

3-[(2,4-Diethoxyphenyl)(hydroxy)methylene]-1-isopropylpyrrolidine-2,4-dione

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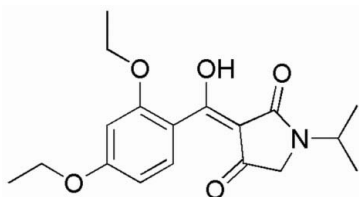
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Key indicators: single-crystal X-ray study; $T = 294$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.052; wR factor = 0.155; data-to-parameter ratio = 16.3.

The title compound, $\text{C}_{18}\text{H}_{23}\text{NO}_5$, is a potential potent new herbicide containing the pyrrolidine-2,4-dione ring system. In the crystalline state, the molecular skeleton contains one enol grouping, which is intramolecularly hydrogen bonded to a neighbouring keto O atom. The dihedral angle between the six-membered ring formed by the enol group and the benzene ring is 41.29 (10)°.

Related literature

For related literature, see: Allen *et al.* (1987); Ellis & Spek (2001); Holzapfel *et al.* (1970); MacKellar *et al.* (1971); Matsuo *et al.* (1980); Rinehart *et al.* (1963); van Rooyen *et al.* (1992); Stickings (1959); Van Der Baan *et al.* (1978); Xu (2005); Zhu, Hu & Yang (2004); Zhu, Song, Li *et al.* (2004); Zhu, Song, Yao *et al.* (2004).



Experimental

Crystal data

$\text{C}_{18}\text{H}_{23}\text{NO}_5$
 $M_r = 333.37$
Monoclinic, $P2_1/n$
 $a = 13.513$ (7) Å
 $b = 8.135$ (4) Å
 $c = 16.312$ (8) Å
 $\beta = 99.518$ (9)°

$V = 1768.5$ (15) Å³
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹
 $T = 294$ (2) K
 $0.28 \times 0.24 \times 0.20$ mm

Data collection

Bruker SMART CCD area-detector diffractometer
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.975$, $T_{\max} = 0.982$
9501 measured reflections
3612 independent reflections
2072 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.052$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.156$
 $S = 1.00$
3612 reflections
222 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.17$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.21$ e Å⁻³

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O3}-\text{H3}\cdots\text{O4}$	0.82	1.74	2.505 (2)	154
$\text{C6}-\text{H6}\cdots\text{O5}$	0.93	2.56	2.994 (3)	109
$\text{C16}-\text{H16}\cdots\text{O4}$	0.98	2.52	2.880 (3)	102

Data collection: SMART (Bruker, 1999); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1999); software used to prepare material for publication: SHELXTL.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SI2023).

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supplementary materials

Acta Cryst. (2007). E63, o3384 [doi:10.1107/S1600536807030759]

3-[(2,4-Diethoxyphenyl)(hydroxy)methylene]-1-isopropylpyrrolidine-2,4-dione

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Comment

Many compounds containing the 3-acylpyrrolidine-2,4-dione system belong to heterocycles with antibiotic activity, such as tenuazonic acid (Stickings, 1959), streptolydigin (Rinehart *et al.*, 1963), tirandamycin (MacKellar *et al.*, 1971), malonomycin (Van Der Baan *et al.*, 1978), α -cyclopiazonic acid (Stickings, 1959; van Rooyen *et al.*, 1992) and β -cyclopiazonic acid (Holzapfel *et al.*, 1970). All these compounds possess a 3-acyltetramic acid grouping as a tricarbonylmethane fragment. Most of the excellent inhibitors of *p*-hydroxyphenylpyruvate dioxygenase also possess similar characteristics, which are crucial for their two kinds of bioactivity (Zhu, Hu & Yang *et al.*, 2004). In order to develop new herbicides, we synthesized the title compound. The molecular structure of the title compound is shown in Fig. 1. Atom H3, involved in intramolecular hydrogen bonding between atoms O3 and O4, was assigned to O3 rather than to O4. The C15=O4 distance is 1.264 (2) Å, which is longer than the normal carbonyl bond length (C13=O1) of 1.227 (3) Å. In contrast, the C11=O3 distance [1.320 (2) Å] is intermediate between a normal carbonyl C=O double bond and a C—O single-bond length (Allen *et al.*, 1987) (Table 1). A similar situation was reported for 3-(1-hydroxyethylidene)-1-phenylpyrrolidine-2,4-dione (Ellis & Spek, 2001), 1-benzyl-3-(α -hydroxybenzylidene)pyrrolidine-2,4-dione, (I) (Zhu, Song, Li *et al.*, 2004), 1-*tert*-butyl-3-(α -hydroxy-4-isopropylbenzylidene)pyrrolidine-2,4-dione, (II) (Xu, 2005), and 3-(α -hydroxyl-2-methoxybenzylidene)-1-isopropylpyrrolidine-2,4-dione, (III) (Zhu, Song, Yao *et al.*, 2004). The dihedral angle formed by the enol ring A with the benzene ring is 41.29 (10)°, which is larger than the dihedral angles for (I), (II) (10 and 21°, respectively) and smaller than the dihedral angle for (III) (53°). The crystal structure of the title compound also involves two weak intramolecular C—H \cdots O hydrogenbonding interactions (Table 2).

