organic compounds

(6-Methoxy-2-oxo-2*H*-chromen-4-yl)methyl piperidine-1-carbodithioate

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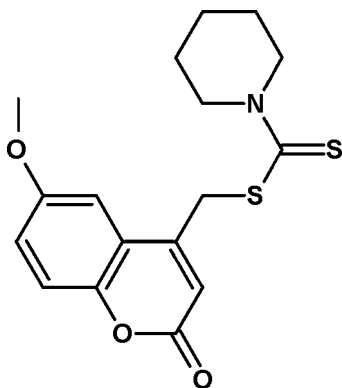Correspondence e-mail: devarajegowda@yahoo.com

In the title compound, $C_{17}H_{19}NO_3S_2$, the maximum deviation of atoms in the 2*H*-chromene ring system is 0.0097 (14) Å and the piperidine ring adopts a chair conformation. The dihedral angle between the 2*H*-chromene ring and the piperidine ring (all atoms) is 87.59 (8)°. In the crystal, inversion dimers linked by pairs of C—H···O interactions generate $R_2^2(22)$ loops. Further C—H···O hydrogen bonds link the dimers into [110] chains and weak aromatic π–π stacking [shortest centroid–centroid distance = 3.824 (8) Å] is also observed.

Related literature

For a related structure and the synthesis, see: Kumar *et al.* (2012).

Experimental

Crystal data

$C_{17}H_{19}NO_3S_2$	γ = 106.497 (1)°
M_r = 349.45	V = 819.96 (4) Å³
Triclinic, $P\bar{1}$	Z = 2
a = 6.9731 (2) Å	Mo $K\alpha$ radiation
b = 10.2310 (3) Å	μ = 0.34 mm⁻¹
c = 11.9955 (3) Å	T = 296 K
α = 92.024 (1)°	0.24 × 0.20 × 0.12 mm
β = 90.176 (1)°	

Data collection

Bruker SMART CCD diffractometer	17441 measured reflections
Absorption correction: multi-scan (*SADABS*; Bruker, 2007)	4261 independent reflections
	3513 reflections with $I > 2\sigma(I)$
T_{min} = 0.770, T_{max} = 1.000	R_{int} = 0.023

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.037	208 parameters
$wR(F^2)$ = 0.109	H-atom parameters constrained
S = 1.04	$\Delta\rho_{max}$ = 0.28 e Å⁻³
4261 reflections	$\Delta\rho_{min}$ = −0.18 e Å⁻³

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
C19—H19B···O5^i	0.97	2.50	3.410 (2)	157
C23—H23A···O3^{ii}	0.97	2.60	3.365 (2)	136

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

Data collection: *SMART* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012); software used to prepare material for publication: *SHELXL97*.

References

Bruker (2007). *SMART, SAINT* and *SADABS*. Bruker AXS Inc., Madison, Wisconsin, USA.
Farrugia, L. J. (2012). *J. Appl. Cryst.* **45**, 849–854.
Kumar, K. M., Kour, D., Kapoor, K., Mahabaleshwaraiah, N. M., Kotresh, O., Gupta, V. K. & Kant, R. (2012). *Acta Cryst.* E**68**, o878–o879.
Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.

supporting information

(6-Methoxy-2-oxo-2*H*-chromen-4-yl)methyl piperidine-1-carbodithioate

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S1. Comment

As part of our ongoing structural studies of coumarin derivatives (Kumar *et al.*, 2012), we now describe the title compound, (I) (Fig. 1).

The 2*H*- chromene ring system is almost planar, with a maximum deviation of 0.0097 (14) Å for atom C9 and the piperidine ring adopts a chair conformation. The dihedral angle between the 2*H*-chromene (O3/C7–C15) ring and the piperidine ring (N6/C19–C23) is 87.59 (8)°. In the crystal structure, C19—H19B⋯O5 and C23—H23A⋯O3 hydrogen bonding (Table 1) and π–π interactions between the fused benzene ring (C10–C15) of 2*H*-chromene and fused pyran ring (O3/C7–C11) [shortest centroid–centroid distance = 3.824 (8) Å] occur.

