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 (S)-6-Methyl- ϵ -caprolactone

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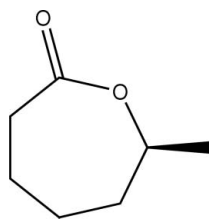
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 Key indicators: single-crystal X-ray study; $T = 150$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.027; wR factor = 0.072; data-to-parameter ratio = 10.7.

The chiral title compound, $\text{C}_7\text{H}_{12}\text{O}_2$, a lactone derivative, features a seven-membered ring that adopts a chair conformation. The crystal structure is stabilized by weak $\text{C}-\text{H}\cdots\text{O}$ interactions occurring in the (100) plane. The absolute configuration was assigned on the basis of the enantioselective synthesis.

Related literature

For related literature, see: van As *et al.* (2005); van Buijtenen *et al.* (2006). For details of the synthesis, see: van As *et al.* (2007). For geometry, see: Cremer & Pople (1975).



Experimental

Crystal data

 $\text{C}_7\text{H}_{12}\text{O}_2$
 $M_r = 128.17$

 Monoclinic, $P2_1$
 $a = 6.757$ (2) Å

 $b = 7.577$ (2) Å

 $c = 7.586$ (2) Å

 $\beta = 110.949$ (13)°

 $V = 362.71$ (17) Å³
 $Z = 2$

 Mo $K\alpha$ radiation

 $\mu = 0.08$ mm⁻¹
 $T = 150$ (2) K

 $0.35 \times 0.15 \times 0.10$ mm

Data collection

Nonius KappaCCD diffractometer

Absorption correction: none

1010 measured reflections

889 independent reflections

 862 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.027$
 $wR(F^2) = 0.072$
 $S = 1.10$

889 reflections

83 parameters

1 restraint

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.12$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.16$ e Å⁻³

Table 1

 Short-contact $\text{C}-\text{H}\cdots\text{O}$ interactions (Å, °) found in the (100) plane.

$\text{C}-\text{H}\cdots\text{O}$	$\text{C}-\text{H}$	$\text{H}\cdots\text{O}$	$\text{C}\cdots\text{O}$	$\text{C}-\text{H}\cdots\text{O}$
$\text{C2}-\text{H2A}\cdots\text{O1}^{\text{i}}$	0.99	2.67	3.573 (2)	152
$\text{C5}-\text{H5A}\cdots\text{O1}^{\text{ii}}$	0.99	2.64	3.616 (2)	166
$\text{C5}-\text{H5B}\cdots\text{O1}^{\text{ii}}$	0.99	2.63	3.601 (2)	168
$\text{C6}-\text{H6}\cdots\text{O1}^{\text{i}}$	1.00	2.54	3.466 (2)	154

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

Data collection: *COLLECT* (Nonius, 1998); cell refinement: *DENZO* (Otwinowski & Minor, 1997); data reduction: *DENZO*; program(s) used to solve structure: *SHELXS86* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *PLATON* and *Mercury* (Macrae *et al.*, 2006).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: TK2248).

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supporting information

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(S)-6-Methyl- ϵ -caprolactone

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S1. Comment

The enantiomers (*R*)- and (*S*)-6-methyl- ϵ -caprolactone (6-MeCL) have been recently used as monomer units for chiral oligomerization and chiral polymerization reactions (van As *et al.*, 2005; van Buijtenen *et al.*, 2006). A new two-step enantioselective synthesis of (*R*)- and (*S*)-6-MeCL from a racemic mixture of 6-MeCL has been described recently (van As *et al.*, 2007). The present paper describes the crystal structure of (*S*)-6-MeCL, (I).

The structure of (I) (Fig. 1) was solved in the non-centrosymmetric space group $P2_1$ with $Z' = 1$. The stereochemistry at the chiral center, C6, was assigned S based on the enantioselective synthesis of (*S*)-6-MeCL, which was reported to yield an enantiomeric excess greater than 99% (van As *et al.*, 2007).

The seven-membered ring (O2/C1—C6) adopts a chair conformation with puckering parameters: $Q_2 = 0.453(2) \text{ \AA}$, $\varphi_2 = 130.6(2)^\circ$, $Q_3 = 0.653(2) \text{ \AA}$, $\varphi_3 = 102.6(2)^\circ$ and with a total puckering amplitude $Q = 0.795(2) \text{ \AA}$ (Cremer & Pople, 1975).

The crystal structure (I) is stabilized by weak C—H \cdots O interactions (Table 1) with the four shortest contacts involving the O1 atom. These short contacts occur between molecules in the (1 0 0) plane, Fig. 2.