Experimental

The title compound was obtained according to the procedure reported by Matsuo *et al.* (1980). Colourless single crystals of the title compound were obtained by recrystallization of 1-isopropyl-3-(α -hydroxy-2,4-diethoxybenzylidene)pyrrolidine-2,4-dione from petroleum ether and ethyl acetate (1:3).

Refinement

The hydroxyl H atom (O3)H3 was found in a difference map and the coordinates were fixed. The other H atoms were placed in calculated positions, with C—H=0.93–0.98 Å, and included in the final cycles of refinement using a riding model, with $U_{\text{iso}}(\text{H})=1.2U_{\text{eq}}(\text{C})$.

Figures

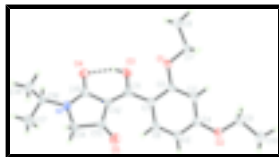


Fig. 1. View of the title compound with 30% probability ellipsoid.

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Crystal data

$C_{18}H_{23}NO_5$

$M_r = 333.37$

Monoclinic, $P2_1/n$

Hall symbol: $-P\ 2_1n$

$a = 13.513\ (7)\ \text{\AA}$

$b = 8.135\ (4)\ \text{\AA}$

$c = 16.312\ (8)\ \text{\AA}$

$\beta = 99.518\ (9)^\circ$

$V = 1768.5\ (15)\ \text{\AA}^3$

$Z = 4$

$F_{000} = 712$

$D_x = 1.252\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation

$\lambda = 0.71073\ \text{\AA}$

Cell parameters from 2383 reflections

$\theta = 2.8\text{--}23.7^\circ$

$\mu = 0.09\ \text{mm}^{-1}$

$T = 294\ (2)\ \text{K}$

Prism, colorless

$0.28 \times 0.24 \times 0.20\ \text{mm}$

Data collection

Bruker SMART CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Monochromator: graphite

$T = 294\ (2)\ \text{K}$

φ and ω scans

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

$T_{\min} = 0.975$, $T_{\max} = 0.982$

9501 measured reflections

3612 independent reflections

2072 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.052$

$\theta_{\max} = 26.4^\circ$

$\theta_{\min} = 1.8^\circ$

$h = -9 \rightarrow 16$

$k = -10 \rightarrow 8$

$l = -18 \rightarrow 20$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.052$

$wR(F^2) = 0.156$

$S = 1.00$

3612 reflections

Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0745P)^2 + 0.2597P]$$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.17\ \text{e \AA}^{-3}$