S2. Experimental

This compound was prepared according to the reported method (Kumar *et al.*, 2012). Colourless needles of the title compound were grown from a mixed solution of EtOH / CHCl$_3$(V/V = 1/1) by slow evaporation at room temperature. Colour: yellowish. Yield= 79%, m.p.395 K. IR (KBr) 655 cm^{-1}(C—S), 1243 cm^{-1}(C=S), 1006 cm^{-1}(C—O), 887 cm^{-1} (C—N), 1161 cm^{-1}(C—O—C), 1719 cm^{-1}(C=O). GCMS: m/e: 349. 1H NMR (400 MHz, CDCl$_3$, \?, p.p.m.): 1.65(s, 6H, Piperidine-CH$_2$), 3.15(s, 3H, –OCH$_3$), 3.89(s, 2H, Piperidine-CH$_2$), 4.20(s, 2H Piperidine-CH$_2$), 4.70(d, 2H, Methylene-CH$_2$), 6.45(s, 1H, Ar—H), 7.14(d, 1H, Ar—H), 7.30(s, 1H, Ar—H), 7.50(s,1*H*, Ar—H). Elemental analysis for C$_{17}$H$_{19}$NO$_3$S$_2$: C, 58.36; H, 5.42; N, 3.93.

S3. Refinement

All H atoms were positioned geometrically, with C—H = 0.93 Å for aromatic H, C—H = 0.97 Å for methylene H and C—H = 0.96 Å for methyl H,and refined using a riding model with U_{iso}(H) = 1.5U_{eq}(C) for methyl H and U_{iso}(H) = 1.2U_{eq}(C) for all other H
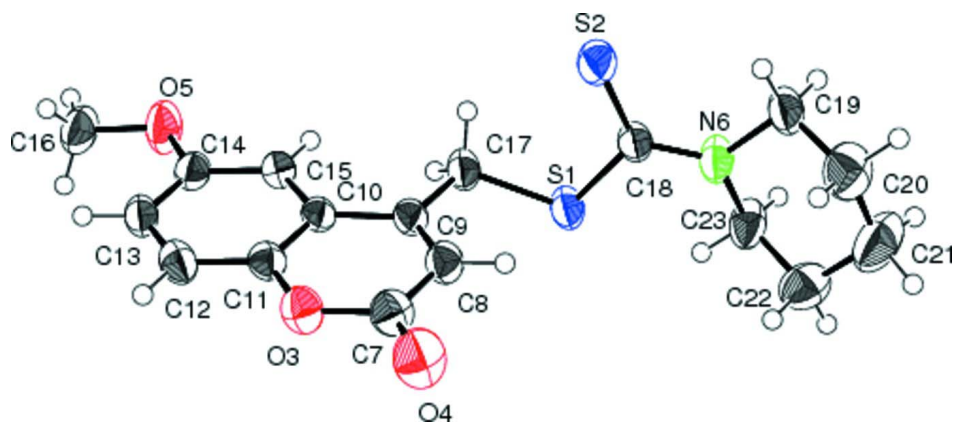
Figure 1

