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c-5-Hydroxy-r-2,c-4-bis(methoxycarbonyl)-t-5-methyl-t-3-(3-nitrophenyl)cyclohexanone

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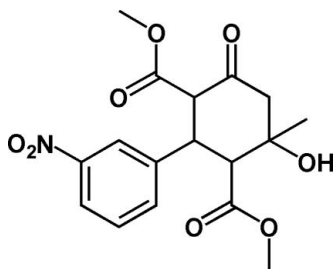
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Key indicators: single-crystal X-ray study; $T = 273$ K; mean $\sigma(\text{C}-\text{C}) = 0.005$ Å; R factor = 0.044; wR factor = 0.112; data-to-parameter ratio = 6.3.

In the title compound, $\text{C}_{17}\text{H}_{19}\text{NO}_8$ [systematic name = dimethyl 4-hydroxy-4-methyl-2-(3-nitrophenyl)-6-oxocyclohexane-1,3-dicarboxylate], the cyclohexanone ring exhibits a chair conformation. The methoxycarbonyl groups are oriented in opposite directions with respect to the cyclohexanone ring. In the crystal, $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds links the molecules into chains running parallel to the a axis. These chains are connected by weak $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds, forming sheets parallel to the ab plane.

Related literature

For the pharmacological activity of cyclohexanone derivatives, see: Puetz *et al.* (2003); Danyi *et al.* (1989); For related structure, see: Hema *et al.* (2006). For conformational analysis, see: Allinger (1977); Cremer & Pople (1975). For graph-set analysis, see: Bernstein *et al.* (1995).



Experimental

Crystal data

$\text{C}_{17}\text{H}_{19}\text{NO}_8$	$V = 1712.3$ (3) Å ³
$M_r = 365.33$	$Z = 4$
Monoclinic, Cc	Mo $K\alpha$ radiation
$a = 20.1842$ (18) Å	$\mu = 0.11$ mm ⁻¹
$b = 5.7380$ (5) Å	$T = 273$ K
$c = 15.5771$ (14) Å	$0.3 \times 0.18 \times 0.15$ mm
$\beta = 108.357$ (1)°	

Data collection

Bruker SMART CCD area-detector diffractometer	1513 independent reflections
7818 measured reflections	1469 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.028$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.044$	2 restraints
$wR(F^2) = 0.112$	H-atom parameters constrained
$S = 1.17$	$\Delta\rho_{\text{max}} = 0.27$ e Å ⁻³
1513 reflections	$\Delta\rho_{\text{min}} = -0.15$ e Å ⁻³
239 parameters	

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C4}-\text{H4}\cdots\text{O2}^i$	0.98	2.46	3.374 (5)	154
$\text{C36}-\text{H36}\cdots\text{O2}^i$	0.93	2.51	3.414 (5)	164
$\text{O8}-\text{H8}\cdots\text{O5}^{ii}$	0.82	2.22	2.969 (5)	152

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

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINTE* (Bruker, 2001); data reduction: *SAINTE*; program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1999); software used to prepare material for publication: *SHELXL97* and *PLATON* (Spek, 2009).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GO2056).

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supplementary materials

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c-5-Hydroxy-r-2,c-4-bis(methoxycarbonyl)-t-5-methyl-t-3-(3-nitrophenyl)cyclohexanone

S. Rizwana Begum, R. Hema, K. Pandiarajan, Sridhar Balasubramanian and A.G. Anitha

Comment

Cyclohexanone derivatives have potent pharmacological activity in the treatment of a broad spectrum of medical conditions (Puetz *et al.*, 2003). Cyclohexanone derivatives penetrate into the stratum corneum and alter the skin permeability of indomethacin by fluidizing or modifying the hard hydrophobic barrier of the corneum (Danyi *et al.*, 1989), thus giving an alternative method of administration of this compound which can cause serious gastric upsets.

