

Crystal structure of *N'*-[(1*E*)-1-(6-methyl-2,4-dioxo-3,4-dihydro-2*H*-pyran-3-ylidene)ethyl]benzenesulfonohydrazide

Jonnie N. Asegbeloyin,^a Oguejiofo T. Ujam,^{a*} Chizoba M. Ngige,^b Valentine I. Onwukeme^b and Tania Grouso^c

^aDepartment of Pure and Industrial Chemistry, University of Nigeria, Nsukka, Enugu State, Nigeria, ^bDepartment of Pure and Industrial Chemistry, Nnamdi Azikiwe University, PMB 5025, Awka, Anambra State, Nigeria, and ^cSchool of Chemical Sciences, the University of Auckland Private Bag 92019, Auckland 1142, New Zealand. *Correspondence e-mail: oguejiofo.ujam@unn.edu.ng

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In the title compound, C₁₄H₁₄N₂O₅S, the molecule exists in the enamine (C=C–NH) tautomeric form. The hydrazone fragment derived from the 3-acetyl-4-hydroxy-6-methyl-2*H*-pyran-2-one moiety is approximately planar, with a maximum deviation of 0.1291 (11) Å for the N atom bound to the S atom of the benzenesulfonohydrazide group. The latter adopts a *gauche* conformation relative to the hydrazone N–N bond, with an N–N–S angle of 113.54 (10)°. There is an intramolecular N–H···O=C hydrogen bond that stabilizes the tautomeric form. In the crystal, molecules are linked by N–H···O=C hydrogen bonds into chains extending parallel to [100].

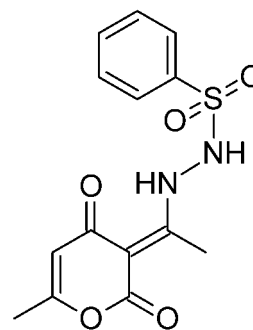
Keywords: crystal structure; hydrazone; benzenesulfonohydrazide; enamine tautomeric form; hydrogen bonds.

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1. Related literature

3-Acetyl-4-hydroxy-6-methyl-2*H*-pyran-2-one and its derivatives have received attention due to their coordination chemistry, pharmaceutical significance and biological activities (Battaini *et al.*, 2000; Puerta & Cohen, 2003; Rao & Narasaiah, 2003; Zucolotto Chalaça *et al.*, 2002; Fouad *et al.*, 2010; Kubaisi & Ismail, 1994; Rao *et al.*, 1985; Deshmukh *et al.*, 2010*a,b*; Munde *et al.*, 2009, 2010; Faidallah *et al.*, 2011; Jadhav *et al.*, 2010). 3-Acetyl-4-hydroxy-6-methyl-2*H*-pyran-2-one is also well-noted for its fungicidal (Rao *et al.*, 1978), herbicidal and antimicrobial activities (Zucolotto Chalaça *et al.*, 2002). The title compound is a new hydrazone prepared as part of an on-going research to study the ligating ability and anti-

microbial properties of 3-acetyl-4-hydroxy-6-methyl-2*H*-pyran-2-one hydrazones and their derivatives. For the crystal structure of a related thiosemicarbazone, see: Vrdoljak *et al.* (2008). For a benzenesulfonohydrazide derivative of a similar tautomeric enamine form, see: Ukwueze *et al.* (2014).



2. Experimental

2.1. Crystal data

C ₁₄ H ₁₄ N ₂ O ₅ S	<i>V</i> = 1453.51 (12) Å ³
<i>M_r</i> = 322.33	<i>Z</i> = 4
Monoclinic, <i>P</i> 2 ₁ / <i>n</i>	Mo <i>K</i> α radiation
<i>a</i> = 7.4797 (4) Å	<i>μ</i> = 0.25 mm ⁻¹
<i>b</i> = 15.2458 (7) Å	<i>T</i> = 99 K
<i>c</i> = 12.7820 (6) Å	0.30 × 0.30 × 0.12 mm
<i>β</i> = 94.282 (3)°	