The molecular structure of the title compound. Displacement ellipsoids are drawn at the 50% probability level. Hydrogen atoms are shown as spheres of arbitrary radius.
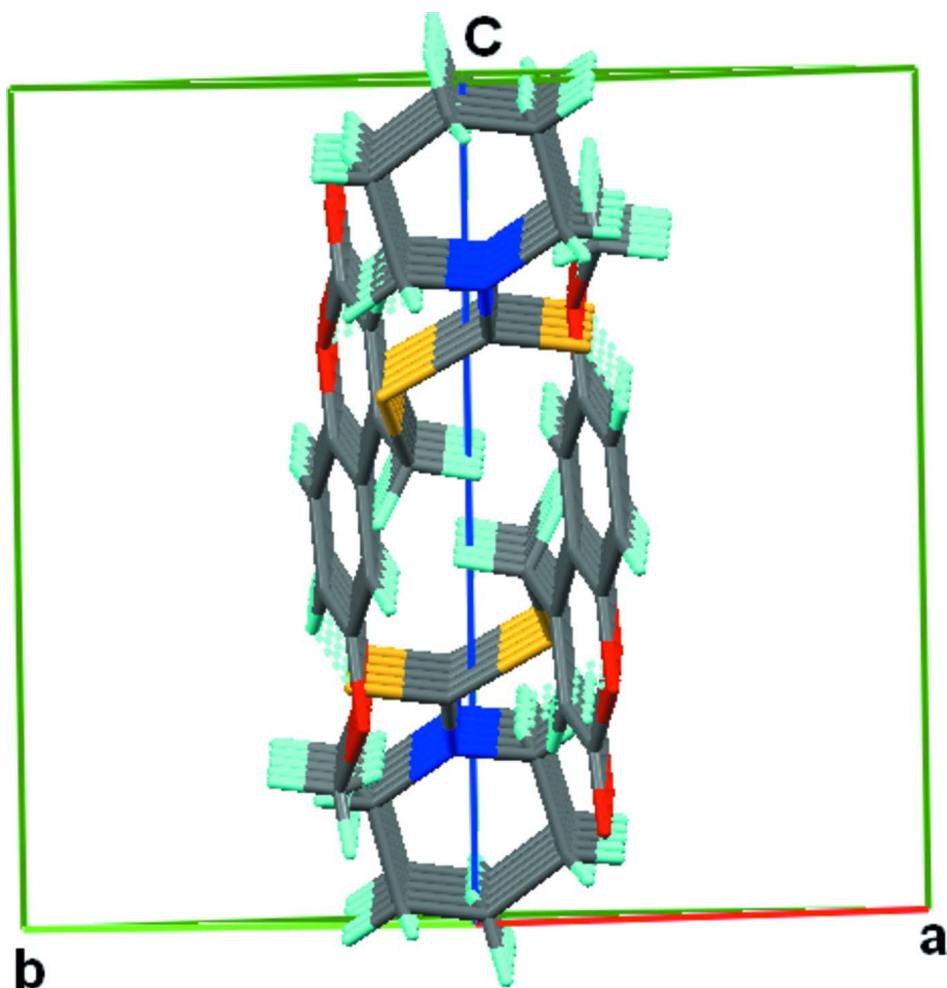
Figure 2

The packing of molecules.

(6-Methoxy-2-oxo-2*H*-chromen-4-yl)methyl piperidine-1-carbodithioate

Crystal data

$C_{17}H_{19}NO_3S_2$	$F(000) = 368$
$M_r = 349.45$	$D_x = 1.415$ Mg m^{-3}
Triclinic, $P\overline{1}$	Melting point: 395 K
$a = 6.9731$ (2) Å	Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
$b = 10.2310$ (3) Å	Cell parameters from 4261 reflections
$c = 11.9955$ (3) Å	$\theta = 1.7$–28.8°
$\alpha = 92.024$ (1)°	$\mu = 0.34$ mm^{-1}
$\beta = 90.176$ (1)°	$T = 296$ K
$\gamma = 106.497$ (1)°	Plate, colourless
$V = 819.96$ (4) Å^3	$0.24 \times 0.20 \times 0.12$ mm
$Z = 2$	

Data collection

Bruker SMART CCD diffractometer	17441 measured reflections
Radiation source: fine-focus sealed tube	4261 independent reflections
Graphite monochromator	3513 reflections with $I > 2\sigma(I)$
ω and φ scans	$R_{int} = 0.023$
Absorption correction: multi-scan (*SADABS*; Bruker, 2007)	$\theta_{max} = 28.8°$, $\theta_{min} = 1.7°$
	$h = -9 \rightarrow 9$
	$k = -11 \rightarrow 13$
$T_{min} = 0.770$, $T_{max} = 1.000$	$l = -16 \rightarrow 16$

Refinement

Refinement on F^2	Hydrogen site location: inferred from neighbouring sites
Least-squares matrix: full	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.037$	$w = 1/[\sigma^2(F_o^2) + (0.0576P)^2 + 0.1837P]$
$wR(F^2) = 0.109$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.04$	$(\Delta/\sigma)_{max} = 0.001$
4261 reflections	$\Delta\rho_{max} = 0.28$ e Å^{-3}
208 parameters	$\Delta\rho_{min} = -0.18$ e Å^{-3}
0 restraints	