In $C_{17}H_{19}NO_8$, (I), (Fig. 1), the cyclohexanone ring adopts a chair conformation [$Q=0.589$ (4) Å, $\theta=173.0$ (4)° and $\varphi=16$ (3)° (Cremer & Pople, 1975)]. The mean value [56.55 (16)°] of the endocyclic torsion angles of the cyclohexanone ring in (I) shows that it is slightly more puckered than the idealized cyclohexanone ring [54.1 (3)°, *MM2* calculation; (Allinger, 1977)]. The two methoxycarbonyl groups at C2 (C4—C3—C2—C21 = -178.4 (2)°) and at C4 (C2—C3—C4—C41 = -179.2 (3)°) are substituted in β -equatorial positions. The nitrophenyl ring (ring A) attached to C3 adopts an α equatorial orientation. The methyl and hydroxyl groups at C5 are oriented in β axial and equatorial positions respectively. The mean planes through C1, C3, C4 and C6 and ring A make a dihedral angle of 82.06 (12)°. This value is greater than that reported for a similar structure (73.76°) (Hema *et al.*, 2006). The dihedral angle between ring A and the carboxy groups O2—C21—O3 and O6—C41—O4 are 61.14 (24)° and 74.71 (27)°, respectively. These two carbonyl groups in (I) are twisted in opposite direction with C5—C4—C41—O7 and C1—C2—C21—O2 torsion angles of 75.8 (5)° and -73.0 (4)°, respectively.

The hydroxyl group forms a strong intermolecular hydrogen bond, O8—H8 \cdots O5(-0.5+x, 0.5+y, z) linking the molecules into C(10) chains, (Bernstein *et al.*, 1995), Table 1 and Figure 2. The weak hydrogen bonds C4—H2 \cdots O2(x, -1+y, z) and C36—H36 \cdots O2(x, -1+y, z) link these chains into sheets which lie parallel to the *ab*-plane.

Experimental

A mixture of methyl acetoacetate (11.6 g, 100 mmol), 3-nitrobenzaldehyde (7.55 g, 50 mmol) and methylamine (1.55 g, 50 mmol) in ethanol (50 ml) was heated to boiling. The reaction mixture was allowed to stand overnight. The separated solid was filtered and purified by recrystallization from ethanol to yield the title compound (yield 11.2 g, 75%, mp 475K).

Refinement

H atoms were treated as riding atoms with C—H (aromatic), 0.93 Å and C—H (aliphatic), 0.98 Å with $U_{iso} = 1.2U_{eq}(C)$ and C—H (methyl), 0.96 Å, O—H, 0.82 Å with $U_{iso} = 1.5U_{eq}(O)$. Friedel pairs were merged.

Computing details

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINTE* (Bruker, 2001); data reduction: *SAINTE* (Bruker, 2001); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97*

(Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1999); software used to prepare material for publication: *SHELXL97* and *PLATON* (Spek, 2009).

