

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

3-*tert*-Butyl-1*H*-isochromene-1-thioneF. Nawaz Khan,^a P. Manivel,^a S. Mohana Roopan,^a
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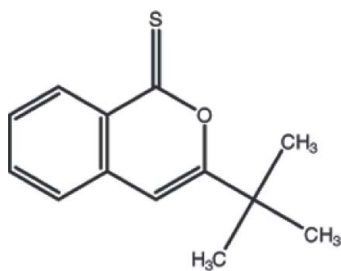
Received 18 May 2010; accepted 24 May 2010

Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.074; wR factor = 0.161; data-to-parameter ratio = 21.9.

The title compound, $\text{C}_{13}\text{H}_{14}\text{OS}$, crystallizes with two independent molecules in the asymmetric unit. The unit cell contains three voids of 197 Å³, but the residual electron density (highest peak = 0.24 e Å⁻³ and deepest hole = -0.18 e Å⁻³) in the difference Fourier map suggests no solvent molecule occupies this void. The crystal structure is stabilized by π - π interactions between the isocoumarin ring systems, with centroid-centroid distances of 3.6793 (14) and 3.6566 (15) Å.

Related literature

For the crystal structure and synthesis of isocoumarin and its thioanalogues, see: Hathwar *et al.* (2007*a,b*, 2009); Manivel *et al.* (2008); Basvanag *et al.* (2009); Henerson & Hill (1982).



Experimental

Crystal data

$\text{C}_{13}\text{H}_{14}\text{OS}$
 $M_r = 218.31$

Trigonal, $R\bar{3}$
 $a = 43.2799$ (16) Å

$c = 6.9025$ (5) Å
 $V = 11197.2$ (10) Å³
 $Z = 36$
Mo $K\alpha$ radiation

$\mu = 0.23$ mm⁻¹
 $T = 293$ K
 $0.31 \times 0.18 \times 0.15$ mm

Data collection

Bruker SMART CCD area-detector diffractometer
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.932$, $T_{\max} = 0.966$

33028 measured reflections
5943 independent reflections
3409 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.047$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.074$
 $wR(F^2) = 0.161$
 $S = 1.09$
5943 reflections

271 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.24$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.18$ e Å⁻³

Data collection: SMART (Bruker, 2004); cell refinement: SAINT (Bruker, 2004); data reduction: SAINT; 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, 2009).

We thank the Department of Science and Technology, India, for use of the CCD facility set up under the IRHPA-DST program at IISc. We thank Professor T. N. Guru Row, IISc, Bangalore, for the data collection. FNK thanks the DST for Fast Track Proposal funding.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FJ2306).

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supporting information

Acta Cryst. (2010). E66, o1470 [https://doi.org/10.1107/S1600536810019422]

3-*tert*-Butyl-1*H*-isochromene-1-thione

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S1. Comment

Isocoumarins are isolated in a great variety of microorganisms, plants, and insects, and have been shown to have considerable biological activity. Isocoumarins and its derivatives are secondary metabolites of a wide variety of microbial plant and insect sources and in synthesis of other medicinal compounds (Manivel *et al.*, 2008, Basvanag *et al.*, 2009). Sulfur containing isocoumarins have been known and a number of substituted thioisocoumarins, (Henerson *et al.*, 1982) have been prepared. Most methods available for the construction of thioisocoumarin nucleus suffer from one or more drawbacks, such as the long reaction time required obtaining a good yield of the desired product or the use of expensive and hazardous reagents and solvents. Herewith, we are reporting the synthesis of 3-*tert*-butyl-1*H*-isochromene-1-thione using Lawessons reagent.

The isocoumarin moieties in the symmetric unit of the title compound, (I), (Fig. 1a and Fig. 1 b), are essentially parallel to each other with a small dihedral angle of 1.20 (7) °.