Special details

Experimental. IR (KBr) 655 cm^{-1}(C—S), 1243 cm^{-1}(C=S), 1006 cm^{-1}(C—O), 887 cm^{-1} (C—N), 1161 cm^{-1}(C—O—C), 1719 cm^{-1}(C=O). GCMS: m/e: 349. 1H NMR (400 MHz, CDCl$_3$, \?, p.p.m.): 1.65(s, 6H, Piperidine-CH$_2$), 3.15(s, 3H, –OCH$_3$), 3.89(s, 2H, Piperidine-CH$_2$), 4.20(s, 2H Piperidine-CH$_2$), 4.70(d, 2H, Methylene-CH$_2$), 6.45(s, 1H, Ar—H), 7.14(d, 1H, Ar—H), 7.30(s, 1H, Ar—H), 7.50(s,1*H*, Ar—H). Elemental analysis for C$_{17}$H$_{19}$NO$_3$S$_2$: C, 58.36; H, 5.42; N, 3.93.

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.

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 (Å^2)

	x	*y*	*z*	U_{iso}*/U_{eq}
S1	−0.11287 (5)	0.07096 (4)	0.65164 (3)	0.04415 (12)

S2	0.27902 (6)	0.01452 (4)	0.70354 (4)	0.05410 (14)
O3	0.28766 (17)	0.60135 (11)	0.70830 (9)	0.0468 (3)
O4	0.2364 (2)	0.52695 (14)	0.87805 (10)	0.0691 (4)
O5	0.2788 (2)	0.51786 (12)	0.25306 (9)	0.0642 (4)
N6	−0.07603 (19)	−0.11995 (14)	0.78182 (12)	0.0498 (3)
C7	0.2257 (2)	0.49686 (16)	0.78009 (13)	0.0461 (3)
C8	0.1519 (2)	0.36058 (15)	0.73062 (12)	0.0417 (3)
H8	0.1090	0.2881	0.7778	0.050*
C9	0.14239 (19)	0.33355 (13)	0.62055 (11)	0.0348 (3)
C10	0.21349 (18)	0.44638 (13)	0.54632 (11)	0.0331 (3)
C11	0.2836 (2)	0.57753 (14)	0.59446 (11)	0.0366 (3)
C12	0.3533 (2)	0.69027 (14)	0.53032 (13)	0.0432 (3)
H12	0.3988	0.7768	0.5643	0.052*
C13	0.3553 (2)	0.67440 (14)	0.41558 (13)	0.0425 (3)
H13	0.4036	0.7500	0.3721	0.051*
C14	0.2849 (2)	0.54501 (15)	0.36542 (12)	0.0410 (3)
C15	0.2157 (2)	0.43231 (14)	0.43028 (11)	0.0380 (3)
H15	0.1700	0.3460	0.3960	0.046*
C16	0.3336 (3)	0.62831 (19)	0.18046 (14)	0.0585 (4)
H16A	0.3220	0.5938	0.1045	0.088*
H16B	0.4694	0.6806	0.1961	0.088*
H16C	0.2468	0.6854	0.1916	0.088*
C17	0.0638 (2)	0.19172 (14)	0.56997 (12)	0.0415 (3)
H17A	0.1767	0.1563	0.5551	0.050*
H17B	0.0011	0.1976	0.4988	0.050*
C18	0.0326 (2)	−0.02097 (14)	0.72018 (11)	0.0380 (3)
C19	0.0146 (3)	−0.20858 (17)	0.84504 (14)	0.0516 (4)
H19A	0.1589	−0.1777	0.8384	0.062*
H19B	−0.0318	−0.3013	0.8143	0.062*
C20	−0.0410 (3)	−0.2057 (2)	0.96626 (16)	0.0684 (5)
H20A	0.0082	−0.2715	1.0051	0.082*
H20B	0.0233	−0.1160	0.9995	0.082*
C21	−0.2638 (4)	−0.2379 (3)	0.98128 (18)	0.0868 (8)
H21A	−0.3259	−0.3330	0.9601	0.104*
H21B	−0.2915	−0.2246	1.0594	0.104*
C22	−0.3539 (3)	−0.1499 (2)	0.91250 (16)	0.0643 (5)
H22A	−0.3084	−0.0563	0.9413	0.077*
H22B	−0.4984	−0.1805	0.9179	0.077*
C23	−0.2955 (2)	−0.15637 (19)	0.79241 (15)	0.0540 (4)
H23A	−0.3544	−0.2479	0.7612	0.065*
H23B	−0.3471	−0.0940	0.7505	0.065*