2.2. Data collection

Bruker APEXII CCD diffractometer	17170 measured reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 2003)	3484 independent reflections
<i>T</i> _{min} = 0.659, <i>T</i> _{max} = 0.746	2817 reflections with <i>I</i> > 2σ(<i>I</i>)
	<i>R</i> _{int} = 0.056

2.3. Refinement

<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)] = 0.043	H atoms treated by a mixture of independent and constrained refinement
<i>wR</i> (<i>F</i> ²) = 0.126	<i>Δρ</i> _{max} = 0.35 e Å ⁻³
<i>S</i> = 1.05	<i>Δρ</i> _{min} = -0.42 e Å ⁻³
3484 reflections	
209 parameters	

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>
N1–HN1···O4 ⁱ	0.89 (2)	1.89 (2)	2.7837 (19)	175 (2)
N2–HN2···O3	0.90 (2)	1.74 (2)	2.5194 (18)	144 (2)

Symmetry code: (i) *x* – 1, *y*, *z*.

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL2013* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012) and *Mercury* (Macrae *et al.*, 2006); software used to prepare material for publication: *WinGX* (Farrugia, 2012).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: WM5072).

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Crystal structure of *N'*-[(1*E*)-1-(6-methyl-2,4-dioxo-3,4-dihydro-2*H*-pyran-3-ylidene)ethyl]benzenesulfonohydrazide

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S1. Synthesis and Crystallisation

A solution of 3-acetyl-4-hydroxy-6-methyl-(2*H*)-pyran-2-one [168 mg, 1 mmol] in methanol (10 ml) was mixed with a solution of benzenesulfonohydrazide [172 mg, 1 mmol] in methanol (10 ml). The mixture was refluxed for 3 h, and the resulting solution cooled to obtain a white precipitate. The product was filtered, dried and recrystallized from methanol. Crystals suitable for X-ray crystallographic analysis were obtained by slow evaporation of a methanolic solution at room temperature for 48 h.

S2. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. Hydrogen atoms were placed in calculated positions with C—H = 0.95 - 0.98 Å and refined using a riding model with displacement parameters $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$ for aromatic and methylene groups and $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{C})$ for methyl groups. The two N—H hydrogen atoms were located in a difference map and were refined freely.