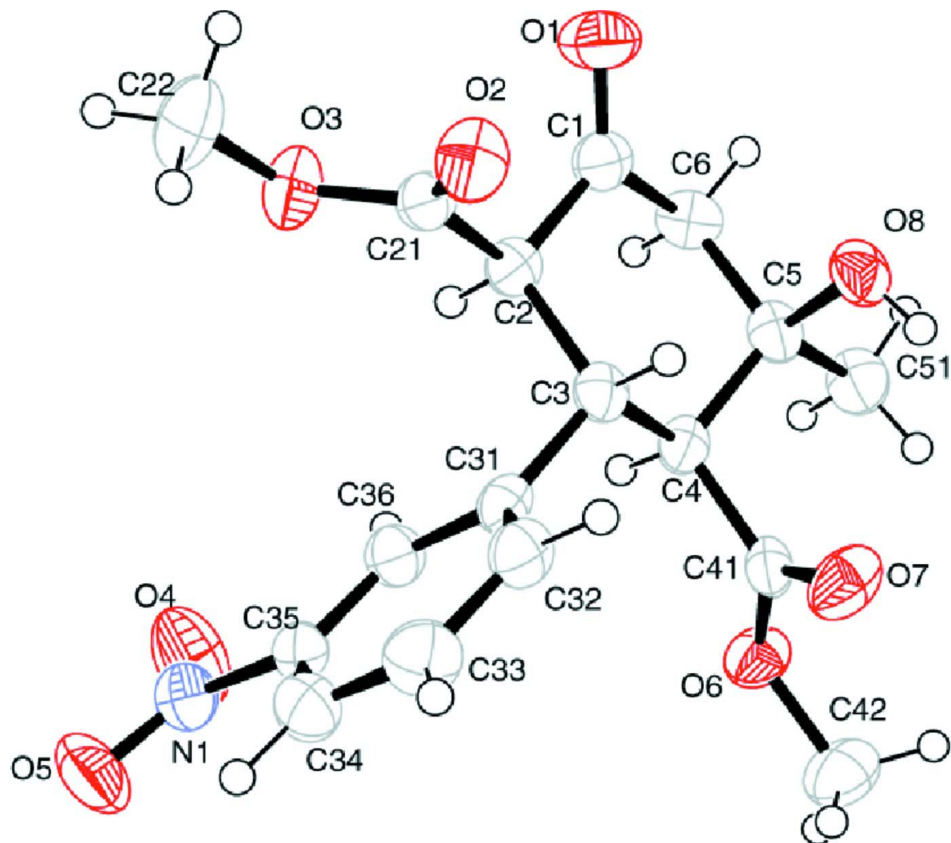


Figure 1

View of the molecule of the title compound showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented by circles of arbitrary radii.

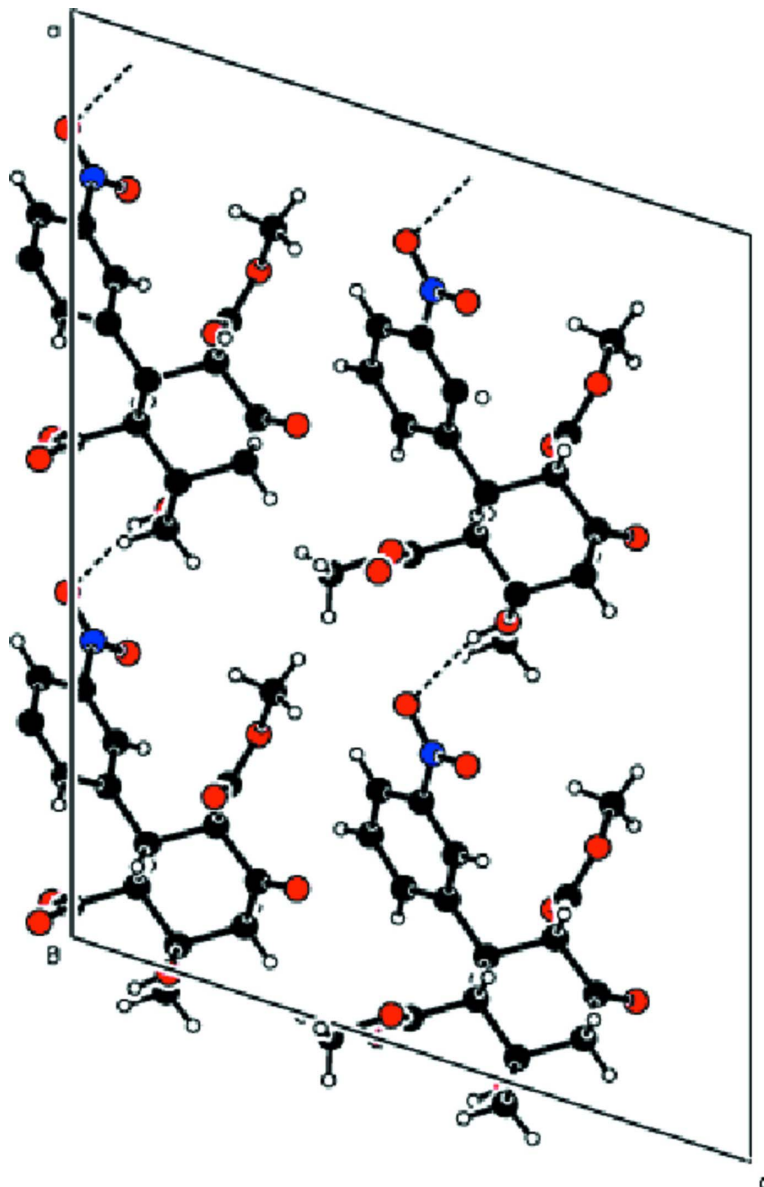


Figure 2

The O8–H8···O5 CHAINS, viewed along *b* axis. Dashed lines indicate hydrogen bonds.