Atomic displacement parameters (Å^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0397 (2)	0.03896 (19)	0.0547 (2)	0.01085 (15)	0.00079 (15)	0.01674 (15)
S2	0.0365 (2)	0.0502 (2)	0.0774 (3)	0.01264 (17)	0.01009 (18)	0.0240 (2)
O3	0.0547 (6)	0.0401 (5)	0.0431 (5)	0.0095 (5)	−0.0055 (5)	0.0000 (4)

O4	0.0994 (11)	0.0649 (8)	0.0402 (6)	0.0193 (7)	−0.0079 (6)	−0.0026 (5)
O5	0.0992 (10)	0.0489 (7)	0.0404 (6)	0.0126 (6)	0.0149 (6)	0.0142 (5)
N6	0.0354 (6)	0.0536 (8)	0.0617 (8)	0.0112 (5)	0.0035 (5)	0.0300 (6)
C7	0.0487 (8)	0.0491 (8)	0.0412 (8)	0.0150 (7)	−0.0044 (6)	0.0037 (6)
C8	0.0443 (8)	0.0421 (7)	0.0403 (7)	0.0140 (6)	0.0002 (6)	0.0095 (6)
C9	0.0323 (6)	0.0349 (6)	0.0395 (7)	0.0126 (5)	0.0008 (5)	0.0081 (5)
C10	0.0286 (6)	0.0332 (6)	0.0399 (6)	0.0120 (5)	0.0011 (5)	0.0075 (5)
C11	0.0321 (6)	0.0367 (7)	0.0424 (7)	0.0114 (5)	−0.0018 (5)	0.0047 (5)
C12	0.0397 (7)	0.0326 (7)	0.0560 (8)	0.0075 (6)	−0.0037 (6)	0.0057 (6)
C13	0.0372 (7)	0.0364 (7)	0.0546 (8)	0.0100 (6)	0.0049 (6)	0.0163 (6)
C14	0.0397 (7)	0.0430 (7)	0.0424 (7)	0.0138 (6)	0.0066 (6)	0.0118 (6)
C15	0.0406 (7)	0.0336 (6)	0.0411 (7)	0.0121 (5)	0.0051 (5)	0.0073 (5)
C16	0.0623 (11)	0.0646 (11)	0.0490 (9)	0.0158 (9)	0.0132 (7)	0.0250 (8)
C17	0.0499 (8)	0.0344 (7)	0.0402 (7)	0.0109 (6)	0.0042 (6)	0.0102 (5)
C18	0.0385 (7)	0.0338 (6)	0.0416 (7)	0.0095 (5)	0.0008 (5)	0.0080 (5)
C19	0.0476 (9)	0.0489 (8)	0.0616 (9)	0.0161 (7)	0.0026 (7)	0.0253 (7)
C20	0.0691 (12)	0.0856 (14)	0.0536 (10)	0.0250 (11)	−0.0086 (9)	0.0175 (9)
C21	0.0761 (14)	0.133 (2)	0.0569 (11)	0.0354 (14)	0.0170 (10)	0.0390 (13)
C22	0.0486 (10)	0.0764 (13)	0.0676 (11)	0.0174 (9)	0.0071 (8)	0.0037 (9)
C23	0.0351 (7)	0.0615 (10)	0.0624 (10)	0.0061 (7)	0.0018 (7)	0.0264 (8)

Geometric parameters (Å, º)