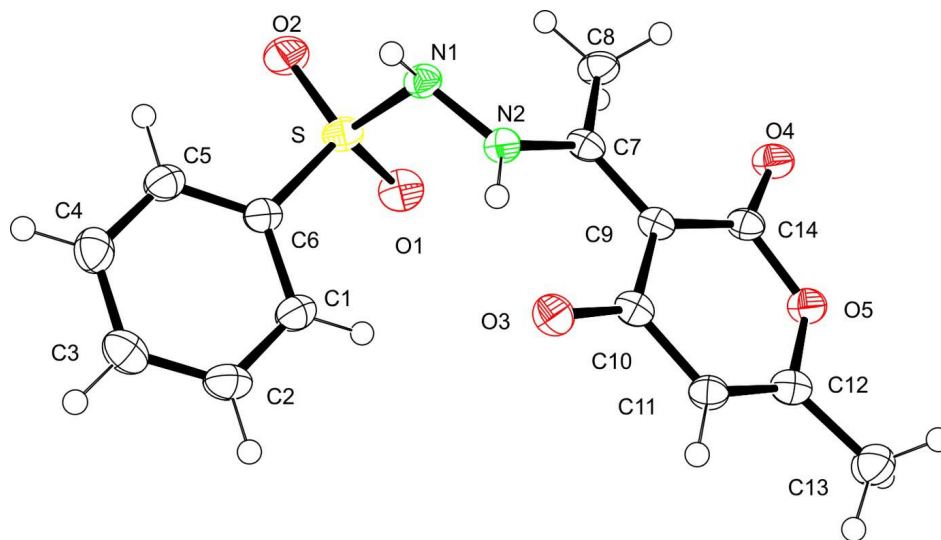
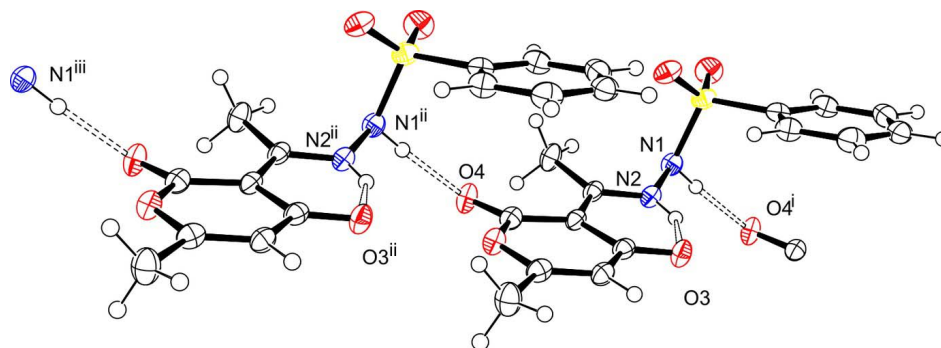
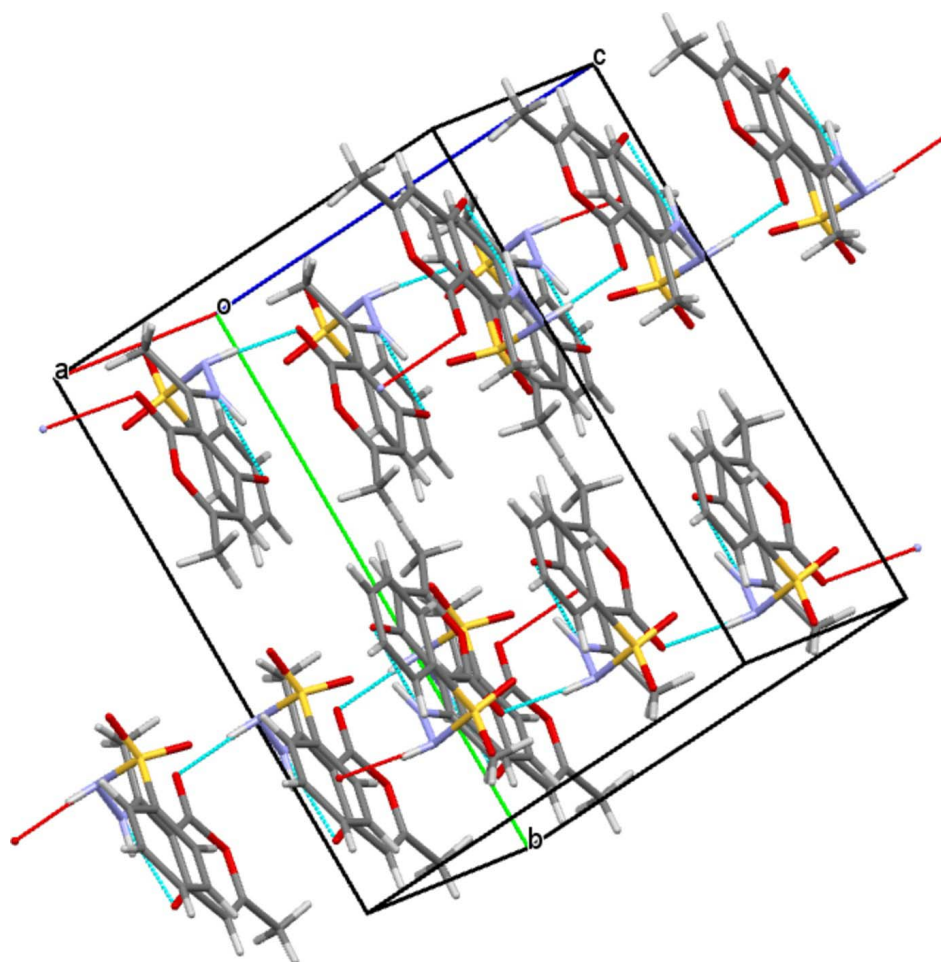


Figure 1

The molecular structure and atom numbering of the title compound with displacement ellipsoids drawn at the 50% probability level for non-H atoms.

**Figure 2**

Intramolecular N2-Hn2...O3 and intermolecular N1-Hn1...O4 (dotted lines) hydrogen bonding interactions in the title compound. [Symmetry codes: i) 1+x, y, z; ii) -1+x, y, z; iii) -2+x, y, z.]