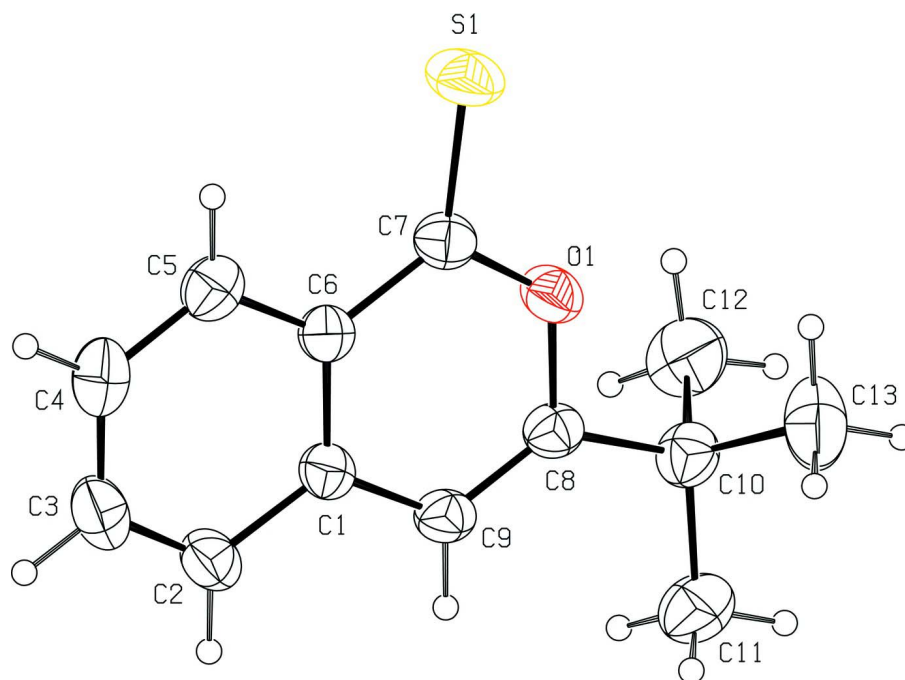
In the molecular structure of (I), there exist C—H···S and C—H···O intramolecular interactions (Table 1, Fig. 2). In addition, π - π interactions are observed between the isocoumarin ring systems [$Cg2\cdots Cg4(x, y, z) = 3.6793$ (14) Å and $Cg2\cdots Cg5(x, y, 1 + z) = 3.6566$ (15) Å; where $Cg2$, $Cg4$ and $Cg5$ are the centroids of the C1–C6, O2'/C1'/C6'–C9' and C1'–C6' rings, respectively]. There is no classic hydrogen bonds in the structure.

S2. Experimental

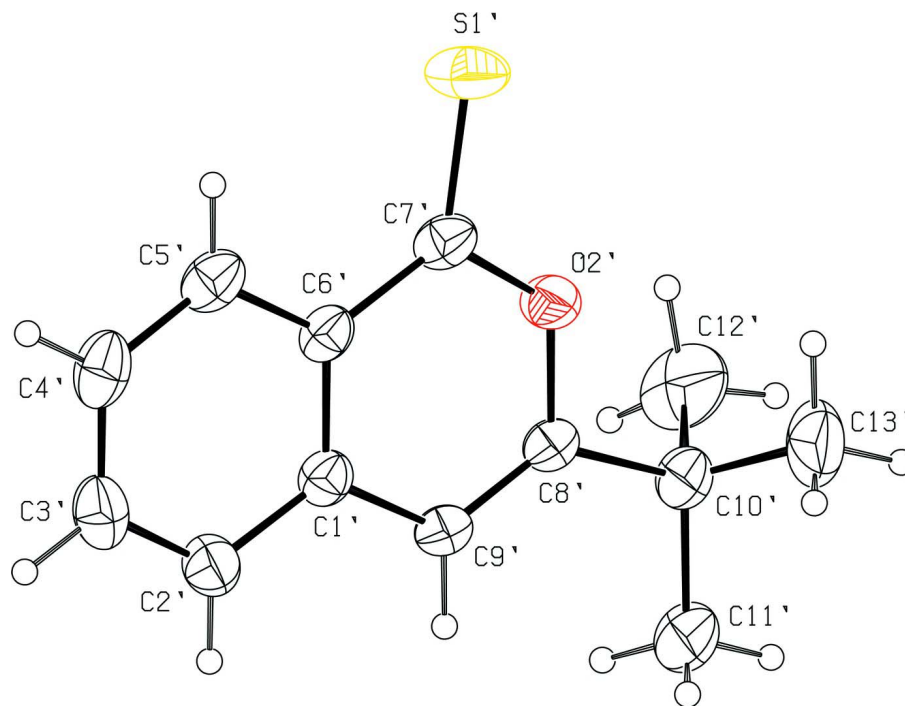
The 3-*tert*-butyl-1*H*-isochromen-1-one and Lawessons reagent were taken in toluene (1:1 ratio) and refluxed for 1 h. Then the reaction mass was quenched with water, extracted with dichloromethane, washed with water, dried, concentrated and purified by column chromatography to get the titled compound. Single crystals of the title compound were obtained *via* recrystallization from a chloroform solution.

