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Dimethyl 2-butyl-2-(3,5-di-*tert*-butyl-4-hydroxybenzyl)malonate

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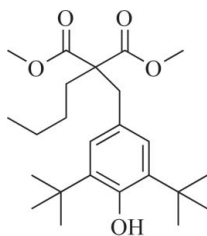
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 Key indicators: single-crystal X-ray study; $T = 294$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.048; wR factor = 0.142; data-to-parameter ratio = 16.5.

The title compound, $\text{C}_{25}\text{H}_{38}\text{O}_5$, was formed by the reaction of dimethyl 2-butylmalonate and 2,6-di-*tert*-butyl-4-[(dimethyl-amino)methyl]phenol. In the crystal structure, molecules are linked by intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds into chains along [010].

Related literature

For background to hindered phenols and hindered amines, see: Denisov (1991); Klemchuk & Gande (1998); Yamazaki & Seguchi (1997); Rasberger (1980); Eggensperger *et al.* (1974, 1976). For a related structure, see: Zeng & Chen (2006).



Experimental

Crystal data

 $\text{C}_{25}\text{H}_{38}\text{O}_5$
 $M_r = 406.54$

 Monoclinic, $P2_1/n$
 $a = 10.854$ (2) Å

 $b = 10.341$ (2) Å

 $c = 22.899$ (5) Å

 $\beta = 98.838$ (4)°

 $V = 2539.7$ (9) Å³
 $Z = 4$

 Mo $K\alpha$ radiation

 $\mu = 0.07$ mm⁻¹
 $T = 294$ K

 $0.24 \times 0.20 \times 0.16$ mm

Data collection

 Bruker SMART CCD
diffractometer

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

 $T_{\min} = 0.983$, $T_{\max} = 0.988$

12778 measured reflections

4475 independent reflections

 2448 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.048$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.048$
 $wR(F^2) = 0.142$
 $S = 1.01$

4475 reflections

272 parameters

12 restraints

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.16$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.18$ e Å⁻³
Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O1}-\text{H1}\cdots\text{O2}^i$	0.82	2.23	2.832 (2)	130

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

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1997); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008) and *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH2893).

References

- Bruker (1997). *SMART* and *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Denisov, E. T. (1991). *Polym. Degrad. Stab.* **34**, 325-332.
- Eggensperger, H., Franzen, V. & Kloss, W. (1974). US Patent No. 3 950 382.
- Eggensperger, H., Franzen, V. & Kloss, W. (1976). US Patent No. 3 856 846.
- Klemchuk, P. P. & Gande, M. E. (1998). *Polym. Degrad. Stab.* **22**, 241-274.
- Rasberger, M. (1980). US Patent No. 4 198 334.
- Sheldrick, G. M. (1996). *SADABS*. University of Göttingen, Germany.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112-122.
- Spek, A. L. (2009). *Acta Cryst.* **D65**, 148-155.
- Yamazaki, T. & Seguchi, T. (1997). *J. Polym. Sci. A*, **35**, 2431-2439.
- Zeng, T. & Chen, L.-G. (2006). *Acta Cryst.* **E62**, o2914-o2915.

supporting information

Acta Cryst. (2009). E65, o2429 [doi:10.1107/S1600536809035697]

Dimethyl 2-butyl-2-(3,5-di-*tert*-butyl-4-hydroxybenzyl)malonate

Tao Zeng and Wan-Zhong Ren

S1. Comment

Hindered phenols are widely used as antioxidants while hindered amines are used as light stabilizers in polymers and lubricants because of their special structures (Denisov, 1991; Klemchuk & Gande, 1998; Yamazaki & Seguchi, 1997). In a former paper (Zeng & Chen, 2006), we reported the crystal structure of bis(1,2,2,6,6-pentamethylpiperidin-4-yl)butylmalonate, a key intermediate in the preparation of 'Tinuvin 144' which is a light stabilizer of the hindered amine class that also contains an oxidant unit of the sterically hindered phenol type (Rasberger, 1980; Eggensperger *et al.*, 1974;1976).

The title compound was prepared as an intermediate when we attempted to synthesize 'Tinuvin 144' by a different route.