***c*-5-Hydroxy-*r*-2,*c*-4-bis(methoxycarbonyl)-*t*-5-methyl-*t*-3-(3-nitrophenyl)cyclohexanone**

Crystal data

$C_{17}H_{19}NO_8$

$M_r = 365.33$

Monoclinic, *Cc*

Hall symbol: C -2yc

$a = 20.1842$ (18) Å

$b = 5.7380$ (5) Å

$c = 15.5771$ (14) Å

$\beta = 108.357$ (1)°

$V = 1712.3$ (3) Å³

$Z = 4$

$F(000) = 768$

$D_x = 1.417$ Mg m⁻³

Melting point: 475 K

Mo *K*α radiation, $\lambda = 0.71073$ Å

Cell parameters from 2970 reflections

$\theta = 2.8$ – 25 °

$\mu = 0.11$ mm⁻¹

$T = 273$ K

Prism, colourless

$0.3 \times 0.18 \times 0.15$ mm

Data collection

Bruker SMART CCD area-detector diffractometer	1469 reflections with $I > 2\sigma(I)$
Radiation source: fine-focus sealed tube	$R_{\text{int}} = 0.028$
Graphite monochromator	$\theta_{\text{max}} = 25.0^\circ$, $\theta_{\text{min}} = 2.1^\circ$
ω scan	$h = -24 \rightarrow 24$
7818 measured reflections	$k = -6 \rightarrow 6$
1513 independent reflections	$l = -18 \rightarrow 18$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.044$	H-atom parameters constrained
$wR(F^2) = 0.112$	$w = 1/[\sigma^2(F_o^2) + (0.0657P)^2 + 0.665P]$
$S = 1.17$	where $P = (F_o^2 + 2F_c^2)/3$
1513 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
239 parameters	$\Delta\rho_{\text{max}} = 0.27 \text{ e } \text{\AA}^{-3}$
2 restraints	$\Delta\rho_{\text{min}} = -0.15 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

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	$U_{\text{iso}}^*/U_{\text{eq}}$
C3	0.12737 (18)	0.5989 (6)	0.1103 (2)	0.0308 (7)
H3	0.0992	0.7412	0.0959	0.037*
C2	0.17193 (17)	0.6084 (6)	0.2116 (2)	0.0316 (8)
H2	0.2008	0.4675	0.2250	0.038*
C1	0.12635 (19)	0.6086 (6)	0.2729 (2)	0.0370 (8)
C6	0.0735 (2)	0.4139 (7)	0.2538 (3)	0.0400 (9)
H61	0.0438	0.4325	0.2916	0.048*
H62	0.0975	0.2657	0.2686	0.048*
C5	0.02901 (18)	0.4137 (6)	0.1554 (3)	0.0348 (8)
C4	0.07746 (18)	0.3893 (6)	0.0952 (2)	0.0311 (7)
H4	0.1052	0.2467	0.1126	0.037*
C31	0.17510 (18)	0.5939 (6)	0.0524 (2)	0.0312 (7)
C32	0.1713 (2)	0.7685 (7)	-0.0109 (2)	0.0381 (8)
H32	0.1375	0.8839	-0.0197	0.046*
C33	0.2168 (2)	0.7735 (8)	-0.0610 (3)	0.0468 (10)
H33	0.2127	0.8903	-0.1037	0.056*
C34	0.2678 (2)	0.6082 (8)	-0.0482 (3)	0.0444 (9)
H34	0.2992	0.6120	-0.0809	0.053*