S3. Refinement

All H-atoms were placed in calculated positions (C—H = 0.93 and 0.96 Å) and were included in the refinement in the riding model approximation, with $U_{iso}(H)$ set to 1.2 or 1.5 $U_{eq}(C)$. In the crystal structure, there is an 197 Å³ void, but the low electron density (0.24 e.Å⁻³) in the difference Fourier map suggests no solvent molecule occupying this void.

**Figure 1**

The first molecule in the asymmetric unit with the atom numbering scheme. Displacement ellipsoids for non-H atoms are drawn at the 30% probability level.

**Figure 2**

The second molecule in the asymmetric unit with the atom numbering scheme. Displacement ellipsoids for non-H atoms are drawn at the 30% probability level.

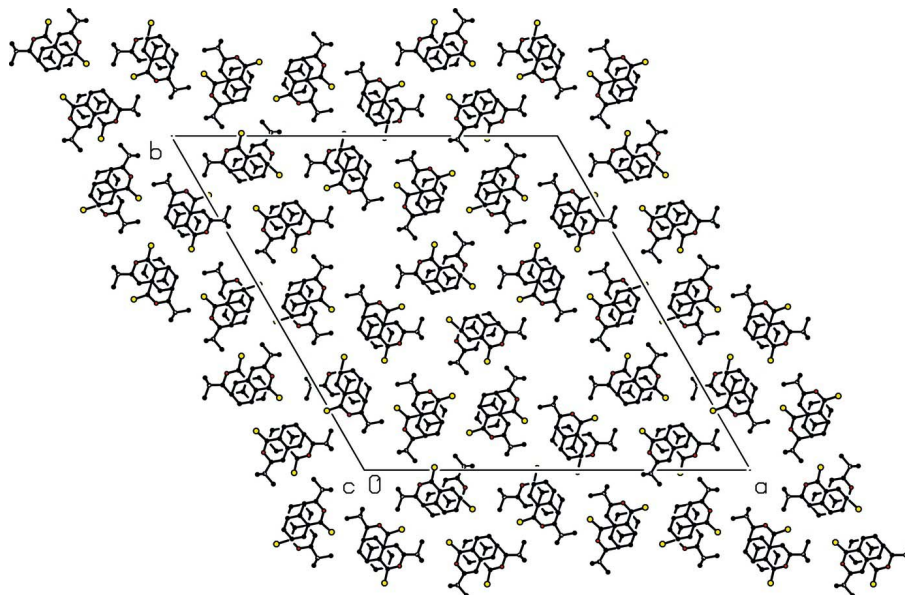


Figure 3

The packing diagram of the title compound down *c* axis in the unitcell.

3-*tert*-Butyl-1*H*-isochromene-1-thione

Crystal data

$C_{13}H_{14}OS$

$M_r = 218.31$

Trigonal, $R\bar{3}$

Hall symbol: $-R\bar{3}$

$a = 43.2799$ (16) Å

$c = 6.9025$ (5) Å

$V = 11197.2$ (10) Å³

$Z = 36$

$F(000) = 4176$

$D_x = 1.166$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 856 reflections

$\theta = 1.9$ – 24.7°

$\mu = 0.23$ mm⁻¹

$T = 293$ K

Block, yellow

$0.31 \times 0.18 \times 0.15$ mm

Data collection

Bruker SMART CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.932$, $T_{\max} = 0.966$

33028 measured reflections

5943 independent reflections

3409 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.047$

$\theta_{\max} = 28.0^\circ$, $\theta_{\min} = 1.6^\circ$

$h = -57 \rightarrow 57$

$k = -56 \rightarrow 56$

$l = -9 \rightarrow 9$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.074$

$wR(F^2) = 0.161$

$S = 1.09$

5943 reflections

271 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

$$w = 1/[\sigma^2(F_o^2) + (0.0599P)^2 + 4.5485P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$

$$\Delta\rho_{\max} = 0.24 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.18 \text{ e } \text{\AA}^{-3}$$