**Figure 3**

The packing diagram of the title compound showing intra- and intermolecular N—H...O=C hydrogen bonds as dotted lines.

N'*-[*(1E)*-1-(6-Methyl-2,4-dioxo-3,4-dihydro-2*H*-pyran-3-ylidene)ethyl]benzenesulfonohydrazideCrystal data*C₁₄H₁₄N₂O₅S $M_r = 322.33$ Monoclinic, $P2_1/n$ Hall symbol: $-P\ 2_1n$ $a = 7.4797\ (4)\ \text{\AA}$ $b = 15.2458\ (7)\ \text{\AA}$ $c = 12.7820\ (6)\ \text{\AA}$ $\beta = 94.282\ (3)^\circ$ $V = 1453.51\ (12)\ \text{\AA}^3$ $Z = 4$ $F(000) = 672$ $D_x = 1.473\ \text{Mg m}^{-3}$

Melting point: 473 K

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

Cell parameters from 5140 reflections

 $\theta = 2.7\text{--}28.0^\circ$ $\mu = 0.25\ \text{mm}^{-1}$ $T = 99\ \text{K}$

Needle, colourless

 $0.30 \times 0.30 \times 0.12\ \text{mm}$ *Data collection*

Bruker APEXII CCD

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

 φ and ω scans

Absorption correction: multi-scan

(SADABS; Sheldrick, 2003)

 $T_{\min} = 0.659$, $T_{\max} = 0.746$

17170 measured reflections

3484 independent reflections

2817 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.056$ $\theta_{\max} = 28.0^\circ$, $\theta_{\min} = 2.1^\circ$ $h = -9 \rightarrow 9$ $k = -20 \rightarrow 19$ $l = -16 \rightarrow 16$ *Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.043$ $wR(F^2) = 0.126$ $S = 1.05$

3484 reflections

209 parameters

0 restraints

Hydrogen site location: mixed

H atoms treated by a mixture of independent and constrained refinement

 $w = 1/[\sigma^2(F_o^2) + (0.0681P)^2 + 0.4046P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.35\ \text{e \AA}^{-3}$ $\Delta\rho_{\min} = -0.42\ \text{e \AA}^{-3}$ *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.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	-0.1162 (3)	0.17867 (12)	-0.16895 (13)	0.0281 (4)
H1	0.0092	0.1698	-0.1554	0.034*
C2	-0.2300 (3)	0.10824 (12)	-0.19238 (14)	0.0326 (4)
H2	-0.1824	0.0505	-0.1945	0.039*
C3	-0.4125 (3)	0.12162 (13)	-0.21263 (13)	0.0321 (4)
H3	-0.4890	0.0732	-0.2298	0.038*
C4	-0.4844 (3)	0.20534 (13)	-0.20807 (13)	0.0322 (4)
H4	-0.6099	0.2139	-0.2211	0.039*
C5	-0.3730 (3)	0.27633 (12)	-0.18449 (12)	0.0280 (4)

