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1,3-Diethyl-2-sulfanylidene-5-(2,4,5-trimethoxybenzylidene)-1,3-diazinane-4,6-dione

 Abdullah M. Asiri,^{a,b*} Muhammad Nadeem Arshad,^{a,b} Muhammad Zia-ur-Rehman^{c*} and Tariq R. Sobahi^a

^aChemistry Department, Faculty of Science, King Abdulaziz University, PO Box 80203, Jeddah 21589, Saudi Arabia, ^bCenter of Excellence for Advanced Materials Research (CEAMR), Faculty of Science, King Abdulaziz University, PO Box 80203, Jeddah 21589, Saudi Arabia, and ^cApplied Chemistry Research Centre, PCSIR Laboratories Complex, Ferozpur Road, Lahore 54600, Pakistan
Correspondence e-mail: aasiri2@kau.edu.sa, rehman_pcsir@yahoo.com

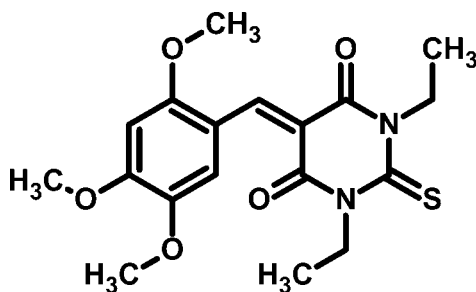
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 Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.041; wR factor = 0.121; data-to-parameter ratio = 15.7.

The title compound, $\text{C}_{18}\text{H}_{22}\text{N}_2\text{O}_5\text{S}$, is largely planar, with an r.m.s. deviation of 0.0546 (1) Å of atoms from the mean plane through all non-H atoms except for the methyl groups. The benzene and pyrimidinedione rings are inclined to one another at a dihedral angle of 1.41 (7)°. In the crystal, weak $\text{C}-\text{H}\cdots\text{O}$ interactions connect the molecules into chains propagating along the b -axis direction.

Related literature

For the synthesis of the title compound, see: Asiri *et al.* (2004). For a related structure, see: Asiri *et al.* (2009).



Experimental

Crystal data

$\text{C}_{18}\text{H}_{22}\text{N}_2\text{O}_5\text{S}$
 $M_r = 378.44$
 Monoclinic, $P2_1/c$
 $a = 7.9711$ (1) Å
 $b = 17.4106$ (3) Å
 $c = 13.5265$ (2) Å
 $\beta = 99.237$ (2)°
 $V = 1852.89$ (5) Å³
 $Z = 4$
 Cu $K\alpha$ radiation
 $\mu = 1.83$ mm⁻¹
 $T = 296$ K
 $0.29 \times 0.10 \times 0.09$ mm

Data collection

Agilent SuperNova (Dual, Cu at zero, Atlas, CCD) diffractometer
 Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2012)
 $T_{\min} = 0.875$, $T_{\max} = 1.000$
 14850 measured reflections
 3777 independent reflections
 3083 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.121$
 $S = 1.05$
 3777 reflections
 240 parameters
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.25$ e Å⁻³
 $\Delta\rho_{\min} = -0.20$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C13}-\text{H13C}\cdots\text{O2}^i$	0.96	2.58	3.455 (2)	152

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

Data collection: *CrysAlis PRO* (Agilent, 2012); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; 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: *WinGX* (Farrugia, 2012) and *X-SEED* (Barbour, 2001).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SJ5286).