C35	0.27115 (19)	0.4360 (7)	0.0145 (3)	0.0380 (8)
C36	0.22569 (18)	0.4240 (6)	0.0639 (2)	0.0361 (8)
H36	0.2289	0.3029	0.1047	0.043*
C41	0.03373 (17)	0.3699 (6)	-0.0021 (2)	0.0317 (7)
C42	-0.0151 (3)	0.1008 (8)	-0.1194 (3)	0.0558 (11)
H421	-0.0626	0.1288	-0.1220	0.084*
H422	-0.0100	-0.0590	-0.1345	0.084*
H423	-0.0032	0.2008	-0.1617	0.084*
C21	0.22025 (18)	0.8146 (6)	0.2304 (2)	0.0335 (8)
C22	0.3365 (2)	0.9385 (9)	0.2956 (4)	0.0607 (13)
H221	0.3412	0.9999	0.2405	0.091*
H222	0.3806	0.8782	0.3326	0.091*
H223	0.3217	1.0603	0.3277	0.091*
C51	-0.0253 (2)	0.2189 (7)	0.1379 (3)	0.0452 (9)
H511	-0.0524	0.2361	0.1782	0.068*
H512	-0.0021	0.0706	0.1479	0.068*
H513	-0.0555	0.2278	0.0765	0.068*
N1	0.32670 (18)	0.2619 (7)	0.0308 (2)	0.0492 (9)
O1	0.13259 (17)	0.7525 (5)	0.3311 (2)	0.0556 (8)
O2	0.20216 (15)	1.0111 (5)	0.2094 (2)	0.0485 (7)
O3	0.28532 (14)	0.7538 (5)	0.2748 (2)	0.0472 (7)
O7	0.00392 (16)	0.5267 (5)	-0.04872 (19)	0.0502 (7)
O8	-0.00399 (14)	0.6359 (4)	0.14196 (19)	0.0415 (6)
H8	-0.0331	0.6410	0.0915	0.062*
O6	0.03014 (14)	0.1479 (4)	-0.03018 (19)	0.0434 (7)
O4	0.32638 (19)	0.1005 (7)	0.0817 (3)	0.0780 (12)
O5	0.37036 (18)	0.2827 (8)	-0.0070 (2)	0.0742 (12)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C3	0.0308 (17)	0.0229 (15)	0.0379 (19)	0.0038 (13)	0.0098 (14)	0.0007 (14)
C2	0.0335 (18)	0.0236 (16)	0.0343 (19)	0.0063 (13)	0.0057 (15)	0.0026 (14)
C1	0.039 (2)	0.0347 (19)	0.035 (2)	0.0031 (16)	0.0090 (16)	0.0009 (17)
C6	0.050 (2)	0.0342 (19)	0.039 (2)	-0.0011 (16)	0.0178 (17)	0.0015 (15)
C5	0.0335 (18)	0.0222 (17)	0.050 (2)	-0.0015 (13)	0.0144 (17)	-0.0028 (14)
C4	0.0281 (16)	0.0228 (16)	0.0383 (19)	0.0020 (13)	0.0045 (14)	-0.0022 (14)
C31	0.0296 (16)	0.0269 (16)	0.0316 (17)	-0.0010 (13)	0.0017 (14)	-0.0028 (13)
C32	0.038 (2)	0.034 (2)	0.0384 (19)	0.0075 (15)	0.0061 (16)	0.0040 (15)
C33	0.054 (2)	0.049 (2)	0.038 (2)	0.0008 (19)	0.0149 (19)	0.0114 (17)
C34	0.039 (2)	0.056 (2)	0.041 (2)	-0.0011 (17)	0.0170 (17)	0.0022 (18)
C35	0.0299 (18)	0.044 (2)	0.0379 (19)	0.0031 (15)	0.0074 (15)	-0.0089 (16)
C36	0.0356 (19)	0.0345 (19)	0.0371 (19)	0.0040 (15)	0.0099 (16)	0.0010 (15)
C41	0.0226 (16)	0.0292 (17)	0.0418 (19)	0.0021 (13)	0.0081 (14)	-0.0025 (15)
C42	0.062 (3)	0.045 (2)	0.050 (3)	-0.0019 (19)	0.004 (2)	-0.012 (2)
C21	0.0342 (18)	0.0328 (19)	0.0324 (18)	0.0013 (15)	0.0090 (14)	-0.0007 (15)
C22	0.042 (2)	0.067 (3)	0.066 (3)	-0.017 (2)	0.007 (2)	0.006 (2)
C51	0.042 (2)	0.036 (2)	0.058 (3)	-0.0063 (16)	0.0160 (19)	-0.0048 (18)
N1	0.0377 (18)	0.060 (2)	0.047 (2)	0.0110 (16)	0.0095 (16)	-0.0060 (17)
O1	0.066 (2)	0.0561 (18)	0.0502 (17)	-0.0150 (15)	0.0260 (15)	-0.0196 (15)