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 esds 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	0.81419 (2)	-0.00817 (2)	0.78291 (13)	0.0811 (3)
O1	0.87826 (4)	0.04385 (4)	0.7757 (3)	0.0606 (7)
C1	0.86386 (7)	0.09888 (6)	0.7809 (3)	0.0502 (8)
C2	0.85585 (8)	0.12659 (7)	0.7850 (4)	0.0631 (10)
C3	0.82145 (9)	0.11923 (8)	0.7911 (4)	0.0729 (11)
C4	0.79363 (8)	0.08436 (9)	0.7932 (4)	0.0720 (11)
C5	0.80057 (7)	0.05677 (7)	0.7903 (4)	0.0635 (10)
C6	0.83556 (6)	0.06358 (6)	0.7841 (3)	0.0492 (8)
C7	0.84331 (7)	0.03469 (6)	0.7812 (4)	0.0548 (9)
C8	0.90635 (6)	0.07871 (6)	0.7703 (3)	0.0516 (8)
C9	0.89948 (7)	0.10530 (6)	0.7739 (3)	0.0535 (8)
C10	0.94129 (7)	0.07913 (7)	0.7620 (4)	0.0639 (10)
C11	0.97229 (7)	0.11727 (8)	0.7400 (5)	0.0878 (13)
C12	0.94624 (9)	0.06357 (9)	0.9523 (5)	0.0979 (16)
C13	0.94106 (9)	0.05714 (9)	0.5875 (5)	0.1021 (16)
S1'	0.77849 (2)	0.11691 (2)	0.28420 (12)	0.0772 (3)
O2'	0.76501 (4)	0.05251 (4)	0.2971 (3)	0.0574 (6)
C1'	0.83413 (6)	0.06544 (6)	0.2874 (3)	0.0465 (8)
C2'	0.86926 (6)	0.07253 (7)	0.2822 (4)	0.0574 (9)
C3'	0.89714 (7)	0.10670 (8)	0.2764 (4)	0.0649 (10)
C4'	0.89068 (7)	0.13502 (7)	0.2747 (4)	0.0660 (10)
C5'	0.85652 (7)	0.12899 (6)	0.2800 (3)	0.0579 (9)
C6'	0.82767 (6)	0.09412 (6)	0.2863 (3)	0.0459 (8)
C7'	0.79136 (6)	0.08729 (6)	0.2900 (3)	0.0520 (8)
C8'	0.77110 (6)	0.02397 (6)	0.2971 (4)	0.0521 (8)
C9'	0.80418 (6)	0.02988 (6)	0.2925 (3)	0.0524 (9)
C10'	0.73624 (7)	-0.01086 (6)	0.3019 (5)	0.0680 (10)
C11'	0.74338 (8)	-0.04203 (7)	0.2923 (5)	0.0903 (13)
C12'	0.71642 (9)	-0.01317 (8)	0.4895 (6)	0.1127 (16)
C13'	0.71430 (8)	-0.01215 (8)	0.1256 (6)	0.1090 (16)
H2	0.87430	0.15020	0.78360	0.0760*
H3	0.81660	0.13790	0.79390	0.0880*