222 parameters

$$\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$$

Primary atom site location: structure-invariant direct methods

Extinction correction: none

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.71221 (12)	-0.20692 (19)	0.38809 (10)	0.0520 (5)
O2	0.96831 (12)	-0.2405 (2)	0.21363 (10)	0.0562 (5)
O3	0.78362 (14)	-0.06279 (19)	0.53402 (10)	0.0586 (5)
H3	0.7691	-0.0117	0.5737	0.088*
O4	0.76454 (13)	0.16486 (19)	0.63361 (10)	0.0558 (5)
O5	0.88238 (16)	0.3662 (2)	0.39616 (10)	0.0679 (6)
N1	0.81988 (15)	0.4165 (2)	0.59623 (11)	0.0468 (5)
C1	0.85415 (16)	-0.0350 (3)	0.41169 (12)	0.0375 (5)
C2	0.80068 (16)	-0.1612 (3)	0.36515 (13)	0.0390 (5)
C3	0.83646 (17)	-0.2313 (3)	0.29811 (13)	0.0432 (6)
H3A	0.7996	-0.3126	0.2666	0.052*
C4	0.92752 (17)	-0.1795 (3)	0.27832 (13)	0.0415 (5)
C5	0.98245 (17)	-0.0563 (3)	0.32420 (13)	0.0435 (6)
H5	1.0437	-0.0221	0.3112	0.052*
C6	0.94493 (16)	0.0141 (3)	0.38891 (13)	0.0414 (5)
H6	0.9813	0.0978	0.4189	0.050*
C7	0.65284 (19)	-0.3318 (3)	0.34169 (16)	0.0567 (7)
H7A	0.6918	-0.4315	0.3400	0.068*
H7B	0.6299	-0.2952	0.2851	0.068*
C8	0.5653 (2)	-0.3624 (4)	0.38525 (19)	0.0799 (10)
H8A	0.5891	-0.4007	0.4407	0.120*
H8B	0.5223	-0.4440	0.3554	0.120*
H8C	0.5284	-0.2622	0.3876	0.120*
C9	0.9102 (2)	-0.3523 (3)	0.15804 (17)	0.0652 (8)
H9A	0.8444	-0.3063	0.1381	0.078*
H9B	0.9018	-0.4551	0.1862	0.078*
C10	0.9652 (3)	-0.3810 (4)	0.08683 (19)	0.0993 (12)
H10A	0.9695	-0.2798	0.0574	0.149*
H10B	0.9297	-0.4611	0.0499	0.149*
H10C	1.0315	-0.4206	0.1075	0.149*

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C11	0.82072 (16)	0.0394 (3)	0.48406 (13)	0.0397 (5)
C12	0.82908 (16)	0.2029 (3)	0.50607 (12)	0.0383 (5)
C13	0.85927 (18)	0.3479 (3)	0.46522 (14)	0.0468 (6)
C14	0.8572 (2)	0.4900 (3)	0.52583 (14)	0.0521 (6)
H14A	0.9238	0.5352	0.5428	0.062*
H14B	0.8128	0.5766	0.5010	0.062*
C15	0.80136 (16)	0.2574 (3)	0.58414 (13)	0.0416 (6)
C16	0.79723 (19)	0.5096 (3)	0.66811 (14)	0.0510 (6)
H16	0.7915	0.4313	0.7127	0.061*
C17	0.8813 (2)	0.6273 (4)	0.69922 (18)	0.0773 (9)
H17A	0.9433	0.5677	0.7116	0.116*
H17B	0.8677	0.6809	0.7486	0.116*
H17C	0.8863	0.7081	0.6572	0.116*
C18	0.6977 (2)	0.5972 (4)	0.64567 (19)	0.0783 (9)
H18A	0.7012	0.6732	0.6012	0.118*
H18B	0.6829	0.6558	0.6933	0.118*
H18C	0.6457	0.5181	0.6282	0.118*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0456 (9)	0.0513 (10)	0.0616 (10)	-0.0146 (8)	0.0164 (8)	-0.0173 (8)
O2	0.0573 (10)	0.0580 (11)	0.0573 (10)	-0.0122 (9)	0.0220 (9)	-0.0213 (8)
O3	0.0876 (13)	0.0431 (10)	0.0493 (10)	-0.0178 (10)	0.0238 (10)	-0.0030 (8)
O4	0.0748 (12)	0.0498 (10)	0.0489 (9)	-0.0123 (9)	0.0276 (9)	0.0002 (8)
O5	0.1104 (16)	0.0489 (11)	0.0530 (11)	0.0001 (10)	0.0384 (11)	0.0057 (8)
N1	0.0619 (13)	0.0400 (11)	0.0428 (11)	-0.0033 (10)	0.0213 (10)	-0.0041 (9)
C1	0.0396 (12)	0.0346 (12)	0.0370 (12)	-0.0010 (10)	0.0027 (10)	-0.0004 (9)
C2	0.0370 (12)	0.0358 (12)	0.0439 (12)	-0.0028 (10)	0.0056 (10)	-0.0020 (10)
C3	0.0450 (13)	0.0390 (13)	0.0447 (13)	-0.0064 (11)	0.0048 (10)	-0.0084 (10)
C4	0.0464 (13)	0.0398 (13)	0.0385 (12)	0.0009 (11)	0.0075 (10)	-0.0036 (10)
C5	0.0395 (12)	0.0457 (14)	0.0448 (13)	-0.0032 (11)	0.0057 (10)	-0.0027 (11)
C6	0.0434 (13)	0.0379 (12)	0.0408 (12)	-0.0053 (10)	0.0012 (10)	-0.0038 (10)
C7	0.0503 (14)	0.0570 (16)	0.0634 (16)	-0.0183 (13)	0.0112 (13)	-0.0172 (13)
C8	0.0602 (18)	0.091 (2)	0.094 (2)	-0.0343 (17)	0.0280 (17)	-0.0295 (18)
C9	0.0723 (18)	0.0656 (18)	0.0611 (16)	-0.0120 (15)	0.0208 (14)	-0.0276 (14)
C10	0.116 (3)	0.108 (3)	0.086 (2)	-0.037 (2)	0.052 (2)	-0.053 (2)
C11	0.0410 (12)	0.0398 (13)	0.0376 (12)	-0.0057 (10)	0.0047 (10)	0.0028 (10)
C12	0.0418 (12)	0.0383 (12)	0.0350 (11)	-0.0009 (10)	0.0068 (10)	-0.0004 (9)
C13	0.0560 (15)	0.0431 (13)	0.0438 (13)	0.0018 (12)	0.0155 (11)	0.0014 (11)
C14	0.0705 (17)	0.0396 (14)	0.0508 (14)	-0.0038 (12)	0.0237 (13)	-0.0017 (11)
C15	0.0445 (13)	0.0407 (14)	0.0402 (12)	-0.0008 (11)	0.0082 (10)	0.0005 (10)
C16	0.0675 (17)	0.0460 (14)	0.0444 (13)	-0.0033 (13)	0.0236 (13)	-0.0085 (11)
C17	0.0680 (19)	0.095 (2)	0.0686 (18)	-0.0088 (17)	0.0092 (15)	-0.0351 (17)
C18	0.0592 (17)	0.095 (2)	0.087 (2)	0.0036 (17)	0.0316 (16)	-0.0250 (18)