O2	0.0460 (16)	0.0306 (15)	0.0640 (18)	0.0004 (11)	0.0102 (13)	0.0031 (13)
O3	0.0317 (13)	0.0435 (15)	0.0584 (17)	-0.0014 (11)	0.0027 (12)	0.0060 (13)
O7	0.0573 (17)	0.0346 (14)	0.0435 (15)	0.0095 (13)	-0.0060 (13)	0.0020 (12)
O8	0.0380 (14)	0.0319 (13)	0.0539 (16)	0.0078 (11)	0.0135 (12)	-0.0042 (12)
O6	0.0487 (16)	0.0309 (14)	0.0432 (14)	0.0031 (12)	0.0039 (12)	-0.0083 (11)
O4	0.066 (2)	0.067 (2)	0.109 (3)	0.0325 (18)	0.039 (2)	0.023 (2)
O5	0.0514 (19)	0.118 (3)	0.061 (2)	0.033 (2)	0.0295 (17)	0.005 (2)

Geometric parameters (Å, °)

C3—C31	1.513 (5)	C34—C35	1.376 (6)
C3—C4	1.539 (5)	C34—H34	0.9300
C3—C2	1.551 (5)	C35—C36	1.372 (5)
C3—H3	0.9800	C35—N1	1.463 (5)
C2—C21	1.502 (5)	C36—H36	0.9300
C2—C1	1.521 (5)	C41—O7	1.193 (4)
C2—H2	0.9800	C41—O6	1.341 (4)
C1—O1	1.203 (4)	C42—O6	1.428 (5)
C1—C6	1.508 (5)	C42—H421	0.9600
C6—C5	1.513 (5)	C42—H422	0.9600
C6—H61	0.9700	C42—H423	0.9600
C6—H62	0.9700	C21—O2	1.199 (5)
C5—O8	1.423 (4)	C21—O3	1.324 (4)
C5—C51	1.529 (5)	C22—O3	1.444 (5)
C5—C4	1.559 (5)	C22—H221	0.9600
C4—C41	1.499 (5)	C22—H222	0.9600
C4—H4	0.9800	C22—H223	0.9600
C31—C36	1.382 (5)	C51—H511	0.9600
C31—C32	1.392 (5)	C51—H512	0.9600
C32—C33	1.378 (5)	C51—H513	0.9600
C32—H32	0.9300	N1—O5	1.209 (5)
C33—C34	1.368 (6)	N1—O4	1.221 (5)
C33—H33	0.9300	O8—H8	0.8200
C31—C3—C4	113.8 (3)	C32—C33—H33	119.7
C31—C3—C2	109.4 (3)	C33—C34—C35	117.9 (4)
C4—C3—C2	109.0 (3)	C33—C34—H34	121.1
C31—C3—H3	108.2	C35—C34—H34	121.1
C4—C3—H3	108.2	C36—C35—C34	122.8 (3)
C2—C3—H3	108.2	C36—C35—N1	118.7 (3)
C21—C2—C1	111.2 (3)	C34—C35—N1	118.5 (4)
C21—C2—C3	111.0 (3)	C35—C36—C31	119.3 (3)
C1—C2—C3	111.5 (3)	C35—C36—H36	120.3
C21—C2—H2	107.6	C31—C36—H36	120.3
C1—C2—H2	107.6	O7—C41—O6	123.5 (3)
C3—C2—H2	107.6	O7—C41—C4	125.7 (3)
O1—C1—C6	123.8 (3)	O6—C41—C4	110.8 (3)
O1—C1—C2	122.2 (3)	O6—C42—H421	109.5
C6—C1—C2	113.9 (3)	O6—C42—H422	109.5
C1—C6—C5	111.0 (3)	H421—C42—H422	109.5