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supporting information

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1,3-Diethyl-2-sulfanylidene-5-(2,4,5-trimethoxybenzylidene)-1,3-diazinane-4,6-dione

Abdullah M. Asiri, Muhammad Nadeem Arshad, Muhammad Zia-ur-Rehman and Tariq R. Sobahi

S1. Comment

The title compound is related to the arylidene 5-[3-(2,5-dimethoxyphenyl)prop-2-enylidene]-1,3-diethyl-2-thioxohexahydropyrimidine-4,6-dione (II) already reported by our group (Asiri *et al.*, 2009). The molecule is largely planar with a dihedral angle between the benzene and pyrimidine dione rings of 1.41 (7)°. The r. m. s. deviation of atoms from the best fit plane through the non-hydrogen atoms C1/N1/C2/C3/C4/N2/S1/O1/O2/C5/C6/C7/C8/C9/C10/C11/C12/C14/O3/O4/O5 is 0.0546 (1) Å. Atoms S1 and O1 are displaced from this plane by -0.1187 (1) Å and 0.1131 (2) Å respectively. Non-classical C13–H13C···O2 hydrogen bonds connect the molecule into chains along the *b* axis (Table. 1, Fig. 2).

S2. Experimental

1,3-Diethyl-2-thiobarbituric acid (0.005 mol) and 2,4,5-trimethoxybenzaldehyde (0.005 mol) were heated in ethanol (15 ml) for 3 h; several drops of diethylamine were added. The progress of reaction was monitored by TLC. The mixture was cooled and the resulting solid was recrystallized from methanol (Asiri *et al.* 2004) by slow evaporation at room temperature.

S3. Refinement

All the H-atoms bound to C were positioned with idealized geometry with C—H = 0.93 Å for aromatic, C—H = 0.97 Å for methylene & C—H = 0.96 Å for methyl groups. H-atoms were refined as riding with $U_{\text{iso}}(\text{H}) = kU_{\text{eq}}(\text{C})$, where $k = 1.5$ for methyl H-atoms & $k = 1.2$ for other H-atoms .

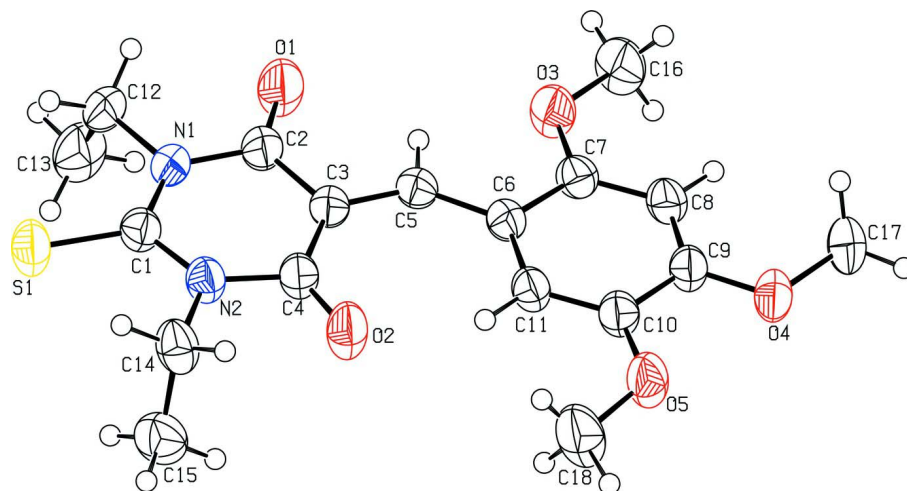


Figure 1

The structure of (I) with 50% probability ellipsoids.

