

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

Methyl 2-((2Z,5Z)-2-[(E)-2-[1-(4-hydroxyphenyl)ethylidene]hydrazin-1-ylidene]-4-oxo-3-phenyl-1,3-thiazolidin-5-ylidene)-acetate

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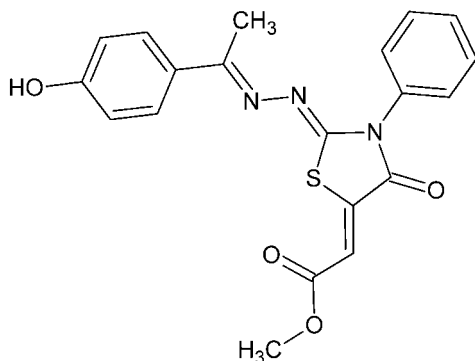
Received 29 July 2013; accepted 11 September 2013

Key indicators: single-crystal X-ray study; $T = 150$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.044; wR factor = 0.118; data-to-parameter ratio = 17.7.

In the title compound, $\text{C}_{20}\text{H}_{17}\text{N}_3\text{O}_4\text{S}$, all non-H atoms, except those of the phenyl ring, are approximately coplanar [maximum deviation = 0.2214 (1) Å], and the dihedral angle between their best plane and the benzene ring is 53.13 (1)°. A short intramolecular $\text{O}\cdots\text{S}$ contact of 2.838 (1) Å is formed between the ester carbonyl O atom and the S atom of the thiazolidine ring. In the crystal, molecules associated *via* $\text{O}-\text{H}\cdots\text{O}$, $\text{C}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{S}$ hydrogen bonds form layers parallel to (010), with only $\text{C}-\text{H}\cdots\text{O}$ -type short contacts between the molecules in adjacent layers.

Related literature

For the biological activity of 4-thiazolidinones, see: Dayam *et al.* (2006); Srivastava *et al.* (2005), Look *et al.* (1996), Barreca *et al.* (2001); Diurno *et al.* (1992).



Experimental

Crystal data

$\text{C}_{20}\text{H}_{17}\text{N}_3\text{O}_4\text{S}$
 $M_r = 395.42$
 Monoclinic, $P2_1/n$
 $a = 9.5049$ (9) Å
 $b = 20.656$ (2) Å
 $c = 10.1364$ (10) Å
 $\beta = 107.637$ (1)°

$V = 1896.6$ (3) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.20$ mm⁻¹
 $T = 150$ K
 $0.19 \times 0.11 \times 0.05$ mm

Data collection

Bruker SMART APEX CCD diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2013)
 $T_{\min} = 0.82$, $T_{\max} = 0.99$

16907 measured reflections
 4582 independent reflections
 3740 reflections with $i > 2\sigma(i)$
 $R_{\text{int}} = 0.039$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.118$
 $S = 1.06$
 4582 reflections
 259 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.34$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.44$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O4}-\text{H4O}\cdots\text{O1}^{\text{i}}$	0.84 (2)	1.96 (2)	2.7901 (16)	174.8 (19)
$\text{C8}-\text{H8}\cdots\text{S1}^{\text{ii}}$	0.95	2.82	3.7272 (17)	160
$\text{C10}-\text{H10}\cdots\text{O4}^{\text{iii}}$	0.95	2.52	3.452 (2)	167
$\text{C19}-\text{H19}\cdots\text{O1}^{\text{i}}$	0.95	2.47	3.200 (2)	133

Symmetry codes: (i) $x - \frac{3}{2}, -y + \frac{1}{2}, z - \frac{1}{2}$; (ii) $x + \frac{1}{2}, -y + \frac{1}{2}, z + \frac{1}{2}$; (iii) $-x + \frac{3}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$.

Data collection: APEX2 (Bruker, 2013); cell refinement: SAINT (Bruker, 2013); data reduction: SAINT; program(s) used to solve structure: SHELXT (Sheldrick, 2008); program(s) used to refine structure: SHELXL2013 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 2012); software used to prepare material for publication: WinGX (Farrugia, 2012) and PLATON (Spek, 2009).

Manchester Metropolitan University, Tulane University and Erciyes University are gratefully acknowledged for supporting this study.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GK2589).