H5	-0.4214	0.3338	-0.1812	0.034*
C6	-0.1896 (2)	0.26269 (11)	-0.16563 (12)	0.0237 (4)
C7	0.2970 (2)	0.31837 (11)	0.04228 (12)	0.0216 (3)
C8	0.3518 (2)	0.41187 (11)	0.02935 (15)	0.0289 (4)
H8A	0.2446	0.4487	0.0183	0.043*
H8B	0.4239	0.4169	-0.0313	0.043*
H8C	0.4228	0.4313	0.0926	0.043*
C9	0.4199 (2)	0.24839 (10)	0.06583 (11)	0.0206 (3)
C10	0.3574 (2)	0.15821 (10)	0.06591 (12)	0.0227 (3)
C11	0.4934 (2)	0.09113 (11)	0.08080 (12)	0.0231 (3)
H11	0.4581	0.0313	0.0802	0.028*
C12	0.6670 (2)	0.11099 (11)	0.09535 (12)	0.0256 (4)
C13	0.8158 (2)	0.04793 (13)	0.11667 (16)	0.0351 (4)
H13A	0.8639	0.0538	0.1898	0.053*
H13B	0.9108	0.0602	0.0699	0.053*
H13C	0.7713	-0.0119	0.1045	0.053*
C14	0.6075 (2)	0.26625 (11)	0.08402 (12)	0.0231 (3)
N1	-0.0028 (2)	0.35893 (9)	-0.00836 (11)	0.0236 (3)
N2	0.12527 (19)	0.29758 (9)	0.03004 (10)	0.0223 (3)
O1	0.12058 (18)	0.33764 (9)	-0.18100 (10)	0.0336 (3)
O2	-0.14865 (19)	0.43127 (8)	-0.16350 (10)	0.0361 (3)
O3	0.19542 (17)	0.13680 (8)	0.05307 (10)	0.0289 (3)
O4	0.68037 (16)	0.33852 (8)	0.09030 (10)	0.0293 (3)
O5	0.72498 (16)	0.19616 (8)	0.09491 (9)	0.0260 (3)
S	-0.04830 (6)	0.35372 (3)	-0.13822 (3)	0.02574 (15)
HN1	-0.102 (3)	0.3549 (13)	0.0260 (16)	0.033 (5)*
HN2	0.100 (3)	0.2408 (16)	0.0391 (17)	0.046 (6)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0314 (10)	0.0252 (8)	0.0276 (8)	0.0063 (7)	0.0026 (7)	0.0010 (6)
C2	0.0455 (11)	0.0217 (8)	0.0306 (9)	0.0031 (8)	0.0026 (8)	-0.0002 (7)
C3	0.0413 (11)	0.0308 (9)	0.0238 (8)	-0.0067 (8)	0.0004 (8)	0.0004 (7)
C4	0.0305 (10)	0.0400 (10)	0.0255 (8)	0.0011 (8)	-0.0013 (7)	-0.0011 (7)
C5	0.0327 (10)	0.0274 (9)	0.0237 (8)	0.0071 (8)	0.0011 (7)	-0.0002 (6)
C6	0.0305 (9)	0.0222 (8)	0.0185 (7)	0.0029 (7)	0.0023 (7)	0.0014 (6)
C7	0.0237 (8)	0.0223 (8)	0.0198 (7)	-0.0050 (6)	0.0076 (6)	-0.0039 (6)
C8	0.0258 (9)	0.0192 (8)	0.0423 (10)	-0.0029 (7)	0.0068 (8)	-0.0026 (7)
C9	0.0222 (8)	0.0201 (8)	0.0200 (7)	-0.0036 (6)	0.0059 (6)	-0.0018 (6)
C10	0.0267 (9)	0.0221 (8)	0.0196 (7)	-0.0037 (7)	0.0051 (6)	-0.0011 (6)
C11	0.0256 (9)	0.0186 (7)	0.0255 (7)	-0.0025 (6)	0.0044 (7)	0.0004 (6)
C12	0.0307 (9)	0.0226 (8)	0.0242 (7)	-0.0012 (7)	0.0061 (7)	-0.0005 (6)
C13	0.0269 (10)	0.0306 (10)	0.0476 (11)	0.0019 (8)	0.0015 (8)	-0.0021 (8)
C14	0.0246 (9)	0.0223 (8)	0.0231 (7)	-0.0015 (7)	0.0061 (6)	-0.0035 (6)
N1	0.0219 (7)	0.0230 (7)	0.0265 (7)	0.0009 (6)	0.0059 (6)	-0.0017 (5)
N2	0.0213 (7)	0.0210 (7)	0.0251 (6)	-0.0014 (6)	0.0051 (5)	0.0000 (5)
O1	0.0352 (7)	0.0369 (7)	0.0303 (6)	-0.0015 (6)	0.0129 (6)	0.0034 (5)

O2	0.0450 (8)	0.0206 (6)	0.0422 (7)	0.0044 (6)	-0.0010 (6)	0.0050 (5)
O3	0.0245 (6)	0.0243 (6)	0.0380 (7)	-0.0073 (5)	0.0031 (5)	0.0003 (5)
O4	0.0221 (6)	0.0243 (6)	0.0421 (7)	-0.0050 (5)	0.0072 (5)	-0.0061 (5)
O5	0.0227 (6)	0.0239 (6)	0.0318 (6)	-0.0020 (5)	0.0051 (5)	-0.0020 (5)
S	0.0308 (3)	0.0203 (2)	0.0266 (2)	0.00152 (17)	0.00559 (18)	0.00282 (14)

