organic compounds

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(2RS,8aRS)-6-Oxo-1,2,3,4,6,7,8,8a-octahydronaphthalene-2-carboxylic acid

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Key indicators: single-crystal X-ray study; T = 100 K; mean σ (C–C) = 0.002 Å; R factor = 0.035; wR factor = 0.088; data-to-parameter ratio = 13.1.

The title racemate, $C_{11}H_{14}O_3$, aggregates in the crystal structure as acid-to-ketone O-H···O hydrogen-bonding catemers whose components are glide-related. The relative stereochemistry at the carboxyl group arises spontaneously during the synthesis. Two intermolecular $C-H \cdots O = C$ close contacts were found, both involving the acid group.

Related literature

For background information, see: Borthwick (1980). For synthetic details see: Finnegan & Bachman (1965); House et al. (1965). For information on weak hydrogen bonds, see: Steiner (1997).



Experimental

Crystal data C11H14O3 $M_r = 194.22$

Monoclinic, $P2_1/c$ a = 6.2315 (11) Å

b = 9.2296 (16) Å
c = 17.234 (3) Å
$\beta = 93.366 \ (3)^{\circ}$
V = 989.5 (3) Å ³
Z = 4

Data collection

Bruker SMART APEXII CCD
area-detector diffractometer
Absorption correction: multi-scan
(SADABS; Sheldrick, 2001)
$T_{\rm min} = 0.768, T_{\rm max} = 0.849$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.035$	H atoms treated by a mixture of
$wR(F^2) = 0.088$	independent and constrained
S = 1.09	refinement
1719 reflections	$\Delta \rho_{\rm max} = 0.22 \text{ e} \text{ Å}^{-3}$
131 parameters	$\Delta \rho_{\rm min} = -0.20 \text{ e } \text{\AA}^{-3}$

Cu $K\alpha$ radiation $\mu = 0.77 \text{ mm}^{-1}$

 $0.36 \times 0.31 \times 0.22$ mm

7466 measured reflections 1719 independent reflections

1684 reflections with $I > 2\sigma(I)$

T = 100 (2) K

 $R_{\rm int} = 0.028$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$\begin{array}{c} O3-H3\cdots O1^{i}\\ C2-H2\cdots O2^{ii}\\ C7-H7A\cdots O2^{iii} \end{array}$	0.888 (19) 1.00 0.99	1.79 (2) 2.40 2.47	2.6797 (13) 3.3191 (15) 3.3708 (15)	174.8 (17) 152 151
Symmetry codes: $x, -y + \frac{3}{2}, z + \frac{1}{2}$.	(i) $x + 1, -y$	$+\frac{3}{2}, z-\frac{1}{2};$ (ii)	$-x+1, y-\frac{1}{2},$	$-z + \frac{1}{2};$ (iii)

Data collection: APEX2 (Bruker, 2006); cell refinement: SAINT (Bruker, 2005); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH2720).

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(2RS,8aRS)-6-Oxo-1,2,3,4,6,7,8,8a-octahydronaphthalene-2-carboxylic acid

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Comment

Among ketocarboxylic acids, we have shown that the usually dominant dimerization can be disfavored by lowering molecular flexibility, as measured by the number of fully rotatable bonds present. Typically this results in increased occurrence of acid-to-ketone catemers, whose occurrence is also favored by fixed "anti-like" arrangements, in which carboxyl and ketone are aimed in opposite directions. In this context, we report here the title compound, (I), whose structure conforms to both of the above criteria.

Fig. 1 shows the asymmetric unit, whose only conformational options lie in the carboxyl side-chain, which is oriented $[C1-C2-C9-O2 \text{ torsion angle} = -37.96 (15)^{\circ}]$ so as to minimize steric interactions with H atoms at C1 and C3.

The disordering of C—O bond lengths and C—C—O angles often seen in carboxyl dimers becomes impossible when the H-bonding mode precludes the required averaging mechanisms. Because (I) is not dimeric the distances and angles here are fully ordered and thus typical of those in highly ordered dimeric carboxyls (Borthwick, 1980).

Fig. 2 shows the packing of the cell, with extra molecules included to illustrate the acid-to-ketone H-bonding scheme. Each carboxylic acid is linked to the ketone in a molecule glide related in the c direction. Glide relationships for intra-chain units in catemers is far less common than screw-related schemes. Each of the four molecules in the chosen cell participates in a separate H-bonding chain and these pass through the cell in counterdirectional pairs related by centrosymmetry, with the chains advancing by one cell in a and one-half cell in c for each H bond.