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supporting information

Acta Cryst. (2013). E69, o1553–o1554 [doi:10.1107/S1600536813025270]

Methyl 2-((2*Z*,5*Z*)-2-{(*E*)-2-[1-(4-hydroxyphenyl)ethylidene]hydrazin-1-ylidene}-4-oxo-3-phenyl-1,3-thiazolidin-5-ylidene)acetate

Shaaban K. Mohamed, Joel T. Mague, Mehmet Akkurt, Alaa A. Hassan and Mustafa R. Albayati

S1. Comment

Thiazolidinone scaffold compounds have received much attention from organic and medicinal chemists due to their therapeutic diversity coupled with their commercial viability. Recently, 4-thiazolidinones have exhibited many interesting bio-activity profiles such as anti-cancer (Dayam *et al.*, 2006) and anti-mycobacterial agents (Srivastava *et al.*, 2005), COX-1 inhibitors (Look *et al.* 1996), non-nucleoside inhibitors of HIV-RT (Barreca *et al.*, 2001) and anti-histaminic agents (Diurno *et al.*, 1992). In view of these properties the title compound has been synthesized among a series of other 4-thiazolidinones to investigate the relationship between their crystal structures and their antibacterial activity.

In the title compound (Fig. 1), all non-H atoms, except the phenyl group (C7–C12), are approximately coplanar, with the maximum deviations of -0.2214 (1) Å for C6, -0.2097 (1) Å for C14, 0.1651 (1) Å for O2 and -0.1009 (1) Å for O3, and the benzene ring (C7–C12) makes a dihedral angle of 53.13 (1)° with this plane. Molecular conformation is stabilized by a short intramolecular O··S contact of 2.838 (1) Å.

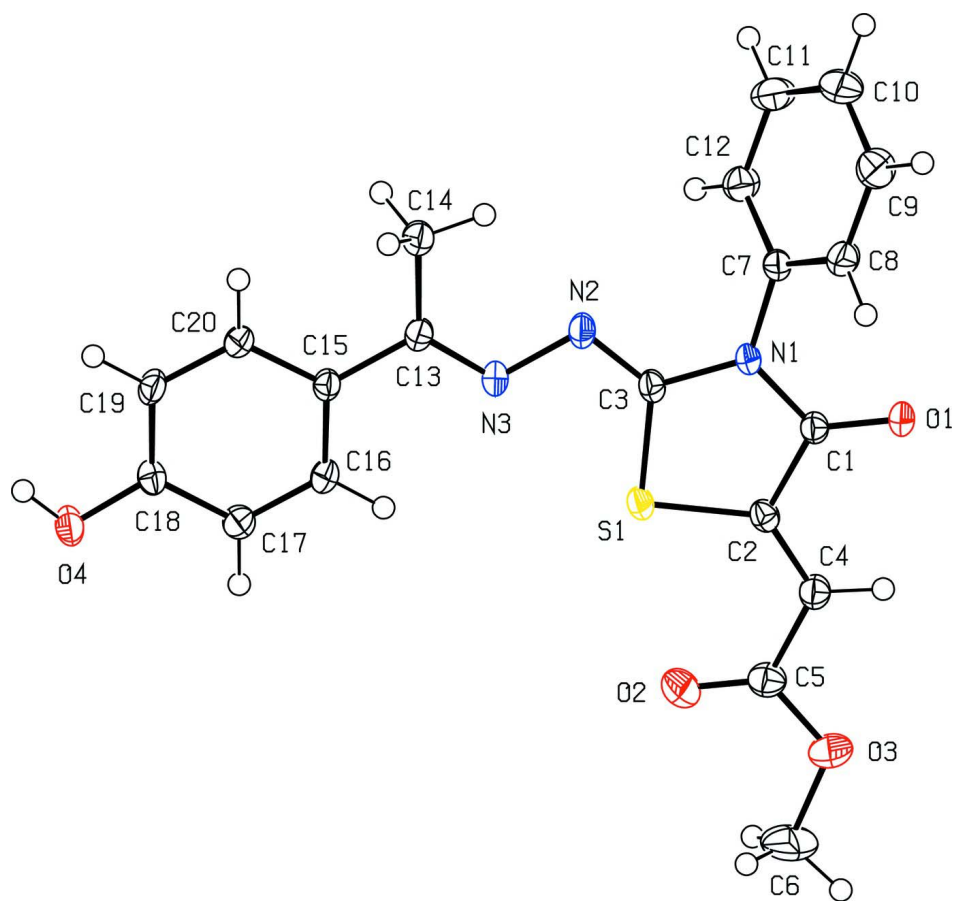
The title compound crystallizes in a layer structure with the layers parallel to the (010) plane (Fig. 2). Molecules within the layers are associated *via* O—H··O, C—H··O and C—H··S hydrogen bonding (Table 1, Fig. 2). One of the C—H··O contacts (C10—H10··O4) in Table 1 is between the layers. The interlayer regions are occupied by the *N*-phenyl and ester groups between which there are no significant interactions.