S2. Experimental

Details about the synthesis of (*S*)-6-methyl- ϵ -caprolactone have been given in a previous paper (van As *et al.*, 2007).

S3. Refinement

In the absence of significant anomalous scattering effects, XXX Friedel pairs were merged prior to the refinement. The H atoms were found in difference Fourier maps and subsequently placed at calculated positions with C—H = 0.99–1.00 \AA , and with $U_{\text{iso}}(\text{H}) = 1.2$ or 1.5 times $U_{\text{eq}}(\text{carrier C})$.

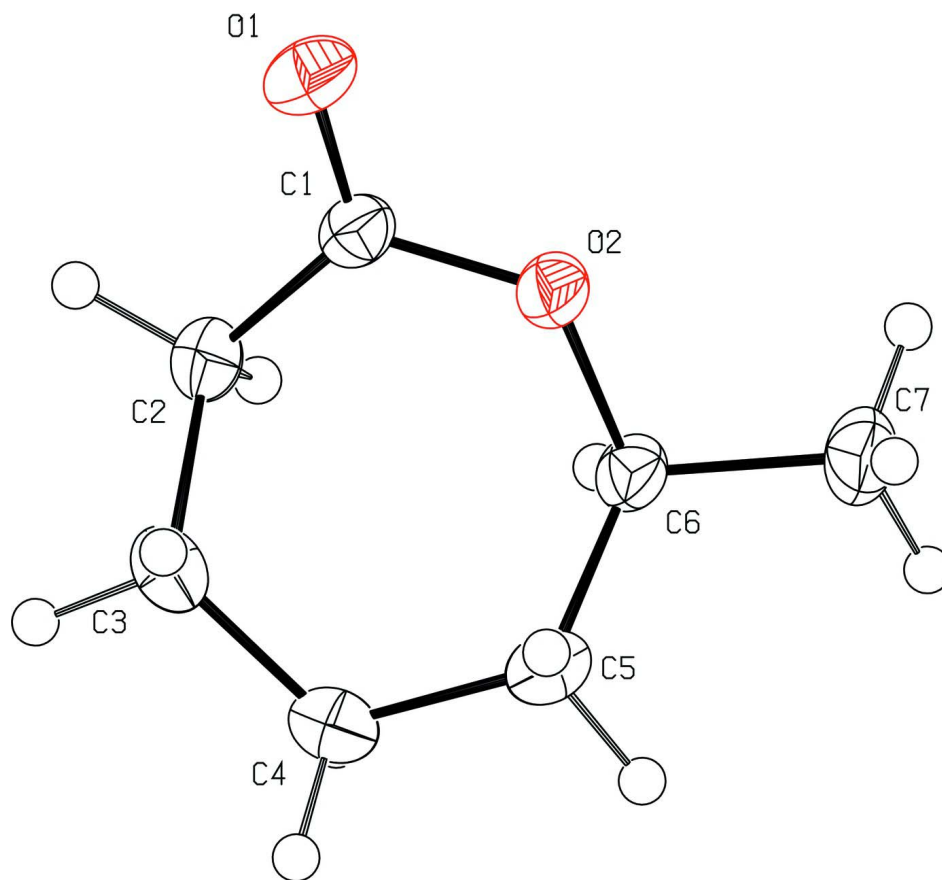
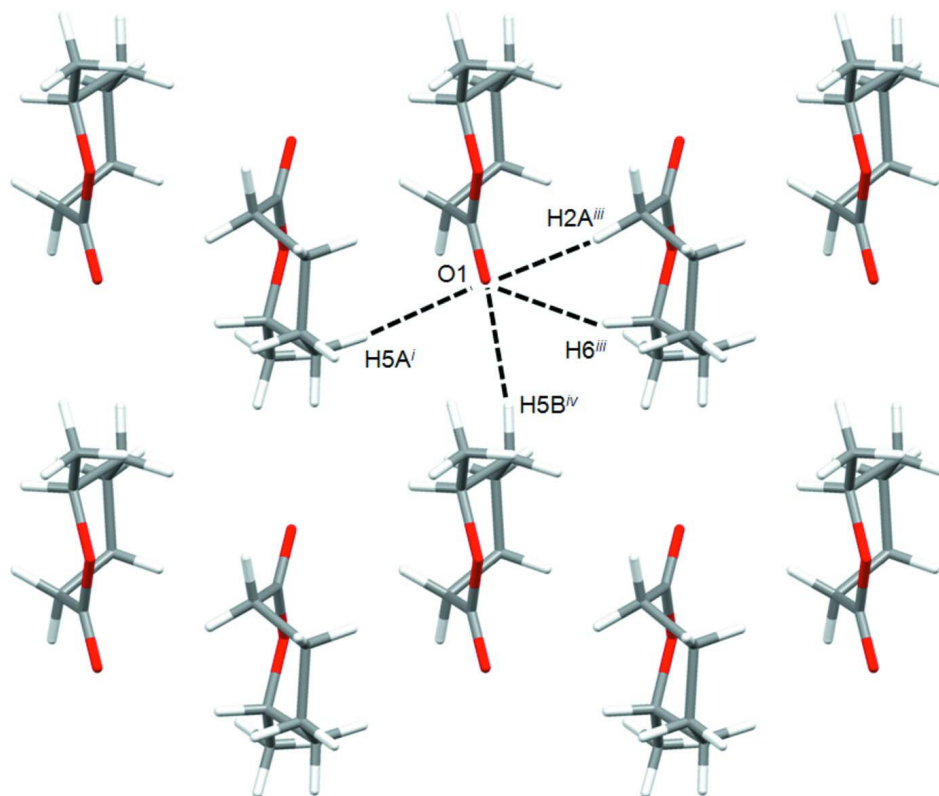