We characterize the geometry of H bonding to carbonyls using a combination of the H···O=C angle and the H···O=C—C torsion angle. These describe the approach of the H atom to the receptor O in terms of its deviation from, respectively, C=O axiality (ideal = 120°) and planarity with the carbonyl (ideal = 0°). In (I), these angles are 131.0 (6) & 0.6 (8)°.

Within the 2.6 Å range we standardly survey for C—H···O packing interactions (Steiner, 1997), two intermolecular close contacts were found, both involving O2, the carboxyl carbonyl (see table).

Experimental

Compound (I) was synthesized by the method of Finnegan & Bachman (1965); crystallization from ethyl acetate yielded material suitable for X-ray, mp 418 K. The C2/C8a stereochemistry clearly represents the stabler of the two epimers possible and probably arises as the result of equilibrations during the synthesis (House *et al.*, 1965).

The solid-state (KBr) infrared spectrum of (I) has C=O absorptions at 1721 & 1640 cm⁻¹, with a peak separation typical of the shifts seen in catemers, due, respectively, to removal of H bonding from the acid C=O and addition of H bonding to the ketone; an alkene peak appears at 1616 cm⁻¹. In CHCl₃ solution, where dimers predominate, these bands appear, respectively, at 1708, 1666 and 1622 cm⁻¹.

Refinement

All H atoms for (I) were found in electron-density difference maps. The positional parameters for the carboxyl H were allowed to refine but the $U_{iso}(H)$ was held at $1.5U_{eq}(O)$. The methylene, methine and vinyl Hs were placed in geometrically idealized positions and constrained to ride on their parent C atoms with C–H distances of 0.99, 1.00 & 0.95 Å, respectively, and $U_{iso}(H) = 1.2U_{eq}(C)$.

Figures



Fig. 1. The asymmetric unit of (I), with its numbering. Displacement ellipsoids are drawn at the 40% probability level.



Fig. 2. A partial packing diagram with extracellular molecules, illustrating the centrosymmetrically related pairs of acid-to-ketone H-bonding chains passing through the cell. All carbonbound H atoms are removed for clarity. Displacement ellipsoids are drawn at the 40% probability level.

(2RS,8aRS)-6-Oxo-1,2,3,4,6,7,8,8a-octahydronaphthalene-2- carboxylic acid

$F_{000} = 416$
$D_{\rm x} = 1.304 {\rm Mg m}^{-3}$
Melting point: 418 K
Cu K α radiation $\lambda = 1.54178$ Å
Cell parameters from 7006 reflections
$\theta = 4.8 - 67.1^{\circ}$
$\mu = 0.77 \text{ mm}^{-1}$
T = 100 (2) K
Parallelepiped, colourless
$0.36 \times 0.31 \times 0.22 \text{ mm}$

Data collection

Bruker SMART CCD APEXII area-detector diffractometer	1719 independent reflections
Radiation source: fine-focus sealed tube	1684 reflections with $I > 2\sigma(I)$
Monochromator: graphite	$R_{\rm int} = 0.028$
T = 100(2) K	$\theta_{\text{max}} = 67.3^{\circ}$
ϕ and ω scans	$\theta_{\min} = 5.1^{\circ}$
Absorption correction: multi-scan (<i>SADABS</i> ; Sheldrick, 2001)	$h = -7 \rightarrow 7$
$T_{\min} = 0.769, \ T_{\max} = 0.849$	$k = -11 \rightarrow 10$

7466 measured reflections	$l = -20 \rightarrow 20$
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Refinement

Refinement on F^2	Hydrogen site location: inferred from neighbouring sites
Least-squares matrix: full	H atoms treated by a mixture of independent and constrained refinement
$R[F^2 > 2\sigma(F^2)] = 0.035$	$w = 1/[\sigma^2(F_o^2) + (0.0417P)^2 + 0.4167P]$ where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.088$	$(\Delta/\sigma)_{max} < 0.001$
<i>S</i> = 1.09	$\Delta \rho_{max} = 0.22 \text{ e} \text{ Å}^{-3}$
1719 reflections	$\Delta \rho_{min} = -0.20 \text{ e } \text{\AA}^{-3}$
131 parameters	Extinction correction: <i>SHELXTL</i> (Sheldrick, 2004), Fc [*] =kFc[1+0.001xFc ² λ^3 /sin(2 θ)] ^{-1/4}
Primary atom site location: structure-invariant direct	