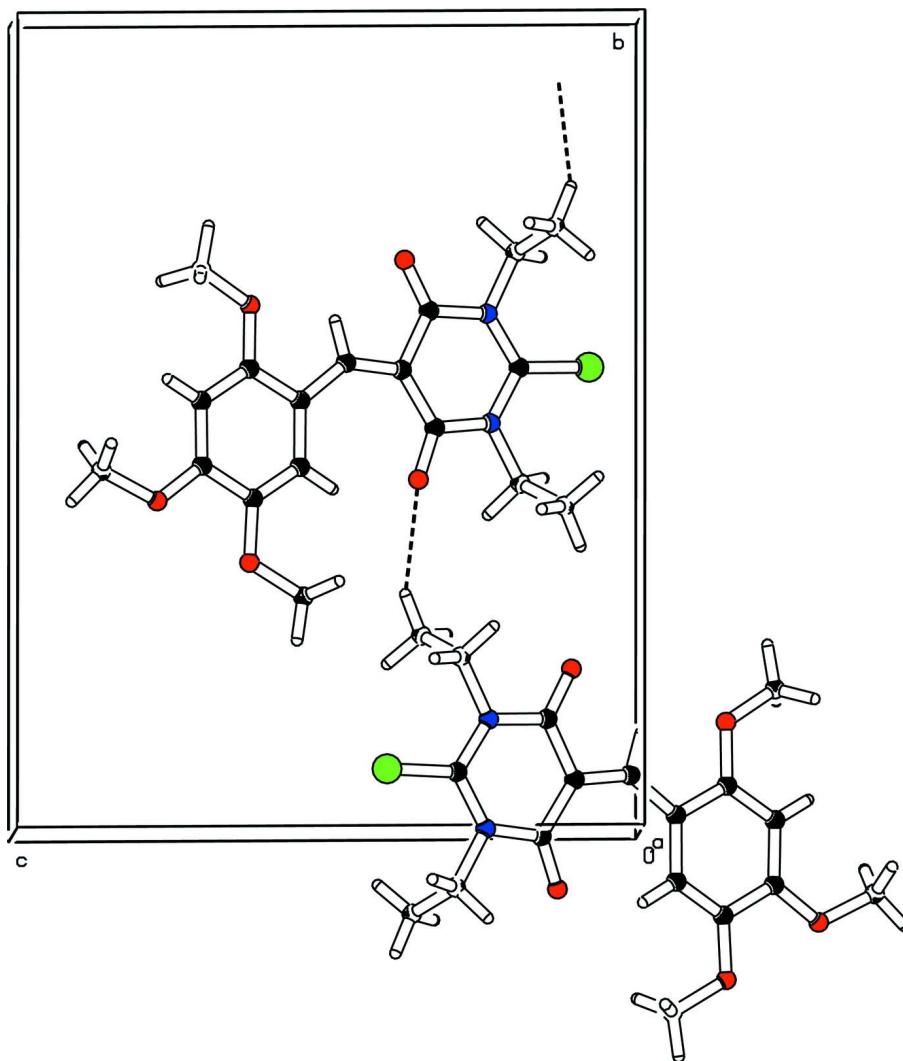


Figure 2

Unit cell diagram showing C—H...O hydrogen bonds, drawn as dashed lines.

1,3-Diethyl-2-sulfanylidene-5-(2,4,5-trimethoxybenzylidene)-1,3-diazinane-4,6-dione

Crystal data

$C_{18}H_{22}N_2O_5S$

$M_r = 378.44$

Monoclinic, $P2_1/c$

Hall symbol: $-P\ 2ybc$

$a = 7.9711(1)\ \text{\AA}$

$b = 17.4106(3)\ \text{\AA}$

$c = 13.5265(2)\ \text{\AA}$

$\beta = 99.237(2)^\circ$

$V = 1852.89(5)\ \text{\AA}^3$

$Z = 4$

$F(000) = 800$

$D_x = 1.357\ \text{Mg m}^{-3}$

Cu $K\alpha$ radiation, $\lambda = 1.54184\ \text{\AA}$

Cell parameters from 6212 reflections

$\theta = 3.3\text{--}75.4^\circ$

$\mu = 1.83\ \text{mm}^{-1}$

$T = 296\ \text{K}$

Needle like, red

$0.29 \times 0.10 \times 0.09\ \text{mm}$

Data collection

Agilent SuperNova (Dual, Cu at zero, Atlas, CCD) diffractometer
 Radiation source: SuperNova (Cu) X-ray Source
 Mirror monochromator
 ω scans
 Absorption correction: multi-scan (CrysAlis PRO; Agilent, 2012)