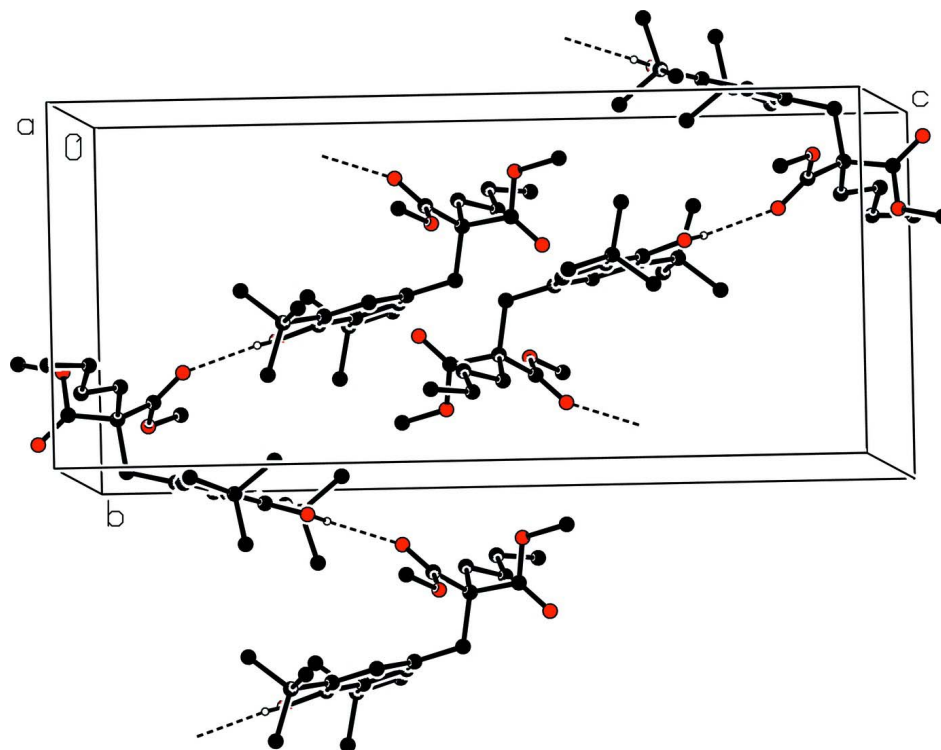
The molecular structure of the title compound (I) is shown in Fig. 1. The phenolic hydroxyl group is sterically hindered by the adjacent bulky *tert*-butyl groups as is indicated by a shorter than normal H...H contact [$H1 \cdots H9B = 1.87 \text{ \AA}$]. In the crystal structure, molecules are linked by intermolecular O-H...O hydrogen bonds into one-dimensional chains along [010] (see Fig. 2).

S2. Experimental

A mixture of bis(1,2,2,6,6-pentamethylpiperidin-4-yl) 2-butylmalonate (11.67 g, 0.025 mol) and 2,6-di-*tert*-butyl-4-((dimethylamino)methyl)phenol (6.59 g, 0.025 mol) was dissolved in toluene (100 ml), stirred and heated to reflux. Then 0.2 g lithium amide was added and stirred for a further 4 h, followed by extraction with ether (30 ml) and drying with anhydrous magnesium sulfate. The solvent was removed by vacuum evaporation at 318 K, and the product was filtered and washed with methanol (10 ml). The title compound (I) (15.05 g) was obtained in 87.9% yield. Suitable crystals (m.p. 420–422 K) were obtained by slow evaporation of a solution of (I) in a mixture of THF and methanol.

S3. Refinement

The H atom of the O—H group was initially located in a difference Fourier map but subsequently included in a calculated position O—H = 0.82 Å and $U_{iso}(H) = 1.5U_{eq}(O)$. All other H atoms were positioned geometrically with C—H distances in the range 0.93–0.97 Å, and they were refined using a riding-model approximation with $U_{iso}(H) = 1.2U_{eq}(C)$ or $1.5U_{eq}(C)$ for methyl H atoms.

**Figure 2**

Part of the crystal structure of (I) with dashed lines indicating O—H...O hydrogen bonds. Only H atoms involved in hydrogen bonds are shown.