C1—C6—H61	109.4	O6—C42—H423	109.5
C5—C6—H61	109.4	H421—C42—H423	109.5
C1—C6—H62	109.4	H422—C42—H423	109.5
C5—C6—H62	109.4	O2—C21—O3	123.9 (3)
H61—C6—H62	108.0	O2—C21—C2	124.4 (3)
O8—C5—C6	104.4 (3)	O3—C21—C2	111.8 (3)
O8—C5—C51	110.6 (3)	O3—C22—H221	109.5
C6—C5—C51	110.1 (3)	O3—C22—H222	109.5
O8—C5—C4	110.2 (3)	H221—C22—H222	109.5
C6—C5—C4	109.0 (3)	O3—C22—H223	109.5
C51—C5—C4	112.2 (3)	H221—C22—H223	109.5
C41—C4—C3	111.2 (3)	H222—C22—H223	109.5
C41—C4—C5	109.5 (3)	C5—C51—H511	109.5
C3—C4—C5	110.0 (3)	C5—C51—H512	109.5
C41—C4—H4	108.7	H511—C51—H512	109.5
C3—C4—H4	108.7	C5—C51—H513	109.5
C5—C4—H4	108.7	H511—C51—H513	109.5
C36—C31—C32	118.2 (3)	H512—C51—H513	109.5
C36—C31—C3	121.3 (3)	O5—N1—O4	123.0 (4)
C32—C31—C3	120.4 (3)	O5—N1—C35	119.0 (4)
C33—C32—C31	121.2 (3)	O4—N1—C35	118.0 (3)
C33—C32—H32	119.4	C21—O3—C22	116.5 (3)
C31—C32—H32	119.4	C5—O8—H8	109.5
C34—C33—C32	120.5 (4)	C41—O6—C42	116.7 (3)
C34—C33—H33	119.7		
C31—C3—C2—C21	56.6 (3)	C36—C31—C32—C33	-0.3 (5)
C4—C3—C2—C21	-178.4 (3)	C3—C31—C32—C33	176.6 (4)
C31—C3—C2—C1	-178.8 (3)	C31—C32—C33—C34	-1.1 (6)
C4—C3—C2—C1	-53.8 (4)	C32—C33—C34—C35	1.2 (6)
C21—C2—C1—O1	-3.4 (5)	C33—C34—C35—C36	0.1 (6)
C3—C2—C1—O1	-128.0 (4)	C33—C34—C35—N1	-178.3 (4)
C21—C2—C1—C6	176.5 (3)	C34—C35—C36—C31	-1.5 (6)
C3—C2—C1—C6	52.0 (4)	N1—C35—C36—C31	176.9 (3)
O1—C1—C6—C5	125.7 (4)	C32—C31—C36—C35	1.5 (5)
C2—C1—C6—C5	-54.2 (4)	C3—C31—C36—C35	-175.3 (3)
C1—C6—C5—O8	-59.9 (4)	C3—C4—C41—O7	-46.0 (5)
C1—C6—C5—C51	-178.7 (3)	C5—C4—C41—O7	75.8 (5)
C1—C6—C5—C4	57.9 (4)	C3—C4—C41—O6	136.2 (3)
C31—C3—C4—C41	-56.8 (4)	C5—C4—C41—O6	-102.0 (3)
C2—C3—C4—C41	-179.2 (3)	C1—C2—C21—O2	-72.9 (4)
C31—C3—C4—C5	-178.2 (3)	C3—C2—C21—O2	51.9 (5)
C2—C3—C4—C5	59.4 (3)	C1—C2—C21—O3	105.8 (3)
O8—C5—C4—C41	-70.4 (3)	C3—C2—C21—O3	-129.4 (3)
C6—C5—C4—C41	175.6 (3)	C36—C35—N1—O5	-173.9 (4)
C51—C5—C4—C41	53.3 (4)	C34—C35—N1—O5	4.6 (5)
O8—C5—C4—C3	52.0 (4)	C36—C35—N1—O4	6.6 (6)
C6—C5—C4—C3	-62.0 (4)	C34—C35—N1—O4	-174.9 (4)
C51—C5—C4—C3	175.8 (3)	O2—C21—O3—C22	-1.9 (6)

C4—C3—C31—C36	-65.9 (4)	C2—C21—O3—C22	179.4 (4)
C2—C3—C31—C36	56.3 (4)	O7—C41—O6—C42	-3.4 (5)
C4—C3—C31—C32	117.4 (3)	C4—C41—O6—C42	174.5 (3)
C2—C3—C31—C32	-120.5 (3)		

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>
C4—H4...O2 ⁱ	0.98	2.46	3.374 (5)	154
C36—H36...O2 ⁱ	0.93	2.51	3.414 (5)	164
O8—H8...O5 ⁱⁱ	0.82	2.22	2.969 (5)	152

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