H4	0.77020	0.07960	0.79660	0.0870*
H5	0.78180	0.03330	0.79240	0.0760*
H9	0.91830	0.12870	0.77170	0.0640*
H11A	0.96880	0.12780	0.62590	0.1320*
H11B	0.99420	0.11690	0.72860	0.1320*
H11C	0.97330	0.13100	0.85160	0.1320*
H12A	0.94830	0.07890	1.05820	0.1460*
H12B	0.96750	0.06190	0.94440	0.1460*
H12C	0.92600	0.04030	0.97290	0.1460*
H13A	0.92160	0.03310	0.59980	0.1540*
H13B	0.96320	0.05700	0.58270	0.1540*
H13C	0.93820	0.06750	0.47070	0.1540*
H2'	0.87380	0.05370	0.28280	0.0690*
H3'	0.92050	0.11100	0.27370	0.0780*
H4'	0.90970	0.15830	0.26980	0.0790*
H5'	0.85250	0.14820	0.27940	0.0700*
H9'	0.80810	0.01060	0.29270	0.0630*
H11D	0.75560	-0.04070	0.17380	0.1360*
H11E	0.72110	-0.06410	0.29670	0.1360*
H11F	0.75790	-0.04080	0.40050	0.1360*
H12D	0.72880	-0.01600	0.59730	0.1690*
H12E	0.69270	-0.03330	0.48330	0.1690*
H12F	0.71530	0.00830	0.50580	0.1690*
H13D	0.71030	0.00780	0.12930	0.1630*
H13E	0.69180	-0.03400	0.12730	0.1630*
H13F	0.72700	-0.01110	0.00950	0.1630*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0696 (5)	0.0415 (4)	0.1141 (7)	0.0142 (3)	-0.0067 (4)	-0.0026 (4)
O1	0.0547 (11)	0.0442 (10)	0.0807 (13)	0.0231 (8)	-0.0041 (9)	-0.0046 (8)
C1	0.0629 (16)	0.0480 (14)	0.0398 (14)	0.0278 (13)	0.0024 (11)	-0.0030 (11)
C2	0.082 (2)	0.0539 (16)	0.0593 (17)	0.0383 (15)	0.0031 (14)	-0.0037 (13)
C3	0.100 (2)	0.079 (2)	0.0639 (19)	0.063 (2)	0.0030 (16)	-0.0045 (15)
C4	0.0701 (19)	0.097 (2)	0.0658 (19)	0.0544 (19)	0.0005 (15)	-0.0057 (17)
C5	0.0550 (16)	0.0694 (18)	0.0613 (17)	0.0275 (14)	0.0006 (13)	-0.0017 (14)
C6	0.0540 (14)	0.0494 (14)	0.0420 (14)	0.0243 (12)	-0.0004 (11)	-0.0034 (11)
C7	0.0536 (15)	0.0482 (14)	0.0561 (16)	0.0206 (12)	-0.0047 (12)	-0.0036 (12)
C8	0.0493 (14)	0.0467 (14)	0.0497 (15)	0.0171 (12)	-0.0020 (11)	-0.0034 (11)
C9	0.0575 (15)	0.0429 (13)	0.0497 (15)	0.0173 (12)	0.0015 (12)	-0.0028 (11)
C10	0.0535 (16)	0.0663 (17)	0.0679 (19)	0.0270 (14)	-0.0014 (13)	-0.0039 (14)
C11	0.0572 (18)	0.082 (2)	0.107 (3)	0.0220 (16)	0.0055 (17)	0.0042 (18)
C12	0.080 (2)	0.115 (3)	0.111 (3)	0.058 (2)	0.0003 (19)	0.028 (2)
C13	0.081 (2)	0.117 (3)	0.121 (3)	0.059 (2)	-0.004 (2)	-0.037 (2)
S1'	0.0796 (5)	0.0522 (4)	0.1108 (7)	0.0413 (4)	0.0313 (4)	0.0127 (4)
O2'	0.0444 (9)	0.0391 (9)	0.0862 (13)	0.0191 (8)	0.0055 (8)	-0.0002 (8)
C1'	0.0473 (13)	0.0459 (13)	0.0432 (14)	0.0209 (11)	0.0008 (11)	-0.0034 (10)

C2'	0.0506 (15)	0.0635 (16)	0.0586 (17)	0.0288 (13)	-0.0024 (12)	-0.0038 (13)
C3'	0.0444 (15)	0.077 (2)	0.0610 (18)	0.0212 (14)	-0.0013 (12)	-0.0015 (14)
C4'	0.0506 (16)	0.0582 (17)	0.0604 (18)	0.0055 (13)	0.0034 (13)	-0.0005 (13)
C5'	0.0616 (17)	0.0445 (14)	0.0561 (16)	0.0178 (12)	0.0086 (13)	-0.0021 (12)
C6'	0.0447 (13)	0.0410 (12)	0.0430 (14)	0.0147 (11)	0.0056 (10)	-0.0042 (10)
C7'	0.0559 (15)	0.0414 (13)	0.0536 (16)	0.0205 (12)	0.0112 (12)	-0.0004 (11)
C8'	0.0507 (14)	0.0364 (12)	0.0675 (17)	0.0206 (11)	-0.0047 (12)	-0.0061 (11)
C9'	0.0539 (15)	0.0396 (13)	0.0665 (17)	0.0255 (12)	-0.0043 (12)	-0.0069 (11)
C10'	0.0500 (15)	0.0382 (14)	0.105 (2)	0.0140 (12)	-0.0103 (16)	-0.0065 (14)
C11'	0.0727 (19)	0.0383 (15)	0.147 (3)	0.0181 (14)	-0.023 (2)	-0.0098 (17)
C12'	0.078 (2)	0.061 (2)	0.158 (4)	0.0039 (17)	0.041 (2)	0.006 (2)
C13'	0.072 (2)	0.063 (2)	0.168 (4)	0.0157 (17)	-0.047 (2)	-0.012 (2)

Geometric parameters (Å, °)