Dimethyl 2-butyl-2-(3,5-di-*tert*-butyl-4-hydroxybenzyl)malonate

Crystal data

$C_{24}H_{38}O_5$

$M_r = 406.54$

Monoclinic, $P2_1/n$

Hall symbol: $-P 2_1/n$

$a = 10.854 (2) \text{ \AA}$

$b = 10.341 (2) \text{ \AA}$

$c = 22.899 (5) \text{ \AA}$

$\beta = 98.838 (4)^\circ$

$V = 2539.7 (9) \text{ \AA}^3$

$Z = 4$

$F(000) = 888$

$D_x = 1.063 \text{ Mg m}^{-3}$

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

Cell parameters from 2194 reflections

$\theta = 2.2\text{--}21.1^\circ$

$\mu = 0.07 \text{ mm}^{-1}$

$T = 294 \text{ K}$

Block, colourless

$0.24 \times 0.20 \times 0.16 \text{ mm}$

Data collection

Bruker SMART CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

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

$T_{\min} = 0.983$, $T_{\max} = 0.988$

12778 measured reflections

4475 independent reflections

2448 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.048$

$\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 1.8^\circ$

$h = -11 \rightarrow 12$

$k = -12 \rightarrow 11$

$l = -27 \rightarrow 21$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.048$
 $wR(F^2) = 0.142$
 $S = 1.01$
 4475 reflections
 272 parameters
 12 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
 $w = 1/[\sigma^2(F_o^2) + (0.0654P)^2 + 0.1568P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.16 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.18 \text{ e } \text{\AA}^{-3}$

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.

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 $F^2 > \sigma(F^2)$ 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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.41570 (13)	1.09610 (18)	0.27640 (6)	0.0596 (5)
H1	0.3622	1.1119	0.2971	0.089*
O2	0.11836 (16)	0.68114 (18)	0.10961 (7)	0.0672 (5)
O3	-0.01613 (15)	0.81627 (17)	0.05786 (7)	0.0607 (5)
O4	0.11420 (16)	0.87050 (17)	-0.07043 (7)	0.0645 (5)
O5	0.07880 (19)	0.67112 (18)	-0.04056 (7)	0.0780 (6)
C1	0.35939 (19)	1.0593 (2)	0.22102 (8)	0.0374 (5)
C2	0.44117 (18)	1.0357 (2)	0.18030 (9)	0.0364 (5)
C3	0.3872 (2)	1.0021 (2)	0.12350 (9)	0.0400 (5)
H3	0.4393	0.9863	0.0956	0.048*
C4	0.25997 (19)	0.9908 (2)	0.10623 (8)	0.0366 (5)
C5	0.18367 (19)	1.01343 (19)	0.14827 (8)	0.0362 (5)
H5	0.0978	1.0068	0.1371	0.043*
C6	0.22977 (18)	1.04564 (19)	0.20643 (9)	0.0344 (5)
C7	0.14008 (18)	1.0678 (2)	0.25207 (9)	0.0397 (5)
C8	0.1437 (2)	1.2092 (2)	0.27215 (10)	0.0550 (7)
H8A	0.1170	1.2639	0.2388	0.082*
H8B	0.2273	1.2317	0.2893	0.082*
H8C	0.0892	1.2205	0.3010	0.082*
C9	0.1722 (2)	0.9759 (2)	0.30526 (10)	0.0564 (7)
H9A	0.1136	0.9882	0.3322	0.085*
H9B	0.2548	0.9941	0.3251	0.085*
H9C	0.1681	0.8880	0.2916	0.085*
C10	0.0049 (2)	1.0388 (3)	0.22531 (10)	0.0609 (7)
H10A	-0.0481	1.0514	0.2548	0.091*