Figure 1

Molecular structure of (I) showing atom labelling and displacement ellipsoids at the 50% probability level.

**Figure 2**

The packing of one layer found in the crystal structure of (I) viewed down the a^* direction. Molecules are connected by short C–H \cdots O contacts in the (1 0 0) plane. Only one set of short contacts (dashed lines) is shown for clarity. The symmetry codes are: (i) $2 - x, -1/2 + y, -z$; (iii) $2 - x, 1/2 + y, -z$; (iv) $x, y, z - 1$.

(S)-3-methyl-2-oxepanone*Crystal data*

$C_7H_{12}O_2$
 $M_r = 128.17$
 Monoclinic, $P2_1$
 Hall symbol: P 2yb
 $a = 6.757(2) \text{ \AA}$
 $b = 7.577(2) \text{ \AA}$
 $c = 7.586(2) \text{ \AA}$
 $\beta = 110.949(13)^\circ$
 $V = 362.71(17) \text{ \AA}^3$
 $Z = 2$

$F(000) = 140$
 $D_x = 1.174 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
 Cell parameters from 7835 reflections
 $\theta = 1.0\text{--}27.5^\circ$
 $\mu = 0.08 \text{ mm}^{-1}$
 $T = 150 \text{ K}$
 Prism, colourless
 $0.35 \times 0.15 \times 0.10 \text{ mm}$

Data collection

Nonius KappaCCD
 diffractometer
 Radiation source: rotating anode
 Graphite monochromator
 φ and ω scans
 10010 measured reflections
 889 independent reflections

862 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$
 $\theta_{\text{max}} = 27.5^\circ$, $\theta_{\text{min}} = 3.5^\circ$
 $h = -8 \rightarrow 8$
 $k = -9 \rightarrow 9$
 $l = -9 \rightarrow 9$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: difference Fourier map
$R[F^2 > 2\sigma(F^2)] = 0.027$	H-atom parameters constrained
$wR(F^2) = 0.072$	$w = 1/[\sigma^2(F_o^2) + (0.0431P)^2 + 0.0282P]$
$S = 1.10$	where $P = (F_o^2 + 2F_c^2)/3$
889 reflections	$(\Delta/\sigma)_{\max} < 0.001$
83 parameters	$\Delta\rho_{\max} = 0.12 \text{ e } \text{\AA}^{-3}$
1 restraint	$\Delta\rho_{\min} = -0.16 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	Absolute structure: known chirality of atom C6(S)