S1—C7	1.641 (2)	C13—H13A	0.9600
S1'—C7'	1.634 (3)	C13—H13B	0.9600
O1—C8	1.386 (3)	C13—H13C	0.9600
O1—C7	1.359 (4)	C1'—C2'	1.393 (4)
O2'—C8'	1.386 (3)	C1'—C6'	1.402 (4)
O2'—C7'	1.361 (3)	C1'—C9'	1.434 (3)
C1—C6	1.401 (3)	C2'—C3'	1.364 (4)
C1—C2	1.405 (4)	C3'—C4'	1.387 (4)
C1—C9	1.424 (5)	C4'—C5'	1.367 (5)
C2—C3	1.359 (6)	C5'—C6'	1.398 (3)
C3—C4	1.382 (5)	C6'—C7'	1.447 (4)
C4—C5	1.369 (5)	C8'—C9'	1.323 (4)
C5—C6	1.391 (4)	C8'—C10'	1.509 (4)
C6—C7	1.448 (4)	C10'—C11'	1.529 (4)
C8—C9	1.325 (4)	C10'—C12'	1.529 (5)
C8—C10	1.504 (4)	C10'—C13'	1.527 (5)
C10—C11	1.528 (4)	C2'—H2'	0.9300
C10—C12	1.539 (5)	C3'—H3'	0.9300
C10—C13	1.532 (4)	C4'—H4'	0.9300
C2—H2	0.9300	C5'—H5'	0.9300
C3—H3	0.9300	C9'—H9'	0.9300
C4—H4	0.9300	C11'—H11D	0.9600
C5—H5	0.9300	C11'—H11E	0.9600
C9—H9	0.9300	C11'—H11F	0.9600
C11—H11A	0.9600	C12'—H12D	0.9600
C11—H11B	0.9600	C12'—H12E	0.9600
C11—H11C	0.9600	C12'—H12F	0.9600
C12—H12C	0.9600	C13'—H13D	0.9600
C12—H12A	0.9600	C13'—H13E	0.9600
C12—H12B	0.9600	C13'—H13F	0.9600
S1'...C5' ⁱⁱ	3.681 (2)	H9...H11A	2.4200
S1...H3' ⁱⁱ	3.0200	H9...H11C	2.4000

S1...H11C ⁱⁱⁱ	3.0800	H9'...C11'	2.5800
S1...H5	2.7800	H9'...H2'	2.5000
S1'...H11E ^{iv}	3.1800	H9'...H11D	2.3900
S1'...H5'	2.7900	H9'...H11F	2.3200
S1'...H5' ⁱ	3.0100	H11A...H9	2.4200
S1'...H2' ^v	3.2000	H11A...H13C	2.5000
O1...H13A	2.4700	H11A...C9	2.8400
O1...H12C	2.5400	H11B...H12B	2.5400
O2'...H12F	2.5000	H11B...H13B	2.4600
O2'...H13D	2.4700	H11C...H9	2.4000
C1'...C6	3.431 (3)	H11C...C9	2.8600
C1'...C6 ^{vi}	3.476 (3)	H11C...H12A	2.4200
C2...C5'	3.487 (3)	H11C...S1 ^{ix}	3.0800
C2...C5' ^{vii}	3.418 (3)	H11D...H13F	2.4600
C3'...C9 ^{vi}	3.471 (3)	H11D...C9'	2.8300
C3'...C9	3.437 (3)	H11D...H9'	2.3900
C4...C7' ^{vii}	3.435 (3)	H11E...H12E	2.5700
C4...C7'	3.479 (3)	H11E...H13E	2.5100
C5'...S1' ⁱ	3.681 (2)	H11E...S1' ^{ix}	3.1800
C5'...C2	3.487 (3)	H11F...C9'	2.7900
C5'...C2 ^{vi}	3.418 (3)	H11F...H12D	2.4400
C6...C1' ^{vii}	3.476 (3)	H11F...H9'	2.3200
C6...C1'	3.431 (3)	H12A...H11C	2.4200
C7'...C4 ^{vi}	3.435 (3)	H12B...H13B	2.5000
C7'...C4	3.479 (3)	H12B...H11B	2.5400
C9...C3'	3.437 (3)	H12C...H13A	2.5900
C9...C3' ^{vii}	3.471 (3)	H12C...O1	2.5400
C9...H11A	2.8400	H12D...H11F	2.4400
C9...H11C	2.8600	H12D...H5	2.5900
C9'...H11F	2.7900	H12E...H13E	2.4600
C9'...H11D	2.8300	H12E...H11E	2.5700
C11...H9	2.6300	H12F...O2'	2.5000
C11'...H9'	2.5800	H12F...H13E ^{iv}	2.5900
H2...H9	2.5000	H13A...H12C	2.5900
H2...S1' ^{iv}	3.2000	H13A...O1	2.4700
H2'...H9'	2.5000	H13B...H12B	2.5000
H3'...S1' ^{viii}	3.0200	H13B...H11B	2.4600
H5...H12D	2.5900	H13C...H11A	2.5000
H5...S1	2.7800	H13D...O2'	2.4700
H5'...S1'	2.7900	H13E...H11E	2.5100
H5'...S1' ⁱ	3.0100	H13E...H12E	2.4600
H9...C11	2.6300	H13E...H12F ^x	2.5900
H9...H2	2.5000	H13F...H11D	2.4600
C7—O1—C8	124.1 (2)	H13A—C13—H13B	109.00
C7'—O2'—C8'	123.9 (2)	C2'—C1'—C6'	118.9 (2)
C2—C1—C6	118.4 (3)	C2'—C1'—C9'	122.6 (2)
C2—C1—C9	122.6 (2)	C6'—C1'—C9'	118.5 (3)