H10B	-0.0016	0.9509	0.2118	0.091*
H10C	-0.0205	1.0959	0.1926	0.091*
C11	0.58343 (19)	1.0484 (2)	0.19695 (10)	0.0456 (6)
C12	0.6333 (2)	0.9627 (3)	0.24987 (11)	0.0730 (8)
H12A	0.7217	0.9746	0.2599	0.109*
H12B	0.6158	0.8737	0.2399	0.109*
H12C	0.5936	0.9861	0.2830	0.109*
C13	0.6192 (2)	1.1891 (3)	0.21131 (13)	0.0798 (9)
H13A	0.5885	1.2427	0.1780	0.120*
H13B	0.7083	1.1964	0.2199	0.120*
H13C	0.5831	1.2166	0.2450	0.120*
C14	0.6513 (2)	1.0060 (3)	0.14653 (11)	0.0823 (10)
H14A	0.6255	1.0594	0.1126	0.123*
H14B	0.6314	0.9174	0.1368	0.123*
H14C	0.7396	1.0145	0.1586	0.123*
C15	0.2057 (2)	0.9562 (2)	0.04293 (8)	0.0418 (6)
H15A	0.2616	0.9878	0.0169	0.050*
H15B	0.1268	1.0008	0.0326	0.050*
C16	0.1842 (2)	0.8100 (2)	0.03162 (8)	0.0404 (6)
C17	0.0948 (2)	0.7595 (2)	0.07120 (10)	0.0454 (6)
C18	-0.1102 (3)	0.7799 (3)	0.09247 (13)	0.0861 (10)
H18A	-0.1429	0.6965	0.0800	0.129*
H18B	-0.1762	0.8426	0.0871	0.129*
H18C	-0.0743	0.7765	0.1335	0.129*
C19	0.1224 (2)	0.7911 (2)	-0.03234 (10)	0.0485 (6)
C20	0.0126 (4)	0.6420 (3)	-0.09909 (12)	0.1183 (14)
H20A	-0.0592	0.6970	-0.1075	0.177*
H20B	-0.0136	0.5532	-0.1006	0.177*
H20C	0.0667	0.6565	-0.1279	0.177*
C21	0.3054 (2)	0.7306 (2)	0.04303 (10)	0.0499 (6)
H21A	0.3474	0.7503	0.0825	0.060*
H21B	0.2835	0.6396	0.0423	0.060*
C22	0.3966 (2)	0.7522 (3)	0.00000 (11)	0.0655 (7)
H22A	0.3594	0.7213	-0.0387	0.079*
H22B	0.4109	0.8444	-0.0032	0.079*
C23	0.5202 (3)	0.6857 (3)	0.01765 (13)	0.0813 (9)
H23A	0.5048	0.5961	0.0269	0.098*
H23B	0.5626	0.7259	0.0534	0.098*
C24	0.6056 (3)	0.6890 (3)	-0.02852 (16)	0.1106 (12)
H24A	0.5647	0.6494	-0.0642	0.166*
H24B	0.6808	0.6425	-0.0143	0.166*
H24C	0.6257	0.7771	-0.0364	0.166*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0364 (9)	0.1016 (14)	0.0393 (9)	-0.0034 (9)	0.0012 (7)	-0.0246 (9)
O2	0.0723 (13)	0.0773 (13)	0.0512 (11)	-0.0060 (10)	0.0072 (9)	0.0247 (10)