S2. Experimental

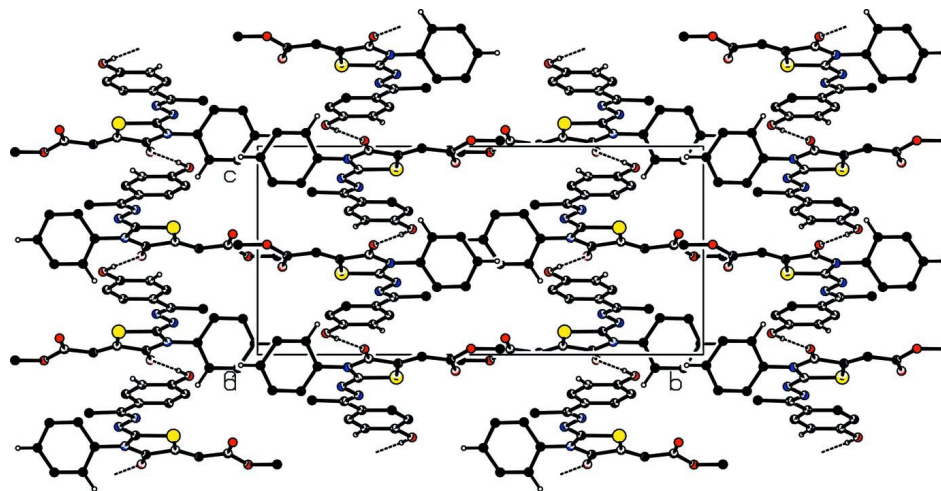
A mixture of 283 mg (1 mmol) (2*Z*)-2-[1-(4-methylphenyl)ethylidene]-*N*-phenylhydrazinecarbothioamide and 142 mg (1 mmol) dimethyl but-2-ynedioate in 50 ml of ethanol was refluxed and monitored by TLC until completion of the reaction. The excess solvent was evaporated under *vacuum* and the solid obtained was recrystallized from ethanol to afford clear yellow plates (*M.p.* 541–543 K) of X-ray quality.

S3. Refinement

The hydroxyl H atom was found from a difference Fourier map [O4—H4O = 0.84 (2) Å] and refined freely. H atoms bonded to C were placed in geometrically idealized positions and constrained to ride on their parent atoms C—H = 0.95 Å (aromatic H) and 0.98 Å (methyl H), with $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{iso}}(\text{C})$ for methyl H atoms and $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{iso}}(\text{C})$ for other H atoms.

**Figure 1**

Perspective view of the title molecule with 50% probability displacement ellipsoids.

**Figure 2**

View of the crystal packing down the *a*-axis showing the layer structure and O-H...O hydrogen bonding interactions as dotted lines.

Methyl 2-((2Z,5Z)-2-((E)-2-[1-(4-hydroxyphenyl)ethylidene]hydrazin-1-ylidene)-4-oxo-3-phenyl-1,3-thiazolidin-5-ylidene)acetate

Crystal data

C₂₀H₁₇N₃O₄S

M_r = 395.42

Monoclinic, *P*2₁/*n*

Hall symbol: -*P* 2₁*y**n*

a = 9.5049 (9) Å

b = 20.656 (2) Å

c = 10.1364 (10) Å

β = 107.637 (1)°

V = 1896.6 (3) Å³

Z = 4

F(000) = 824

D_x = 1.385 Mg m⁻³

Mo *K*α radiation, λ = 0.71073 Å

Cell parameters from 8755 reflections

θ = 2.3–28.6°

μ = 0.20 mm⁻¹

T = 150 K

Plate, clear yellow

0.19 × 0.11 × 0.05 mm

Data collection

Bruker SMART APEX CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 8.3660 pixels mm⁻¹