C6—C1—C9	119.0 (2)	C1'—C2'—C3'	121.1 (3)
C1—C2—C3	120.6 (3)	C2'—C3'—C4'	119.9 (3)
C2—C3—C4	120.7 (3)	C3'—C4'—C5'	120.5 (3)
C3—C4—C5	120.1 (4)	C4'—C5'—C6'	120.3 (3)
C4—C5—C6	120.4 (3)	C1'—C6'—C5'	119.3 (3)
C1—C6—C7	119.2 (3)	C1'—C6'—C7'	119.7 (2)
C5—C6—C7	121.0 (2)	C5'—C6'—C7'	120.9 (2)
C1—C6—C5	119.8 (2)	S1'—C7'—O2'	116.3 (2)
S1—C7—C6	126.7 (2)	S1'—C7'—C6'	126.93 (18)
O1—C7—C6	117.0 (2)	O2'—C7'—C6'	116.8 (2)
S1—C7—O1	116.3 (2)	O2'—C8'—C9'	119.8 (2)
O1—C8—C10	110.1 (2)	O2'—C8'—C10'	110.5 (2)
C9—C8—C10	130.6 (2)	C9'—C8'—C10'	129.7 (2)
O1—C8—C9	119.3 (3)	C1'—C9'—C8'	121.2 (2)
C1—C9—C8	121.5 (2)	C8'—C10'—C11'	109.8 (3)
C8—C10—C12	108.7 (3)	C8'—C10'—C12'	109.4 (2)
C8—C10—C13	109.2 (3)	C8'—C10'—C13'	108.2 (2)
C8—C10—C11	110.6 (2)	C11'—C10'—C12'	109.5 (3)
C11—C10—C13	108.8 (3)	C11'—C10'—C13'	109.2 (3)
C12—C10—C13	111.0 (3)	C12'—C10'—C13'	110.8 (3)
C11—C10—C12	108.6 (3)	C1'—C2'—H2'	120.00
C3—C2—H2	120.00	C3'—C2'—H2'	119.00
C1—C2—H2	120.00	C2'—C3'—H3'	120.00
C2—C3—H3	120.00	C4'—C3'—H3'	120.00
C4—C3—H3	120.00	C3'—C4'—H4'	120.00
C3—C4—H4	120.00	C5'—C4'—H4'	120.00
C5—C4—H4	120.00	C4'—C5'—H5'	120.00
C6—C5—H5	120.00	C6'—C5'—H5'	120.00
C4—C5—H5	120.00	C1'—C9'—H9'	119.00
C1—C9—H9	119.00	C8'—C9'—H9'	119.00
C8—C9—H9	119.00	C10'—C11'—H11D	109.00
C10—C11—H11B	109.00	C10'—C11'—H11E	109.00
C10—C11—H11C	109.00	C10'—C11'—H11F	109.00
C10—C11—H11A	109.00	H11D—C11'—H11E	110.00
H11A—C11—H11C	110.00	H11D—C11'—H11F	109.00
H11B—C11—H11C	110.00	H11E—C11'—H11F	110.00
H11A—C11—H11B	109.00	C10'—C12'—H12D	110.00
C10—C12—H12A	109.00	C10'—C12'—H12E	109.00
C10—C12—H12B	109.00	C10'—C12'—H12F	110.00
H12A—C12—H12B	109.00	H12D—C12'—H12E	109.00
H12A—C12—H12C	109.00	H12D—C12'—H12F	110.00
C10—C12—H12C	109.00	H12E—C12'—H12F	110.00
H12B—C12—H12C	110.00	C10'—C13'—H13D	109.00
C10—C13—H13B	110.00	C10'—C13'—H13E	109.00
C10—C13—H13C	109.00	C10'—C13'—H13F	109.00
C10—C13—H13A	109.00	H13D—C13'—H13E	110.00
H13A—C13—H13C	109.00	H13D—C13'—H13F	110.00
H13B—C13—H13C	110.00	H13E—C13'—H13F	109.00