Geometric parameters (Å, °)

C1—C2	1.389 (3)	C9—C10	1.452 (2)
C1—C6	1.396 (2)	C10—O3	1.254 (2)
C1—H1	0.9500	C10—C11	1.445 (2)
C2—C3	1.386 (3)	C11—C12	1.332 (2)
C2—H2	0.9500	C11—H11	0.9500
C3—C4	1.388 (3)	C12—O5	1.369 (2)
C3—H3	0.9500	C12—C13	1.481 (3)
C4—C5	1.386 (3)	C13—H13A	0.9800
C4—H4	0.9500	C13—H13B	0.9800
C5—C6	1.391 (3)	C13—H13C	0.9800
C5—H5	0.9500	C14—O4	1.229 (2)
C6—S	1.7635 (18)	C14—O5	1.384 (2)
C7—N2	1.321 (2)	N1—N2	1.401 (2)
C7—C9	1.426 (2)	N1—S	1.6712 (15)
C7—C8	1.496 (2)	N1—HN1	0.89 (2)
C8—H8A	0.9800	N2—HN2	0.90 (2)
C8—H8B	0.9800	O1—S	1.4345 (13)
C8—H8C	0.9800	O2—S	1.4246 (13)
C9—C14	1.431 (2)		
C2—C1—C6	118.68 (17)	O3—C10—C9	123.66 (15)
C2—C1—H1	120.7	C11—C10—C9	116.55 (15)
C6—C1—H1	120.7	C12—C11—C10	121.76 (15)
C3—C2—C1	120.40 (17)	C12—C11—H11	119.1
C3—C2—H2	119.8	C10—C11—H11	119.1
C1—C2—H2	119.8	C11—C12—O5	121.38 (16)
C2—C3—C4	120.47 (18)	C11—C12—C13	126.11 (16)
C2—C3—H3	119.8	O5—C12—C13	112.47 (15)
C4—C3—H3	119.8	C12—C13—H13A	109.5
C5—C4—C3	119.92 (18)	C12—C13—H13B	109.5
C5—C4—H4	120.0	H13A—C13—H13B	109.5
C3—C4—H4	120.0	C12—C13—H13C	109.5
C4—C5—C6	119.38 (16)	H13A—C13—H13C	109.5
C4—C5—H5	120.3	H13B—C13—H13C	109.5
C6—C5—H5	120.3	O4—C14—O5	114.24 (15)
C5—C6—C1	121.14 (17)	O4—C14—C9	127.28 (16)
C5—C6—S	119.04 (13)	O5—C14—C9	118.48 (14)
C1—C6—S	119.81 (14)	N2—N1—S	113.54 (10)
N2—C7—C9	116.83 (14)	N2—N1—HN1	110.8 (13)
N2—C7—C8	119.20 (16)	S—N1—HN1	111.7 (14)

C9—C7—C8	123.96 (15)	C7—N2—N1	120.97 (14)
C7—C8—H8A	109.5	C7—N2—HN2	115.3 (16)
C7—C8—H8B	109.5	N1—N2—HN2	123.2 (16)
H8A—C8—H8B	109.5	C12—O5—C14	122.25 (13)
C7—C8—H8C	109.5	O2—S—O1	121.29 (8)
H8A—C8—H8C	109.5	O2—S—N1	104.51 (8)
H8B—C8—H8C	109.5	O1—S—N1	105.48 (8)
C7—C9—C14	120.05 (14)	O2—S—C6	108.08 (8)
C7—C9—C10	120.42 (15)	O1—S—C6	108.75 (8)
C14—C9—C10	119.45 (15)	N1—S—C6	108.02 (7)
O3—C10—C11	119.79 (14)		

Hydrogen-bond geometry (Å, °)

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
N1—HN1...O4 ⁱ	0.89 (2)	1.89 (2)	2.7837 (19)	175 (2)
N2—HN2...O3	0.90 (2)	1.74 (2)	2.5194 (18)	144 (2)

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