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2013)

T_{min} = 0.82, *T_{max}* = 0.99

16907 measured reflections

4582 independent reflections

3740 reflections with *i* > 2σ(*i*)

R_{int} = 0.039

θ_{\max} = 28.7°, θ_{\min} = 2.0°

h = -12→12

k = -27→27

l = -13→13

Refinement

Refinement on *F*²

Least-squares matrix: full

R[*F*² > 2σ(*F*²)] = 0.044

wR(*F*²) = 0.118

S = 1.06

4582 reflections

259 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

H atoms treated by a mixture of independent
and constrained refinement

W = 1/[Σ²(*FO*²) + (0.0643*P*)² + 0.3949*P*]

WHERE *P* = (*FO*² + 2*FC*²)/3

(Δ/σ)_{max} = 0.001

Δρ_{max} = 0.34 e Å⁻³

Δρ_{min} = -0.44 e Å⁻³

Special details

Experimental. The diffraction data were collected in three sets of 606 frames (0.3° width in ω) at φ = 0, 120 and 240°. A scan time of 40 sec/frame was used.

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 > σ(*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 (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U_{iso}</i> */ <i>U_{eq}</i>
S1	1.01222 (4)	0.18869 (2)	0.39033 (4)	0.0203 (1)

O1	1.40358 (11)	0.26023 (5)	0.52637 (11)	0.0240 (3)
O2	1.09079 (13)	0.05575 (6)	0.42140 (13)	0.0347 (4)
O3	1.32263 (14)	0.02111 (6)	0.52834 (14)	0.0378 (4)
O4	0.14035 (12)	0.15348 (6)	0.10230 (13)	0.0285 (3)
N1	1.16412 (13)	0.29748 (6)	0.43935 (12)	0.0189 (3)
N2	0.90771 (13)	0.31123 (6)	0.34670 (13)	0.0222 (4)
N3	0.77907 (13)	0.27335 (6)	0.30473 (13)	0.0219 (4)
C1	1.27118 (16)	0.25046 (7)	0.47990 (15)	0.0190 (4)
C2	1.20296 (16)	0.18459 (7)	0.45802 (14)	0.0193 (4)
C3	1.01955 (15)	0.27395 (7)	0.38850 (14)	0.0190 (4)
C4	1.28646 (17)	0.13149 (7)	0.49043 (16)	0.0229 (4)
C5	1.22088 (18)	0.06672 (8)	0.47422 (16)	0.0260 (5)
C6	1.2661 (3)	-0.04389 (9)	0.5273 (3)	0.0553 (8)
C7	1.20015 (15)	0.36539 (7)	0.44261 (16)	0.0206 (4)
C8	1.29110 (17)	0.39165 (8)	0.56415 (17)	0.0275 (5)
C9	1.3342 (2)	0.45594 (9)	0.5655 (2)	0.0368 (5)
C10	1.2839 (2)	0.49351 (8)	0.4477 (2)	0.0384 (6)
C11	1.1899 (2)	0.46687 (8)	0.32791 (19)	0.0347 (5)
C12	1.14814 (18)	0.40248 (8)	0.32417 (16)	0.0272 (5)
C13	0.65717 (16)	0.30559 (7)	0.27572 (15)	0.0196 (4)
C14	0.64599 (17)	0.37782 (8)	0.28058 (17)	0.0260 (5)
C15	0.52047 (15)	0.26621 (7)	0.23315 (15)	0.0187 (4)
C16	0.52602 (16)	0.19872 (7)	0.23265 (17)	0.0241 (4)
C17	0.39928 (17)	0.16202 (8)	0.18891 (18)	0.0267 (5)
C18	0.26171 (16)	0.19204 (7)	0.14442 (15)	0.0207 (4)
C19	0.25339 (16)	0.25898 (7)	0.14520 (16)	0.0227 (4)
C20	0.38175 (16)	0.29540 (7)	0.18934 (16)	0.0221 (4)
H4	1.39070	0.13580	0.52480	0.0270*
H4O	0.066 (2)	0.1774 (11)	0.077 (2)	0.046 (6)*