C8—O1—C7—S1	178.89 (18)	C10—C8—C9—C1	179.7 (2)
C8—O1—C7—C6	-0.9 (4)	O1—C8—C10—C11	175.6 (2)
C7—O1—C8—C9	1.2 (3)	O1—C8—C10—C12	-65.3 (3)
C7—O1—C8—C10	-179.1 (2)	O1—C8—C10—C13	55.9 (3)
C8'—O2'—C7'—S1'	178.1 (2)	C6'—C1'—C2'—C3'	0.1 (4)
C8'—O2'—C7'—C6'	-1.3 (3)	C9'—C1'—C2'—C3'	179.7 (2)
C7'—O2'—C8'—C9'	0.9 (4)	C2'—C1'—C6'—C5'	0.0 (3)
C7'—O2'—C8'—C10'	-178.9 (2)	C2'—C1'—C6'—C7'	179.5 (2)
C9—C1—C2—C3	179.8 (2)	C9'—C1'—C6'—C5'	-179.56 (19)
C2—C1—C6—C5	0.3 (3)	C9'—C1'—C6'—C7'	-0.1 (3)
C6—C1—C2—C3	-0.3 (4)	C2'—C1'—C9'—C8'	-179.9 (2)
C6—C1—C9—C8	0.0 (3)	C6'—C1'—C9'—C8'	-0.4 (3)
C9—C1—C6—C5	-179.8 (2)	C1'—C2'—C3'—C4'	-0.3 (4)
C9—C1—C6—C7	0.3 (3)	C2'—C3'—C4'—C5'	0.4 (4)
C2—C1—C6—C7	-179.6 (2)	C3'—C4'—C5'—C6'	-0.3 (4)
C2—C1—C9—C8	180.0 (2)	C4'—C5'—C6'—C1'	0.1 (3)
C1—C2—C3—C4	-0.1 (4)	C4'—C5'—C6'—C7'	-179.4 (2)
C2—C3—C4—C5	0.4 (4)	C1'—C6'—C7'—S1'	-178.42 (17)
C3—C4—C5—C6	-0.4 (4)	C1'—C6'—C7'—O2'	0.9 (3)
C4—C5—C6—C7	179.9 (2)	C5'—C6'—C7'—S1'	1.1 (3)
C4—C5—C6—C1	0.0 (4)	C5'—C6'—C7'—O2'	-179.7 (2)
C1—C6—C7—O1	0.1 (3)	O2'—C8'—C9'—C1'	0.0 (4)
C5—C6—C7—S1	0.5 (4)	C10'—C8'—C9'—C1'	179.7 (3)
C5—C6—C7—O1	-179.8 (2)	O2'—C8'—C10'—C11'	177.4 (2)
C1—C6—C7—S1	-179.63 (19)	O2'—C8'—C10'—C12'	-62.5 (3)
O1—C8—C9—C1	-0.8 (3)	O2'—C8'—C10'—C13'	58.3 (3)
C9—C8—C10—C11	-4.8 (4)	C9'—C8'—C10'—C11'	-2.4 (4)
C9—C8—C10—C12	114.3 (3)	C9'—C8'—C10'—C12'	117.8 (3)
C9—C8—C10—C13	-124.5 (3)	C9'—C8'—C10'—C13'	-121.5 (3)

Symmetry codes: (i) $-x+5/3, -y+1/3, -z+1/3$; (ii) $x-y, x-1, -z+1$; (iii) $x-y, x-1, -z+2$; (iv) $-y+2/3, x-y-2/3, z+1/3$; (v) $-x+5/3, -y+1/3, -z+4/3$; (vi) $x, y, z-1$; (vii) $x, y, z+1$; (viii) $y+1, -x+y+1, -z+1$; (ix) $y+1, -x+y+1, -z+2$; (x) $-x+y+4/3, -x+2/3, z-1/3$.

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

<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>
C5—H5 \cdots S1	0.93	2.78	3.148 (3)	105
C5'—H5' \cdots S1'	0.93	2.79	3.149 (3)	104
C12—H12C \cdots O1	0.96	2.54	2.891 (5)	102
C12'—H12F \cdots O2'	0.96	2.50	2.879 (4)	103
C13—H13A \cdots O1	0.96	2.47	2.800 (5)	100
C13'—H13D \cdots O2'	0.96	2.47	2.812 (4)	101