O3	0.0506 (11)	0.0706 (12)	0.0615 (11)	0.0020 (9)	0.0102 (9)	0.0068 (9)
O4	0.0879 (14)	0.0687 (13)	0.0331 (9)	-0.0081 (10)	-0.0027 (9)	0.0055 (9)
O5	0.1241 (17)	0.0569 (13)	0.0447 (10)	-0.0224 (11)	-0.0135 (10)	-0.0085 (8)
C1	0.0365 (13)	0.0437 (14)	0.0309 (12)	-0.0030 (10)	0.0018 (10)	-0.0057 (10)
C2	0.0344 (12)	0.0369 (13)	0.0373 (13)	-0.0036 (10)	0.0040 (10)	-0.0016 (10)
C3	0.0431 (14)	0.0418 (14)	0.0366 (13)	-0.0012 (10)	0.0107 (11)	-0.0020 (10)
C4	0.0412 (13)	0.0359 (13)	0.0326 (12)	0.0011 (10)	0.0050 (10)	0.0016 (10)
C5	0.0345 (12)	0.0369 (13)	0.0357 (12)	-0.0021 (10)	0.0011 (10)	-0.0015 (10)
C6	0.0342 (13)	0.0347 (13)	0.0346 (12)	-0.0002 (9)	0.0057 (10)	-0.0029 (9)
C7	0.0329 (12)	0.0488 (15)	0.0377 (12)	-0.0001 (10)	0.0065 (10)	-0.0055 (11)
C8	0.0520 (16)	0.0604 (18)	0.0524 (15)	0.0115 (12)	0.0077 (12)	-0.0132 (12)
C9	0.0606 (16)	0.0639 (18)	0.0474 (14)	-0.0046 (13)	0.0171 (12)	-0.0003 (12)
C10	0.0394 (14)	0.089 (2)	0.0565 (16)	-0.0098 (13)	0.0135 (12)	-0.0126 (14)
C11	0.0326 (13)	0.0545 (16)	0.0501 (14)	-0.0056 (11)	0.0082 (11)	-0.0040 (11)
C12	0.0417 (16)	0.089 (2)	0.0850 (19)	0.0067 (14)	0.0009 (14)	0.0169 (16)
C13	0.0532 (18)	0.070 (2)	0.115 (2)	-0.0186 (14)	0.0079 (17)	-0.0108 (17)
C14	0.0392 (16)	0.136 (3)	0.0758 (19)	-0.0085 (16)	0.0207 (14)	-0.0216 (19)
C15	0.0512 (14)	0.0433 (14)	0.0294 (12)	-0.0009 (11)	0.0022 (10)	0.0007 (10)
C16	0.0520 (15)	0.0435 (14)	0.0245 (11)	-0.0007 (11)	0.0024 (10)	0.0021 (10)
C17	0.0540 (16)	0.0446 (15)	0.0348 (13)	-0.0055 (12)	-0.0015 (12)	-0.0046 (11)
C18	0.0618 (19)	0.108 (3)	0.095 (2)	-0.0141 (17)	0.0302 (17)	-0.0006 (19)
C19	0.0607 (17)	0.0496 (17)	0.0345 (14)	-0.0012 (13)	0.0048 (12)	-0.0024 (12)
C20	0.180 (4)	0.099 (3)	0.0592 (19)	-0.035 (2)	-0.034 (2)	-0.0227 (18)
C21	0.0611 (16)	0.0472 (15)	0.0400 (13)	0.0031 (12)	0.0034 (12)	-0.0008 (11)
C22	0.0691 (19)	0.0700 (19)	0.0591 (17)	0.0084 (15)	0.0154 (15)	-0.0021 (14)
C23	0.075 (2)	0.074 (2)	0.099 (2)	0.0108 (16)	0.0258 (18)	-0.0095 (17)
C24	0.093 (3)	0.108 (3)	0.142 (3)	0.013 (2)	0.052 (2)	0.002 (2)

Geometric parameters (Å, °)

O1—C1	1.374 (2)	C12—H12A	0.9600
O1—H1	0.8200	C12—H12B	0.9600
O2—C17	1.195 (3)	C12—H12C	0.9600
O3—C17	1.332 (3)	C13—H13A	0.9600
O3—C18	1.435 (3)	C13—H13B	0.9600
O4—C19	1.191 (3)	C13—H13C	0.9600
O5—C19	1.331 (3)	C14—H14A	0.9600
O5—C20	1.452 (3)	C14—H14B	0.9600
C1—C6	1.402 (3)	C14—H14C	0.9600
C1—C2	1.404 (3)	C15—C16	1.546 (3)
C2—C3	1.386 (3)	C15—H15A	0.9700
C2—C11	1.538 (3)	C15—H15B	0.9700
C3—C4	1.382 (3)	C16—C17	1.518 (3)
C3—H3	0.9300	C16—C19	1.527 (3)
C4—C5	1.383 (3)	C16—C21	1.539 (3)
C4—C15	1.521 (3)	C18—H18A	0.9600
C5—C6	1.389 (3)	C18—H18B	0.9600
C5—H5	0.9300	C18—H18C	0.9600