Geometric parameters (\AA , $^\circ$)

O1—C2	1.362 (3)	C8—H8A	0.96
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O1—C7	1.431 (3)	C8—H8B	0.96
O2—C4	1.362 (3)	C8—H8C	0.96
O2—C9	1.425 (3)	C9—C10	1.497 (4)
O3—C11	1.320 (2)	C9—H9A	0.97
O3—H3	0.82	C9—H9B	0.97
O4—C15	1.264 (2)	C10—H10A	0.96
O5—C13	1.227 (3)	C10—H10B	0.96
N1—C15	1.327 (3)	C10—H10C	0.96
N1—C14	1.456 (3)	C11—C12	1.378 (3)
N1—C16	1.470 (3)	C12—C13	1.446 (3)
C1—C6	1.397 (3)	C12—C15	1.455 (3)
C1—C2	1.404 (3)	C13—C14	1.524 (3)
C1—C11	1.463 (3)	C14—H14A	0.97
C2—C3	1.389 (3)	C14—H14B	0.97
C3—C4	1.388 (3)	C16—C17	1.508 (4)
C3—H3A	0.93	C16—C18	1.513 (4)
C4—C5	1.390 (3)	C16—H16	0.98
C5—C6	1.370 (3)	C17—H17A	0.96
C5—H5	0.93	C17—H17B	0.96
C6—H6	0.93	C17—H17C	0.96
C7—C8	1.498 (3)	C18—H18A	0.96
C7—H7A	0.97	C18—H18B	0.96
C7—H7B	0.97	C18—H18C	0.96
C2—O1—C7	119.30 (17)	C9—C10—H10A	109.5
C4—O2—C9	118.34 (18)	C9—C10—H10B	109.5
C11—O3—H3	109.5	H10A—C10—H10B	109.5
C15—N1—C14	111.37 (17)	C9—C10—H10C	109.5
C15—N1—C16	124.19 (18)	H10A—C10—H10C	109.5
C14—N1—C16	124.24 (19)	H10B—C10—H10C	109.5
C6—C1—C2	117.45 (19)	O3—C11—C12	118.06 (19)
C6—C1—C11	119.89 (19)	O3—C11—C1	115.94 (19)
C2—C1—C11	122.59 (19)	C12—C11—C1	125.92 (19)
O1—C2—C3	122.64 (19)	C11—C12—C13	133.2 (2)
O1—C2—C1	116.61 (18)	C11—C12—C15	120.02 (19)
C3—C2—C1	120.7 (2)	C13—C12—C15	106.74 (19)
C4—C3—C2	119.7 (2)	O5—C13—C12	131.0 (2)
C4—C3—H3A	120.1	O5—C13—C14	122.6 (2)
C2—C3—H3A	120.1	C12—C13—C14	106.41 (18)
O2—C4—C3	123.9 (2)	N1—C14—C13	104.27 (18)
O2—C4—C5	115.5 (2)	N1—C14—H14A	110.9
C3—C4—C5	120.6 (2)	C13—C14—H14A	110.9
C6—C5—C4	118.9 (2)	N1—C14—H14B	110.9
C6—C5—H5	120.5	C13—C14—H14B	110.9
C4—C5—H5	120.5	H14A—C14—H14B	108.9
C5—C6—C1	122.6 (2)	O4—C15—N1	124.8 (2)
C5—C6—H6	118.7	O4—C15—C12	124.3 (2)
C1—C6—H6	118.7	N1—C15—C12	110.95 (19)
O1—C7—C8	106.6 (2)	N1—C16—C17	110.5 (2)
O1—C7—H7A	110.4	N1—C16—C18	110.1 (2)