$T_{\min} = 0.875$, $T_{\max} = 1.000$
 14850 measured reflections
 3777 independent reflections
 3083 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$
 $\theta_{\max} = 75.6^\circ$, $\theta_{\min} = 4.2^\circ$
 $h = -9 \rightarrow 7$
 $k = -21 \rightarrow 21$
 $l = -16 \rightarrow 16$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.121$
 $S = 1.05$
 3777 reflections
 240 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
 $w = 1/[\sigma^2(F_o^2) + (0.0597P)^2 + 0.385P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.25 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.20 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 $F^2 > 2\sigma(F^2)$ 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	-0.22340 (6)	0.56530 (3)	0.08605 (3)	0.05970 (16)
O1	0.0851 (2)	0.69234 (8)	0.38430 (11)	0.0875 (5)
O2	0.10560 (19)	0.42242 (7)	0.36247 (9)	0.0651 (4)
O3	0.40703 (17)	0.63202 (7)	0.63548 (9)	0.0591 (3)
O4	0.58803 (17)	0.38883 (7)	0.78593 (9)	0.0595 (3)
O5	0.42004 (19)	0.31584 (7)	0.63615 (9)	0.0646 (4)
N1	-0.05338 (18)	0.62856 (8)	0.25049 (10)	0.0472 (3)
N2	-0.05463 (17)	0.49412 (7)	0.24558 (10)	0.0435 (3)
C1	-0.1055 (2)	0.56302 (9)	0.19924 (12)	0.0431 (4)
C2	0.0503 (2)	0.63052 (10)	0.34507 (13)	0.0527 (4)
C3	0.1117 (2)	0.55692 (9)	0.38958 (11)	0.0415 (3)
C4	0.0594 (2)	0.48671 (9)	0.33594 (11)	0.0428 (3)
C5	0.2186 (2)	0.56421 (8)	0.47980 (11)	0.0411 (3)
H5	0.2339	0.6157	0.4973	0.049*
C6	0.31256 (19)	0.51620 (9)	0.55427 (11)	0.0388 (3)

C7	0.4108 (2)	0.55426 (9)	0.63690 (11)	0.0420 (3)
C8	0.5034 (2)	0.51281 (10)	0.71486 (11)	0.0456 (4)
H8	0.5662	0.5387	0.7686	0.055*
C9	0.5032 (2)	0.43358 (10)	0.71329 (11)	0.0449 (4)
C10	0.4101 (2)	0.39422 (9)	0.63098 (11)	0.0454 (4)
C11	0.3173 (2)	0.43493 (9)	0.55483 (11)	0.0421 (3)
H11	0.2552	0.4083	0.5015	0.050*
C12	-0.1014 (3)	0.70471 (10)	0.20484 (14)	0.0615 (5)
H12A	-0.1120	0.7414	0.2574	0.074*
H12B	-0.2111	0.7007	0.1621	0.074*
C13	0.0275 (3)	0.73339 (11)	0.14422 (17)	0.0764 (6)
H13A	0.0335	0.6987	0.0897	0.115*
H13B	0.1367	0.7364	0.1859	0.115*
H13C	-0.0052	0.7834	0.1182	0.115*
C14	-0.1166 (2)	0.42108 (10)	0.19685 (13)	0.0537 (4)
H14A	-0.2290	0.4291	0.1586	0.064*
H14B	-0.1263	0.3830	0.2480	0.064*
C15	-0.0014 (3)	0.39104 (12)	0.12873 (15)	0.0670 (5)
H15A	0.1113	0.3852	0.1656	0.101*
H15B	0.0009	0.4265	0.0746	0.101*
H15C	-0.0425	0.3422	0.1023	0.101*
C16	0.5046 (3)	0.67409 (11)	0.71491 (15)	0.0659 (5)
H16A	0.6227	0.6615	0.7185	0.099*
H16B	0.4682	0.6610	0.7771	0.099*
H16C	0.4887	0.7281	0.7027	0.099*
C17	0.6693 (3)	0.42503 (13)	0.87605 (14)	0.0652 (5)
H17A	0.5884	0.4565	0.9028	0.098*
H17B	0.7613	0.4564	0.8616	0.098*
H17C	0.7126	0.3865	0.9241	0.098*
C18	0.3502 (4)	0.27476 (12)	0.54961 (16)	0.0889 (8)
H18A	0.3996	0.2927	0.4936	0.133*
H18B	0.2294	0.2826	0.5364	0.133*
H18C	0.3739	0.2210	0.5598	0.133*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0627 (3)	0.0666 (3)	0.0436 (2)	0.0013 (2)	-0.01048 (19)	0.00529 (18)
O1	0.1387 (15)	0.0401 (7)	0.0671 (9)	0.0214 (8)	-0.0336 (9)	-0.0107 (6)
O2	0.0932 (10)	0.0372 (6)	0.0533 (7)	-0.0014 (6)	-0.0234 (7)	0.0007 (5)
O3	0.0709 (8)	0.0411 (6)	0.0572 (7)	-0.0051 (6)	-0.0145 (6)	-0.0083 (5)
O4	0.0676 (8)	0.0606 (7)	0.0434 (6)	0.0106 (6)	-0.0122 (5)	0.0023 (5)
O5	0.0981 (10)	0.0408 (6)	0.0467 (7)	0.0036 (6)	-0.0128 (6)	0.0039 (5)
N1	0.0545 (8)	0.0427 (7)	0.0412 (7)	0.0128 (6)	-0.0015 (6)	-0.0003 (5)
N2	0.0476 (7)	0.0419 (7)	0.0382 (6)	-0.0027 (6)	-0.0019 (5)	-0.0003 (5)
C1	0.0397 (8)	0.0492 (9)	0.0399 (8)	0.0026 (6)	0.0047 (6)	0.0014 (6)
C2	0.0670 (11)	0.0424 (9)	0.0441 (8)	0.0112 (8)	-0.0048 (8)	-0.0045 (7)
C3	0.0467 (8)	0.0389 (8)	0.0374 (7)	0.0037 (6)	0.0023 (6)	-0.0004 (6)