H6A	1.19470	-0.04490	0.57960	0.0830*
H6B	1.34780	-0.07350	0.56990	0.0830*
H6C	1.21740	-0.05730	0.43150	0.0830*
H8	1.32360	0.36600	0.64560	0.0330*
H9	1.39860	0.47420	0.64770	0.0440*
H10	1.31370	0.53750	0.44900	0.0460*
H11	1.15380	0.49300	0.24760	0.0420*
H12	1.08480	0.38410	0.24160	0.0330*
H14A	0.74530	0.39660	0.31100	0.0390*
H14B	0.59250	0.38990	0.34570	0.0390*
H14C	0.59280	0.39420	0.18820	0.0390*
H16	0.61910	0.17760	0.26310	0.0290*
H17	0.40570	0.11610	0.18910	0.0320*
H19	0.16000	0.27990	0.11560	0.0270*
H20	0.37500	0.34130	0.18970	0.0260*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0149 (2)	0.0209 (2)	0.0233 (2)	-0.0034 (1)	0.0030 (1)	-0.0019 (1)
O1	0.0134 (5)	0.0242 (6)	0.0308 (6)	-0.0009 (4)	0.0011 (4)	0.0019 (4)
O2	0.0316 (7)	0.0275 (6)	0.0427 (7)	-0.0071 (5)	0.0078 (5)	-0.0030 (5)
O3	0.0387 (7)	0.0198 (6)	0.0525 (8)	0.0051 (5)	0.0104 (6)	0.0035 (5)
O4	0.0159 (5)	0.0239 (6)	0.0441 (7)	-0.0016 (5)	0.0067 (5)	-0.0074 (5)
N1	0.0131 (6)	0.0191 (6)	0.0222 (6)	-0.0025 (5)	0.0017 (5)	-0.0002 (5)
N2	0.0141 (6)	0.0247 (7)	0.0252 (7)	-0.0007 (5)	0.0022 (5)	-0.0009 (5)
N3	0.0137 (6)	0.0252 (7)	0.0250 (7)	-0.0005 (5)	0.0030 (5)	0.0008 (5)
C1	0.0158 (7)	0.0215 (7)	0.0187 (7)	-0.0012 (5)	0.0036 (5)	0.0005 (5)
C2	0.0180 (7)	0.0220 (7)	0.0172 (7)	-0.0034 (6)	0.0043 (5)	-0.0011 (5)
C3	0.0147 (7)	0.0220 (7)	0.0186 (7)	-0.0026 (5)	0.0025 (5)	-0.0012 (5)
C4	0.0193 (7)	0.0236 (8)	0.0242 (7)	0.0000 (6)	0.0044 (6)	-0.0009 (6)
C5	0.0294 (8)	0.0222 (8)	0.0273 (8)	0.0005 (6)	0.0101 (7)	-0.0014 (6)
C6	0.0658 (15)	0.0195 (9)	0.0817 (17)	0.0013 (9)	0.0238 (13)	0.0051 (10)
C7	0.0153 (6)	0.0181 (7)	0.0273 (8)	-0.0001 (5)	0.0050 (6)	-0.0002 (6)
C8	0.0230 (8)	0.0242 (8)	0.0296 (8)	0.0001 (6)	-0.0005 (6)	0.0014 (6)
C9	0.0334 (9)	0.0251 (9)	0.0423 (10)	-0.0044 (7)	-0.0031 (8)	-0.0061 (7)
C10	0.0401 (10)	0.0182 (8)	0.0524 (12)	-0.0038 (7)	0.0071 (9)	0.0001 (7)
C11	0.0408 (10)	0.0246 (8)	0.0354 (9)	0.0018 (7)	0.0066 (8)	0.0074 (7)
C12	0.0283 (8)	0.0242 (8)	0.0251 (8)	-0.0008 (6)	0.0023 (6)	-0.0004 (6)
C13	0.0157 (7)	0.0239 (8)	0.0185 (7)	0.0017 (6)	0.0041 (5)	0.0016 (5)
C14	0.0198 (7)	0.0224 (8)	0.0344 (9)	0.0006 (6)	0.0060 (6)	-0.0006 (6)
C15	0.0143 (7)	0.0228 (7)	0.0190 (7)	0.0009 (5)	0.0051 (5)	0.0017 (5)
C16	0.0151 (7)	0.0232 (8)	0.0334 (8)	0.0042 (6)	0.0065 (6)	0.0029 (6)
C17	0.0206 (8)	0.0191 (7)	0.0409 (9)	0.0027 (6)	0.0099 (7)	-0.0003 (6)
C18	0.0162 (7)	0.0234 (8)	0.0230 (7)	-0.0013 (6)	0.0069 (6)	-0.0028 (6)
C19	0.0147 (7)	0.0249 (8)	0.0270 (8)	0.0044 (6)	0.0041 (6)	0.0003 (6)
C20	0.0171 (7)	0.0195 (7)	0.0289 (8)	0.0032 (6)	0.0059 (6)	0.0015 (6)

