59.69 (17) | C16—C1—C2—N1 | -109.30 (13) |
| C13—O2—C14—C15 | 59.60 (16) | S1—C1—C2—O1 | -169.45 (11) |
| C2—N1—C10—C11 | 79.30 (14) | N1—C3—C8—C7 | -178.85 (12) |
| C3—N1—C2—O1 | 179.78 (12) | C8—C3—C4—C5 | 52.97 (16) |
| C10—N1—C2—O1 | -4.83 (19) | S1—C3—C8—C7 | 67.13 (14) |
| C3—N1—C10—C11 | -105.38 (13) | N1—C3—C4—C5 | 177.06 (13) |
| C2—N1—C3—C8 | -126.75 (13) | C4—C3—C8—C7 | -54.59 (16) |
| C10—N1—C3—C8 | 58.01 (15) | S1—C3—C4—C5 | -68.06 (15) |
| C10—N1—C2—C1 | 173.42 (10) | C3—C4—C5—C6 | -54.55 (19) |
| C2—N1—C3—C4 | 109.65 (14) | C4—C5—C6—C7 | 55.23 (18) |
| C2—N1—C3—S1 | -9.30 (13) | C4—C5—C6—C9 | 178.97 (16) |
| C10—N1—C3—S1 | 175.45 (9) | C5—C6—C7—C8 | -56.25 (18) |
| C10—N1—C3—C4 | -65.59 (15) | C9—C6—C7—C8 | -179.25 (16) |
| C3—N1—C2—C1 | -1.97 (16) | C6—C7—C8—C3 | 57.30 (18) |
| C11—N2—C12—C13 | 175.83 (12) | N1—C10—C11—N2 | -169.01 (11) |
| C15—N2—C11—C10 | -54.44 (15) | N2—C12—C13—O2 | 58.47 (17) |

| | | | |
|----------------|--------------|---------------|-------------|
| C15—N2—C12—C13 | -56.45 (15) | O2—C14—C15—N2 | -58.52 (16) |
| C11—N2—C15—C14 | -175.38 (11) | | |

Symmetry codes: (i) $x-1, y, z$; (ii) $-x+1, -y+1, -z+1$; (iii) $-x+1, -y, -z+1$; (iv) $x+1, y, z-1$; (v) $x+1, y, z$; (vi) $-x+2, -y+1, -z+1$; (vii) $x, y, z-1$; (viii) $-x+2, -y, -z+1$; (ix) $-x+1, -y, -z+2$; (x) $x-1, y, z+1$; (xi) $x, y, z+1$.

Hydrogen-bond geometry ($\text{\AA}, ^\circ$)

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|-----------------------------------|-------|-------------|-------------|---------------|
| O3—H3 \cdots N2 ⁱⁱ | 0.82 | 2.00 | 2.8104 (14) | 169 |
| C7—H7B \cdots S1 | 0.97 | 2.82 | 3.2235 (18) | 106 |
| C14—H14B \cdots O1 ^v | 0.97 | 2.52 | 3.221 (2) | 129 |

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

Fig. 1

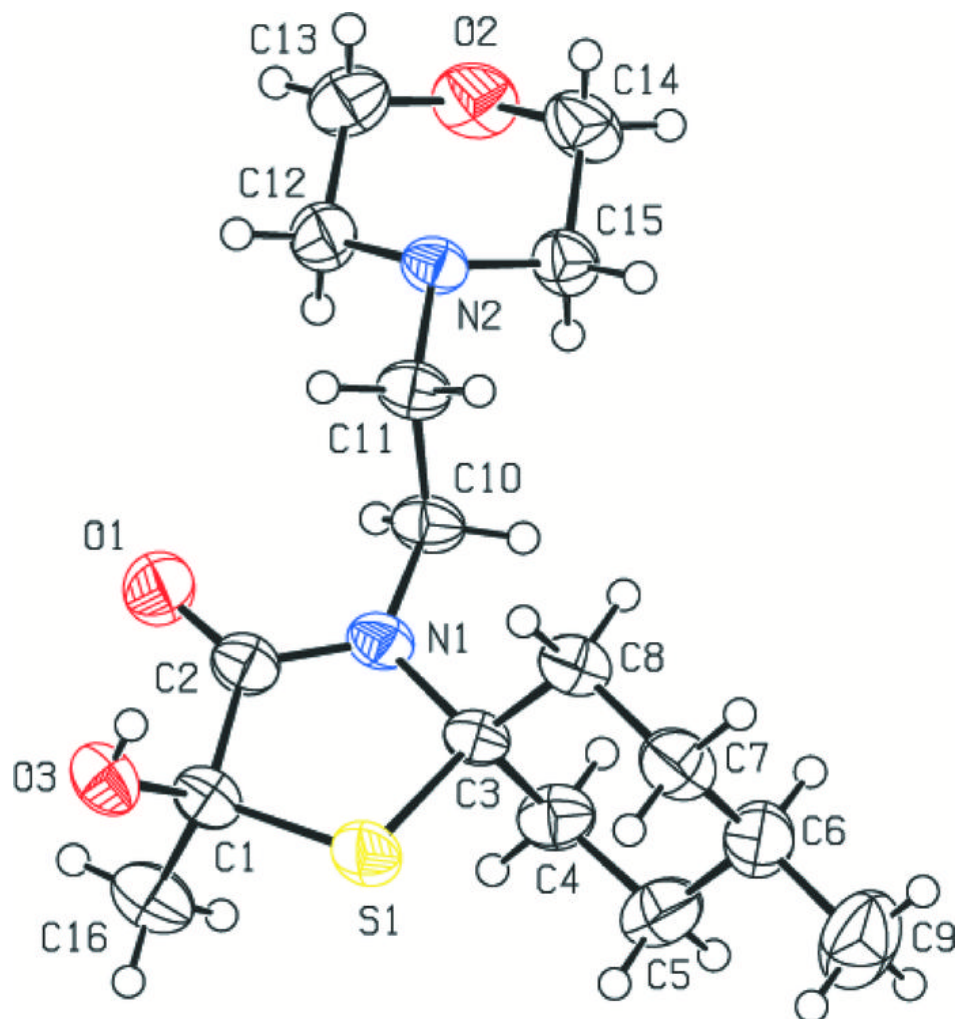


Fig. 2