C4	0.0481 (9)	0.0403 (8)	0.0377 (7)	-0.0016 (6)	-0.0005 (6)	0.0020 (6)
C5	0.0459 (8)	0.0370 (7)	0.0394 (8)	0.0006 (6)	0.0037 (6)	-0.0036 (6)
C6	0.0391 (7)	0.0413 (8)	0.0352 (7)	-0.0016 (6)	0.0033 (6)	-0.0020 (6)
C7	0.0412 (8)	0.0420 (8)	0.0417 (8)	-0.0030 (6)	0.0033 (6)	-0.0041 (6)
C8	0.0417 (8)	0.0520 (9)	0.0401 (8)	-0.0026 (7)	-0.0027 (6)	-0.0073 (6)
C9	0.0427 (8)	0.0536 (9)	0.0363 (8)	0.0034 (7)	-0.0002 (6)	0.0023 (6)
C10	0.0538 (9)	0.0413 (8)	0.0390 (8)	0.0000 (7)	0.0009 (7)	0.0013 (6)
C11	0.0485 (9)	0.0412 (8)	0.0343 (7)	-0.0030 (6)	-0.0001 (6)	-0.0009 (6)
C12	0.0780 (13)	0.0455 (9)	0.0548 (10)	0.0238 (9)	-0.0076 (9)	-0.0010 (8)
C13	0.1027 (17)	0.0430 (10)	0.0783 (14)	0.0035 (11)	-0.0015 (12)	0.0120 (9)
C14	0.0613 (11)	0.0513 (9)	0.0444 (9)	-0.0132 (8)	-0.0039 (8)	0.0005 (7)
C15	0.0834 (15)	0.0549 (11)	0.0608 (11)	-0.0030 (10)	0.0053 (10)	-0.0092 (9)
C16	0.0708 (12)	0.0523 (11)	0.0663 (12)	-0.0143 (9)	-0.0142 (10)	-0.0139 (9)
C17	0.0611 (11)	0.0810 (14)	0.0456 (10)	0.0059 (10)	-0.0158 (8)	-0.0016 (9)
C18	0.157 (2)	0.0432 (10)	0.0560 (11)	0.0000 (13)	-0.0162 (13)	-0.0037 (9)