Geometric parameters (Å, °)

S1—C2	1.7352 (16)	C13—C15	1.482 (2)
S1—C3	1.7628 (15)	C15—C20	1.394 (2)
O1—C1	1.2195 (19)	C15—C16	1.395 (2)
O2—C5	1.211 (2)	C16—C17	1.378 (2)
O3—C5	1.343 (2)	C17—C18	1.393 (2)
O3—C6	1.445 (2)	C18—C19	1.385 (2)
O4—C18	1.360 (2)	C19—C20	1.387 (2)
O4—H4O	0.84 (2)	C4—H4	0.9500
N1—C1	1.376 (2)	C6—H6A	0.9800
N1—C7	1.4420 (19)	C6—H6B	0.9800
N1—C3	1.4003 (19)	C6—H6C	0.9800
N2—N3	1.4046 (18)	C8—H8	0.9500
N2—C3	1.2766 (19)	C9—H9	0.9500
N3—C13	1.291 (2)	C10—H10	0.9500

C1—C2	1.495 (2)	C11—H11	0.9500
C2—C4	1.335 (2)	C12—H12	0.9500
C4—C5	1.464 (2)	C14—H14A	0.9800
C7—C12	1.383 (2)	C14—H14B	0.9800
C7—C8	1.384 (2)	C14—H14C	0.9800
C8—C9	1.389 (3)	C16—H16	0.9500
C9—C10	1.382 (3)	C17—H17	0.9500
C10—C11	1.385 (3)	C19—H19	0.9500
C11—C12	1.385 (2)	C20—H20	0.9500
C13—C14	1.498 (2)		
C2—S1—C3	90.75 (7)	O4—C18—C17	117.69 (13)
C5—O3—C6	115.39 (17)	O4—C18—C19	122.83 (14)
C18—O4—H4O	107.9 (15)	C17—C18—C19	119.48 (14)
C1—N1—C3	114.79 (12)	C18—C19—C20	119.81 (14)
C3—N1—C7	123.08 (12)	C15—C20—C19	121.51 (13)
C1—N1—C7	122.05 (13)	C2—C4—H4	119.00
N3—N2—C3	108.99 (12)	C5—C4—H4	119.00
N2—N3—C13	114.87 (12)	O3—C6—H6A	109.00
O1—C1—C2	123.95 (14)	O3—C6—H6B	109.00
N1—C1—C2	110.48 (13)	O3—C6—H6C	109.00
O1—C1—N1	125.57 (14)	H6A—C6—H6B	110.00
S1—C2—C4	127.54 (12)	H6A—C6—H6C	109.00
C1—C2—C4	120.83 (14)	H6B—C6—H6C	109.00
S1—C2—C1	111.63 (11)	C7—C8—H8	120.00
S1—C3—N2	125.07 (12)	C9—C8—H8	120.00
N1—C3—N2	122.57 (13)	C8—C9—H9	120.00
S1—C3—N1	112.36 (10)	C10—C9—H9	120.00
C2—C4—C5	121.46 (15)	C9—C10—H10	120.00
O2—C5—O3	124.20 (16)	C11—C10—H10	120.00
O2—C5—C4	124.31 (15)	C10—C11—H11	120.00
O3—C5—C4	111.46 (14)	C12—C11—H11	120.00
N1—C7—C12	119.94 (14)	C7—C12—H12	121.00
C8—C7—C12	121.22 (14)	C11—C12—H12	121.00
N1—C7—C8	118.80 (13)	C13—C14—H14A	109.00
C7—C8—C9	119.12 (15)	C13—C14—H14B	109.00
C8—C9—C10	120.31 (17)	C13—C14—H14C	109.00
C9—C10—C11	119.78 (16)	H14A—C14—H14B	110.00
C10—C11—C12	120.59 (16)	H14A—C14—H14C	109.00
C7—C12—C11	118.95 (15)	H14B—C14—H14C	109.00
N3—C13—C15	115.50 (13)	C15—C16—H16	119.00
C14—C13—C15	119.45 (13)	C17—C16—H16	119.00
N3—C13—C14	125.05 (14)	C16—C17—H17	120.00
C13—C15—C20	121.09 (13)	C18—C17—H17	120.00
C16—C15—C20	117.65 (14)	C18—C19—H19	120.00
C13—C15—C16	121.25 (14)	C20—C19—H19	120.00
C15—C16—C17	121.37 (15)	C15—C20—H20	119.00
C16—C17—C18	120.18 (15)	C19—C20—H20	119.00