S1—C18	1.7837 (14)	C13—H13	0.9300
S1—C17	1.7980 (14)	C14—C15	1.3847 (18)
S2—C18	1.6662 (14)	C15—H15	0.9300
O3—C7	1.3697 (18)	C16—H16A	0.9600
O3—C11	1.3774 (17)	C16—H16B	0.9600
O4—C7	1.2008 (19)	C16—H16C	0.9600
O5—C14	1.3647 (18)	C17—H17A	0.9700
O5—C16	1.4169 (18)	C17—H17B	0.9700
N6—C18	1.3293 (17)	C19—C20	1.507 (3)
N6—C19	1.4722 (18)	C19—H19A	0.9700
N6—C23	1.4758 (19)	C19—H19B	0.9700
C7—C8	1.447 (2)	C20—C21	1.507 (3)
C8—C9	1.3370 (19)	C20—H20A	0.9700
C8—H8	0.9300	C20—H20B	0.9700
C9—C10	1.4565 (17)	C21—C22	1.502 (3)
C9—C17	1.503 (2)	C21—H21A	0.9700
C10—C11	1.3942 (19)	C21—H21B	0.9700
C10—C15	1.3949 (18)	C22—C23	1.502 (3)
C11—C12	1.3803 (19)	C22—H22A	0.9700
C12—C13	1.381 (2)	C22—H22B	0.9700
C12—H12	0.9300	C23—H23A	0.9700
C13—C14	1.389 (2)	C23—H23B	0.9700
C18—S1—C17	104.68 (7)	C9—C17—S1	116.29 (10)
C7—O3—C11	121.57 (11)	C9—C17—H17A	108.2

C14—O5—C16	118.89 (14)	S1—C17—H17A	108.2
C18—N6—C19	122.03 (13)	C9—C17—H17B	108.2
C18—N6—C23	125.13 (12)	S1—C17—H17B	108.2
C19—N6—C23	112.81 (12)	H17A—C17—H17B	107.4
O4—C7—O3	117.03 (15)	N6—C18—S2	124.86 (11)
O4—C7—C8	126.12 (15)	N6—C18—S1	113.21 (10)
O3—C7—C8	116.85 (13)	S2—C18—S1	121.91 (8)
C9—C8—C7	123.36 (13)	N6—C19—C20	110.31 (15)
C9—C8—H8	118.3	N6—C19—H19A	109.6
C7—C8—H8	118.3	C20—C19—H19A	109.6
C8—C9—C10	118.56 (12)	N6—C19—H19B	109.6
C8—C9—C17	122.94 (12)	C20—C19—H19B	109.6
C10—C9—C17	118.49 (11)	H19A—C19—H19B	108.1
C11—C10—C15	117.71 (12)	C21—C20—C19	112.10 (16)
C11—C10—C9	117.80 (12)	C21—C20—H20A	109.2
C15—C10—C9	124.49 (12)	C19—C20—H20A	109.2
O3—C11—C12	116.55 (12)	C21—C20—H20B	109.2
O3—C11—C10	121.83 (12)	C19—C20—H20B	109.2
C12—C11—C10	121.62 (13)	H20A—C20—H20B	107.9
C11—C12—C13	119.90 (13)	C22—C21—C20	112.04 (17)
C11—C12—H12	120.0	C22—C21—H21A	109.2
C13—C12—H12	120.0	C20—C21—H21A	109.2
C12—C13—C14	119.66 (12)	C22—C21—H21B	109.2
C12—C13—H13	120.2	C20—C21—H21B	109.2
C14—C13—H13	120.2	H21A—C21—H21B	107.9
O5—C14—C15	115.35 (13)	C21—C22—C23	110.75 (17)
O5—C14—C13	124.52 (12)	C21—C22—H22A	109.5
C15—C14—C13	120.13 (13)	C23—C22—H22A	109.5
C14—C15—C10	120.97 (13)	C21—C22—H22B	109.5
C14—C15—H15	119.5	C23—C22—H22B	109.5
C10—C15—H15	119.5	H22A—C22—H22B	108.1
O5—C16—H16A	109.5	N6—C23—C22	110.94 (14)
O5—C16—H16B	109.5	N6—C23—H23A	109.5
H16A—C16—H16B	109.5	C22—C23—H23A	109.5
O5—C16—H16C	109.5	N6—C23—H23B	109.5
H16A—C16—H16C	109.5	C22—C23—H23B	109.5
H16B—C16—H16C	109.5	H23A—C23—H23B	108.0

Hydrogen-bond geometry (Å, º)

D—H⋯A	D—H	H⋯A	D⋯A	D—H⋯A
C19—H19*B*⋯O5ⁱ	0.97	2.50	3.410 (2)	157
C23—H23*A*⋯O3ⁱⁱ	0.97	2.60	3.365 (2)	136

Symmetry codes: (i) −*x*, −*y*, −*z*+1; (ii) *x*−1, *y*−1, *z*.