Geometric parameters (Å, °)

S1—C1	1.6634 (16)	C9—C10	1.412 (2)
O1—C2	1.212 (2)	C10—C11	1.366 (2)
O2—C4	1.2145 (19)	C11—H11	0.9300
O3—C7	1.3542 (19)	C12—C13	1.499 (3)
O3—C16	1.424 (2)	C12—H12A	0.9700
O4—C9	1.3478 (19)	C12—H12B	0.9700
O4—C17	1.431 (2)	C13—H13A	0.9600
O5—C10	1.368 (2)	C13—H13B	0.9600
O5—C18	1.408 (2)	C13—H13C	0.9600
N1—C1	1.365 (2)	C14—C15	1.496 (3)
N1—C2	1.407 (2)	C14—H14A	0.9700
N1—C12	1.487 (2)	C14—H14B	0.9700
N2—C1	1.384 (2)	C15—H15A	0.9600
N2—C4	1.407 (2)	C15—H15B	0.9600
N2—C14	1.480 (2)	C15—H15C	0.9600
C2—C3	1.466 (2)	C16—H16A	0.9600
C3—C5	1.377 (2)	C16—H16B	0.9600
C3—C4	1.448 (2)	C16—H16C	0.9600
C5—C6	1.426 (2)	C17—H17A	0.9600
C5—H5	0.9300	C17—H17B	0.9600
C6—C11	1.415 (2)	C17—H17C	0.9600
C6—C7	1.421 (2)	C18—H18A	0.9600
C7—C8	1.389 (2)	C18—H18B	0.9600
C8—C9	1.380 (2)	C18—H18C	0.9600
C8—H8	0.9300		
C7—O3—C16	119.70 (14)	N1—C12—C13	111.77 (15)
C9—O4—C17	118.14 (15)	N1—C12—H12A	109.3
C10—O5—C18	116.88 (14)	C13—C12—H12A	109.3
C1—N1—C2	124.63 (13)	N1—C12—H12B	109.3