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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	1.03545 (16)	0.06828 (15)	-0.16671 (13)	0.0355 (3)
O2	1.11573 (13)	0.04935 (14)	0.13758 (12)	0.0288 (2)
C1	0.9754 (2)	0.03155 (18)	-0.03896 (16)	0.0258 (3)
C2	0.7534 (2)	-0.0285 (2)	-0.07029 (18)	0.0301 (3)
H2A	0.7594	-0.1422	-0.0042	0.036*
H2B	0.6778	-0.0491	-0.2069	0.036*
C3	0.6274 (2)	0.1054 (2)	0.0003 (2)	0.0356 (3)
H3A	0.6603	0.2261	-0.0311	0.043*
H3B	0.4740	0.0849	-0.0669	0.043*
C4	0.6759 (2)	0.0938 (2)	0.2120 (2)	0.0365 (3)
H4A	0.6389	-0.0260	0.2421	0.044*
H4B	0.5841	0.1787	0.2461	0.044*
C5	0.9066 (2)	0.1315 (2)	0.33415 (18)	0.0303 (3)
H5A	0.9423	0.2528	0.3071	0.036*
H5B	0.9205	0.1274	0.4685	0.036*
C6	1.06592 (19)	0.00512 (19)	0.30521 (16)	0.0267 (3)
H6	1.0098	-0.1181	0.2938	0.032*
C7	1.2785 (2)	0.0148 (3)	0.46522 (19)	0.0403 (4)
H7A	1.3333	0.1356	0.4759	0.060*
H7B	1.3783	-0.0660	0.4396	0.060*
H7C	1.2615	-0.0190	0.5837	0.060*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0442 (5)	0.0406 (6)	0.0260 (5)	0.0001 (5)	0.0177 (4)	0.0018 (4)
O2	0.0248 (4)	0.0400 (6)	0.0219 (4)	-0.0028 (4)	0.0087 (3)	0.0022 (4)
C1	0.0305 (6)	0.0245 (6)	0.0221 (6)	0.0029 (5)	0.0093 (5)	-0.0007 (5)
C2	0.0272 (6)	0.0341 (7)	0.0255 (6)	-0.0021 (5)	0.0050 (5)	-0.0041 (6)
C3	0.0242 (6)	0.0431 (9)	0.0365 (7)	0.0058 (6)	0.0074 (5)	0.0002 (7)
C4	0.0298 (6)	0.0451 (9)	0.0382 (7)	0.0031 (6)	0.0166 (5)	-0.0030 (7)
C5	0.0360 (7)	0.0322 (7)	0.0254 (6)	-0.0015 (6)	0.0143 (5)	-0.0026 (5)
C6	0.0265 (6)	0.0330 (7)	0.0202 (6)	-0.0027 (5)	0.0079 (5)	0.0028 (5)
C7	0.0311 (6)	0.0598 (11)	0.0252 (6)	-0.0040 (7)	0.0043 (5)	0.0070 (7)

Geometric parameters (Å, °)

O1—C1	1.2095 (16)	C4—H4A	0.9900
O2—C1	1.3428 (15)	C4—H4B	0.9900
O2—C6	1.4659 (14)	C5—C6	1.5140 (19)
C1—C2	1.5028 (18)	C5—H5A	0.9900
C2—C3	1.539 (2)	C5—H5B	0.9900
C2—H2A	0.9900	C6—C7	1.5152 (18)
C2—H2B	0.9900	C6—H6	1.0000
C3—C4	1.523 (2)	C7—H7A	0.9800
C3—H3A	0.9900	C7—H7B	0.9800
C3—H3B	0.9900	C7—H7C	0.9800
C4—C5	1.5288 (19)		
C1—O2—C6	122.88 (10)	C5—C4—H4B	108.6
O1—C1—O2	117.15 (12)	H4A—C4—H4B	107.6
O1—C1—C2	123.02 (11)	C6—C5—C4	114.69 (12)
O2—C1—C2	119.82 (11)	C6—C5—H5A	108.6
C1—C2—C3	112.99 (12)	C4—C5—H5A	108.6
C1—C2—H2A	109.0	C6—C5—H5B	108.6
C3—C2—H2A	109.0	C4—C5—H5B	108.6
C1—C2—H2B	109.0	H5A—C5—H5B	107.6
C3—C2—H2B	109.0	O2—C6—C5	112.13 (11)
H2A—C2—H2B	107.8	O2—C6—C7	103.76 (10)
C4—C3—C2	113.05 (12)	C5—C6—C7	111.89 (12)
C4—C3—H3A	109.0	O2—C6—H6	109.6
C2—C3—H3A	109.0	C5—C6—H6	109.6
C4—C3—H3B	109.0	C7—C6—H6	109.6
C2—C3—H3B	109.0	C6—C7—H7A	109.5
H3A—C3—H3B	107.8	C6—C7—H7B	109.5
C3—C4—C5	114.64 (11)	H7A—C7—H7B	109.5
C3—C4—H4A	108.6	C6—C7—H7C	109.5
C5—C4—H4A	108.6	H7A—C7—H7C	109.5
C3—C4—H4B	108.6	H7B—C7—H7C	109.5

C6—O2—C1—O1	-178.35 (12)	C3—C4—C5—C6	-61.49 (19)
C6—O2—C1—C2	2.78 (18)	C1—O2—C6—C5	-68.78 (16)
O1—C1—C2—C3	-112.90 (16)	C1—O2—C6—C7	170.29 (13)
O2—C1—C2—C3	65.90 (16)	C4—C5—C6—O2	80.45 (14)
C1—C2—C3—C4	-81.25 (16)	C4—C5—C6—C7	-163.43 (12)
C2—C3—C4—C5	61.39 (19)		