C3—S1—C2—C1	0.49 (11)	S1—C2—C4—C5	-1.6 (2)
C3—S1—C2—C4	179.93 (14)	C1—C2—C4—C5	177.79 (14)
C2—S1—C3—N1	-0.41 (11)	C2—C4—C5—O2	6.1 (3)
C2—S1—C3—N2	179.83 (13)	C2—C4—C5—O3	-172.43 (14)
C6—O3—C5—O2	-3.4 (3)	N1—C7—C8—C9	175.77 (15)
C6—O3—C5—C4	175.15 (17)	C12—C7—C8—C9	-2.0 (3)
C3—N1—C1—O1	179.63 (14)	N1—C7—C12—C11	-176.95 (15)
C3—N1—C1—C2	0.14 (17)	C8—C7—C12—C11	0.8 (3)
C7—N1—C1—O1	2.8 (2)	C7—C8—C9—C10	1.5 (3)
C7—N1—C1—C2	-176.72 (12)	C8—C9—C10—C11	0.1 (3)
C1—N1—C3—S1	0.22 (15)	C9—C10—C11—C12	-1.3 (3)
C1—N1—C3—N2	179.98 (14)	C10—C11—C12—C7	0.9 (3)
C7—N1—C3—S1	177.05 (11)	N3—C13—C15—C16	3.4 (2)
C7—N1—C3—N2	-3.2 (2)	N3—C13—C15—C20	-175.24 (14)
C1—N1—C7—C8	-52.6 (2)	C14—C13—C15—C16	-177.37 (14)
C1—N1—C7—C12	125.14 (16)	C14—C13—C15—C20	4.0 (2)
C3—N1—C7—C8	130.78 (15)	C13—C15—C16—C17	-177.90 (15)
C3—N1—C7—C12	-51.5 (2)	C20—C15—C16—C17	0.8 (2)
C3—N2—N3—C13	172.11 (13)	C13—C15—C20—C19	177.96 (14)
N3—N2—C3—S1	0.83 (17)	C16—C15—C20—C19	-0.7 (2)
N3—N2—C3—N1	-178.90 (12)	C15—C16—C17—C18	-0.3 (3)
N2—N3—C13—C14	1.6 (2)	C16—C17—C18—O4	179.96 (15)
N2—N3—C13—C15	-179.18 (12)	C16—C17—C18—C19	-0.3 (2)
O1—C1—C2—S1	-179.96 (13)	O4—C18—C19—C20	-179.92 (14)
O1—C1—C2—C4	0.6 (2)	C17—C18—C19—C20	0.4 (2)
N1—C1—C2—S1	-0.45 (15)	C18—C19—C20—C15	0.1 (2)
N1—C1—C2—C4	-179.95 (15)		

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O4—H4O \cdots O1 ⁱ	0.84 (2)	1.96 (2)	2.7901 (16)	174.8 (19)
C8—H8 \cdots S1 ⁱⁱ	0.95	2.82	3.7272 (17)	160
C10—H10 \cdots O4 ⁱⁱⁱ	0.95	2.52	3.452 (2)	167
C14—H14A \cdots N2	0.98	2.30	2.742 (2)	106
C19—H19 \cdots O1 ⁱ	0.95	2.47	3.200 (2)	133

Symmetry codes: (i) $x-3/2, -y+1/2, z-1/2$; (ii) $x+1/2, -y+1/2, z+1/2$; (iii) $-x+3/2, y+1/2, -z+1/2$.