C1—N1—C12	119.84 (14)	C13—C12—H12B	109.3
C2—N1—C12	115.49 (14)	H12A—C12—H12B	107.9
C1—N2—C4	125.06 (13)	C12—C13—H13A	109.5
C1—N2—C14	119.37 (13)	C12—C13—H13B	109.5
C4—N2—C14	115.52 (13)	H13A—C13—H13B	109.5
N1—C1—N2	116.84 (14)	C12—C13—H13C	109.5
N1—C1—S1	121.91 (12)	H13A—C13—H13C	109.5
N2—C1—S1	121.25 (12)	H13B—C13—H13C	109.5
O1—C2—N1	118.63 (15)	N2—C14—C15	112.36 (15)
O1—C2—C3	123.93 (16)	N2—C14—H14A	109.1
N1—C2—C3	117.44 (14)	C15—C14—H14A	109.1
C5—C3—C4	127.53 (14)	N2—C14—H14B	109.1
C5—C3—C2	113.68 (14)	C15—C14—H14B	109.1
C4—C3—C2	118.78 (14)	H14A—C14—H14B	107.9
O2—C4—N2	117.66 (14)	C14—C15—H15A	109.5
O2—C4—C3	125.48 (15)	C14—C15—H15B	109.5
N2—C4—C3	116.85 (13)	H15A—C15—H15B	109.5
C3—C5—C6	138.77 (14)	C14—C15—H15C	109.5
C3—C5—H5	110.6	H15A—C15—H15C	109.5
C6—C5—H5	110.6	H15B—C15—H15C	109.5
C11—C6—C7	116.80 (14)	O3—C16—H16A	109.5
C11—C6—C5	126.91 (14)	O3—C16—H16B	109.5
C7—C6—C5	116.29 (14)	H16A—C16—H16B	109.5
O3—C7—C8	122.56 (14)	O3—C16—H16C	109.5
O3—C7—C6	116.54 (14)	H16A—C16—H16C	109.5
C8—C7—C6	120.90 (15)	H16B—C16—H16C	109.5
C9—C8—C7	120.58 (14)	O4—C17—H17A	109.5
C9—C8—H8	119.7	O4—C17—H17B	109.5
C7—C8—H8	119.7	H17A—C17—H17B	109.5
O4—C9—C8	124.58 (15)	O4—C17—H17C	109.5
O4—C9—C10	115.65 (15)	H17A—C17—H17C	109.5
C8—C9—C10	119.77 (14)	H17B—C17—H17C	109.5
C11—C10—O5	125.18 (15)	O5—C18—H18A	109.5
C11—C10—C9	119.67 (15)	O5—C18—H18B	109.5
O5—C10—C9	115.14 (14)	H18A—C18—H18B	109.5
C10—C11—C6	122.25 (14)	O5—C18—H18C	109.5
C10—C11—H11	118.9	H18A—C18—H18C	109.5
C6—C11—H11	118.9	H18B—C18—H18C	109.5
C2—N1—C1—N2	1.4 (2)	C3—C5—C6—C7	178.82 (18)
C12—N1—C1—N2	179.25 (15)	C16—O3—C7—C8	1.8 (3)
C2—N1—C1—S1	-178.14 (14)	C16—O3—C7—C6	-178.93 (16)
C12—N1—C1—S1	-0.3 (2)	C11—C6—C7—O3	179.37 (14)
C4—N2—C1—N1	-6.7 (2)	C5—C6—C7—O3	-1.0 (2)
C14—N2—C1—N1	175.91 (14)	C11—C6—C7—C8	-1.3 (2)
C4—N2—C1—S1	172.90 (12)	C5—C6—C7—C8	178.27 (14)
C14—N2—C1—S1	-4.5 (2)	O3—C7—C8—C9	179.83 (15)
C1—N1—C2—O1	-178.32 (19)	C6—C7—C8—C9	0.6 (2)

C12—N1—C2—O1	3.8 (3)	C17—O4—C9—C8	7.1 (3)
C1—N1—C2—C3	2.5 (3)	C17—O4—C9—C10	-173.28 (16)
C12—N1—C2—C3	-175.38 (16)	C7—C8—C9—O4	-179.43 (15)
O1—C2—C3—C5	-2.0 (3)	C7—C8—C9—C10	1.0 (2)
N1—C2—C3—C5	177.12 (15)	C18—O5—C10—C11	10.0 (3)
O1—C2—C3—C4	179.2 (2)	C18—O5—C10—C9	-171.0 (2)
N1—C2—C3—C4	-1.7 (3)	O4—C9—C10—C11	178.64 (15)
C1—N2—C4—O2	-172.69 (16)	C8—C9—C10—C11	-1.7 (2)
C14—N2—C4—O2	4.8 (2)	O4—C9—C10—O5	-0.4 (2)
C1—N2—C4—C3	7.3 (2)	C8—C9—C10—O5	179.17 (15)
C14—N2—C4—C3	-175.18 (14)	O5—C10—C11—C6	179.94 (16)
C5—C3—C4—O2	-1.4 (3)	C9—C10—C11—C6	1.0 (2)
C2—C3—C4—O2	177.22 (18)	C7—C6—C11—C10	0.5 (2)
C5—C3—C4—N2	178.57 (15)	C5—C6—C11—C10	-178.98 (16)
C2—C3—C4—N2	-2.8 (2)	C1—N1—C12—C13	-89.7 (2)
C4—C3—C5—C6	-1.4 (3)	C2—N1—C12—C13	88.3 (2)
C2—C3—C5—C6	179.87 (18)	C1—N2—C14—C15	90.12 (19)
C3—C5—C6—C11	-1.6 (3)	C4—N2—C14—C15	-87.55 (18)

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>
C13—H13C \cdots O2 ⁱ	0.96	2.58	3.455 (2)	152

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