supplementary materials

C8—C7—H7A	110.4	C17—C16—C18	111.9 (2)
O1—C7—H7B	110.4	N1—C16—H16	108.1
C8—C7—H7B	110.4	C17—C16—H16	108.1
H7A—C7—H7B	108.6	C18—C16—H16	108.1
C7—C8—H8A	109.5	C16—C17—H17A	109.5
C7—C8—H8B	109.5	C16—C17—H17B	109.5
H8A—C8—H8B	109.5	H17A—C17—H17B	109.5
C7—C8—H8C	109.5	C16—C17—H17C	109.5
H8A—C8—H8C	109.5	H17A—C17—H17C	109.5
H8B—C8—H8C	109.5	H17B—C17—H17C	109.5
O2—C9—C10	107.6 (2)	C16—C18—H18A	109.5
O2—C9—H9A	110.2	C16—C18—H18B	109.5
C10—C9—H9A	110.2	H18A—C18—H18B	109.5
O2—C9—H9B	110.2	C16—C18—H18C	109.5
C10—C9—H9B	110.2	H18A—C18—H18C	109.5
H9A—C9—H9B	108.5	H18B—C18—H18C	109.5
C7—O1—C2—C3	-0.2 (3)	C1—C11—C12—C13	8.0 (4)
C7—O1—C2—C1	178.3 (2)	O3—C11—C12—C15	2.6 (3)
C6—C1—C2—O1	-179.79 (19)	C1—C11—C12—C15	-174.09 (19)
C11—C1—C2—O1	3.1 (3)	C11—C12—C13—O5	3.6 (5)
C6—C1—C2—C3	-1.3 (3)	C15—C12—C13—O5	-174.5 (3)
C11—C1—C2—C3	-178.3 (2)	C11—C12—C13—C14	-177.0 (2)
O1—C2—C3—C4	-179.7 (2)	C15—C12—C13—C14	4.8 (2)
C1—C2—C3—C4	1.9 (3)	C15—N1—C14—C13	0.1 (3)
C9—O2—C4—C3	6.3 (3)	C16—N1—C14—C13	-174.9 (2)
C9—O2—C4—C5	-172.4 (2)	O5—C13—C14—N1	176.3 (2)
C2—C3—C4—O2	-179.7 (2)	C12—C13—C14—N1	-3.1 (3)
C2—C3—C4—C5	-1.0 (3)	C14—N1—C15—O4	-176.9 (2)
O2—C4—C5—C6	178.33 (19)	C16—N1—C15—O4	-1.9 (4)
C3—C4—C5—C6	-0.4 (3)	C14—N1—C15—C12	3.0 (3)
C4—C5—C6—C1	1.1 (3)	C16—N1—C15—C12	178.0 (2)
C2—C1—C6—C5	-0.2 (3)	C11—C12—C15—O4	-3.5 (3)
C11—C1—C6—C5	176.9 (2)	C13—C12—C15—O4	174.9 (2)
C2—O1—C7—C8	176.1 (2)	C11—C12—C15—N1	176.5 (2)
C4—O2—C9—C10	172.0 (2)	C13—C12—C15—N1	-5.1 (2)
C6—C1—C11—O3	-136.5 (2)	C15—N1—C16—C17	139.1 (2)
C2—C1—C11—O3	40.5 (3)	C14—N1—C16—C17	-46.6 (3)
C6—C1—C11—C12	40.2 (3)	C15—N1—C16—C18	-96.8 (3)
C2—C1—C11—C12	-142.8 (2)	C14—N1—C16—C18	77.5 (3)
O3—C11—C12—C13	-175.4 (2)		

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O3—H3 \cdots O4	0.82	1.74	2.505 (2)	154
C6—H6 \cdots O5	0.93	2.56	2.994 (3)	109
C16—H16 \cdots O4	0.98	2.52	2.880 (3)	102

Fig. 1