C6—C7	1.551 (3)	C20—H20A	0.9600
C7—C8	1.531 (3)	C20—H20B	0.9600
C7—C10	1.531 (3)	C20—H20C	0.9600
C7—C9	1.543 (3)	C21—C22	1.517 (3)
C8—H8A	0.9600	C21—H21A	0.9700
C8—H8B	0.9600	C21—H21B	0.9700
C8—H8C	0.9600	C22—C23	1.507 (3)
C9—H9A	0.9600	C22—H22A	0.9700
C9—H9B	0.9600	C22—H22B	0.9700
C9—H9C	0.9600	C23—C24	1.510 (3)
C10—H10A	0.9600	C23—H23A	0.9700
C10—H10B	0.9600	C23—H23B	0.9700
C10—H10C	0.9600	C24—H24A	0.9600
C11—C14	1.526 (3)	C24—H24B	0.9600
C11—C13	1.528 (3)	C24—H24C	0.9600
C11—C12	1.532 (3)		
C1—O1—H1	109.5	H13B—C13—H13C	109.5
C17—O3—C18	116.9 (2)	C11—C14—H14A	109.5
C19—O5—C20	116.0 (2)	C11—C14—H14B	109.5
O1—C1—C6	122.37 (18)	H14A—C14—H14B	109.5
O1—C1—C2	115.09 (18)	C11—C14—H14C	109.5
C6—C1—C2	122.54 (18)	H14A—C14—H14C	109.5
C3—C2—C1	116.58 (19)	H14B—C14—H14C	109.5
C3—C2—C11	121.17 (19)	C4—C15—C16	114.60 (16)
C1—C2—C11	122.24 (18)	C4—C15—H15A	108.6
C4—C3—C2	123.3 (2)	C16—C15—H15A	108.6
C4—C3—H3	118.3	C4—C15—H15B	108.6
C2—C3—H3	118.3	C16—C15—H15B	108.6
C3—C4—C5	117.74 (18)	H15A—C15—H15B	107.6
C3—C4—C15	121.09 (18)	C17—C16—C19	107.60 (18)
C5—C4—C15	121.17 (19)	C17—C16—C21	108.89 (18)
C4—C5—C6	122.81 (19)	C19—C16—C21	109.42 (18)
C4—C5—H5	118.6	C17—C16—C15	109.27 (18)
C6—C5—H5	118.6	C19—C16—C15	108.57 (17)
C5—C6—C1	116.93 (18)	C21—C16—C15	112.95 (18)
C5—C6—C7	120.70 (18)	O2—C17—O3	123.5 (2)
C1—C6—C7	122.36 (17)	O2—C17—C16	126.0 (2)
C8—C7—C10	106.65 (18)	O3—C17—C16	110.5 (2)
C8—C7—C9	111.05 (18)	O3—C18—H18A	109.5
C10—C7—C9	106.34 (18)	O3—C18—H18B	109.5
C8—C7—C6	110.73 (17)	H18A—C18—H18B	109.5
C10—C7—C6	111.32 (16)	O3—C18—H18C	109.5
C9—C7—C6	110.59 (17)	H18A—C18—H18C	109.5
C7—C8—H8A	109.5	H18B—C18—H18C	109.5
C7—C8—H8B	109.5	O4—C19—O5	123.6 (2)
H8A—C8—H8B	109.5	O4—C19—C16	125.9 (2)
C7—C8—H8C	109.5	O5—C19—C16	110.4 (2)

H8A—C8—H8C	109.5	O5—C20—H20A	109.5
H8B—C8—H8C	109.5	O5—C20—H20B	109.5
C7—C9—H9A	109.5	H20A—C20—H20B	109.5
C7—C9—H9B	109.5	O5—C20—H20C	109.5
H9A—C9—H9B	109.5	H20A—C20—H20C	109.5
C7—C9—H9C	109.5	H20B—C20—H20C	109.5
H9A—C9—H9C	109.5	C22—C21—C16	115.95 (19)
H9B—C9—H9C	109.5	C22—C21—H21A	108.3
C7—C10—H10A	109.5	C16—C21—H21A	108.3
C7—C10—H10B	109.5	C22—C21—H21B	108.3
H10A—C10—H10B	109.5	C16—C21—H21B	108.3
C7—C10—H10C	109.5	H21A—C21—H21B	107.4
H10A—C10—H10C	109.5	C23—C22—C21	113.7 (2)
H10B—C10—H10C	109.5	C23—C22—H22A	108.8
C14—C11—C13	107.5 (2)	C21—C22—H22A	108.8
C14—C11—C12	106.2 (2)	C23—C22—H22B	108.8
C13—C11—C12	109.5 (2)	C21—C22—H22B	108.8
C14—C11—C2	111.75 (18)	H22A—C22—H22B	107.7
C13—C11—C2	110.30 (19)	C22—C23—C24	114.8 (3)
C12—C11—C2	111.46 (18)	C22—C23—H23A	108.6
C11—C12—H12A	109.5	C24—C23—H23A	108.6
C11—C12—H12B	109.5	C22—C23—H23B	108.6
H12A—C12—H12B	109.5	C24—C23—H23B	108.6
C11—C12—H12C	109.5	H23A—C23—H23B	107.5
H12A—C12—H12C	109.5	C23—C24—H24A	109.5
H12B—C12—H12C	109.5	C23—C24—H24B	109.5
C11—C13—H13A	109.5	H24A—C24—H24B	109.5
C11—C13—H13B	109.5	C23—C24—H24C	109.5
H13A—C13—H13B	109.5	H24A—C24—H24C	109.5
C11—C13—H13C	109.5	H24B—C24—H24C	109.5
H13A—C13—H13C	109.5		
O1—C1—C2—C3	-178.16 (19)	C1—C2—C11—C12	56.3 (3)
C6—C1—C2—C3	2.2 (3)	C3—C4—C15—C16	92.5 (2)
O1—C1—C2—C11	0.8 (3)	C5—C4—C15—C16	-88.2 (2)
C6—C1—C2—C11	-178.84 (19)	C4—C15—C16—C17	59.5 (2)
C1—C2—C3—C4	-0.3 (3)	C4—C15—C16—C19	176.56 (18)
C11—C2—C3—C4	-179.3 (2)	C4—C15—C16—C21	-61.9 (2)
C2—C3—C4—C5	-0.6 (3)	C18—O3—C17—O2	0.4 (3)
C2—C3—C4—C15	178.8 (2)	C18—O3—C17—C16	-178.90 (19)
C3—C4—C5—C6	-0.5 (3)	C19—C16—C17—O2	126.1 (2)
C15—C4—C5—C6	-179.80 (19)	C21—C16—C17—O2	7.6 (3)
C4—C5—C6—C1	2.2 (3)	C15—C16—C17—O2	-116.2 (3)
C4—C5—C6—C7	-179.01 (19)	C19—C16—C17—O3	-54.6 (2)
O1—C1—C6—C5	177.25 (19)	C21—C16—C17—O3	-173.07 (18)
C2—C1—C6—C5	-3.1 (3)	C15—C16—C17—O3	63.1 (2)
O1—C1—C6—C7	-1.5 (3)	C20—O5—C19—O4	-3.3 (4)
C2—C1—C6—C7	178.14 (19)	C20—O5—C19—C16	177.3 (2)

C5—C6—C7—C8	-113.8 (2)	C17—C16—C19—O4	131.7 (3)
C1—C6—C7—C8	64.8 (2)	C21—C16—C19—O4	-110.1 (3)
C5—C6—C7—C10	4.6 (3)	C15—C16—C19—O4	13.6 (3)
C1—C6—C7—C10	-176.7 (2)	C17—C16—C19—O5	-48.9 (3)
C5—C6—C7—C9	122.6 (2)	C21—C16—C19—O5	69.3 (2)
C1—C6—C7—C9	-58.7 (3)	C15—C16—C19—O5	-167.05 (19)
C3—C2—C11—C14	-6.2 (3)	C17—C16—C21—C22	169.91 (19)
C1—C2—C11—C14	174.9 (2)	C19—C16—C21—C22	52.6 (3)
C3—C2—C11—C13	113.3 (2)	C15—C16—C21—C22	-68.5 (2)
C1—C2—C11—C13	-65.6 (3)	C16—C21—C22—C23	172.5 (2)
C3—C2—C11—C12	-124.8 (2)	C21—C22—C23—C24	171.4 (2)

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>
O1—H1...O2 ⁱ	0.82	2.23	2.832 (2)	130

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