Primary atom site location: structure-invariant direct methods Extinction coefficient: 0.0064 (8)

Secondary atom site location: difference Fourier map

Special details

Experimental. crystal mounted on a Cryoloop using Paratone-N

Geometry. All e.s.d.'s (except for 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 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 (A^2)

	x	У	Z	$U_{\rm iso}*/U_{\rm eq}$
01	-0.28896 (14)	0.72032 (11)	0.52340 (5)	0.0292 (3)
C1	0.36320 (19)	0.76724 (13)	0.32544 (7)	0.0191 (3)
H1A	0.2697	0.8459	0.3038	0.023*
H1B	0.5053	0.8100	0.3405	0.023*
O2	0.41479 (13)	0.84304 (9)	0.16676 (5)	0.0217 (2)
C2	0.39130 (18)	0.65372 (13)	0.26199 (7)	0.0176 (3)
H2	0.4969	0.5788	0.2816	0.021*
O3	0.61028 (15)	0.64301 (10)	0.15353 (5)	0.0281 (3)
H3	0.652 (3)	0.689 (2)	0.1116 (11)	0.042*
C3	0.17383 (19)	0.58079 (14)	0.24029 (7)	0.0199 (3)
H3A	0.0716	0.6537	0.2178	0.024*
H3B	0.1932	0.5052	0.2006	0.024*
C4	0.08193 (19)	0.51220 (13)	0.31220 (7)	0.0200 (3)

-0.0631	0.4735	0.2978	0.024*
0.1744	0.4299	0.3297	0.024*
0.06671 (19)	0.61740 (13)	0.37828 (7)	0.0175 (3)
-0.11075 (19)	0.62502 (14)	0.41883 (7)	0.0198 (3)
-0.2346	0.5720	0.4009	0.024*
-0.1206 (2)	0.71159 (14)	0.48942 (7)	0.0210 (3)
0.0842 (2)	0.78214 (14)	0.52003 (7)	0.0223 (3)
0.1681	0.7127	0.5534	0.027*
0.0498	0.8668	0.5523	0.027*
0.2190 (2)	0.83140 (14)	0.45355 (7)	0.0219 (3)
0.1415	0.9091	0.4239	0.026*
0.3566	0.8720	0.4754	0.026*
0.26523 (18)	0.70674 (13)	0.39857 (7)	0.0181 (3)
0.3744	0.6421	0.4257	0.022*
0.47226 (18)	0.72473 (13)	0.19025 (7)	0.0176 (3)
	$\begin{array}{c} -0.0631\\ 0.1744\\ 0.06671(19)\\ -0.11075(19)\\ -0.2346\\ -0.1206(2)\\ 0.0842(2)\\ 0.1681\\ 0.0498\\ 0.2190(2)\\ 0.1415\\ 0.3566\\ 0.26523(18)\\ 0.3744\\ 0.47226(18)\end{array}$	-0.0631 0.4735 0.1744 0.4299 $0.06671 (19)$ $0.61740 (13)$ $-0.11075 (19)$ $0.62502 (14)$ -0.2346 0.5720 $-0.1206 (2)$ $0.71159 (14)$ $0.0842 (2)$ $0.78214 (14)$ 0.1681 0.7127 0.0498 0.8668 $0.2190 (2)$ $0.83140 (14)$ 0.1415 0.9091 0.3566 0.8720 $0.26523 (18)$ $0.70674 (13)$ 0.3744 0.6421 $0.47226 (18)$ $0.72473 (13)$	-0.0631 0.4735 0.2978 0.1744 0.4299 0.3297 $0.06671 (19)$ $0.61740 (13)$ $0.37828 (7)$ $-0.11075 (19)$ $0.62502 (14)$ $0.41883 (7)$ -0.2346 0.5720 0.4009 $-0.1206 (2)$ $0.71159 (14)$ $0.48942 (7)$ $0.0842 (2)$ $0.78214 (14)$ $0.52003 (7)$ 0.1681 0.7127 0.5534 0.0498 0.8668 0.5523 $0.2190 (2)$ $0.83140 (14)$ $0.45355 (7)$ 0.1415 0.9091 0.4239 0.3566 0.8720 0.4754 $0.26523 (18)$ $0.70674 (13)$ $0.39857 (7)$ 0.3744 0.6421 0.42257 $0.47226 (18)$ $0.72473 (13)$ $0.19025 (7)$

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
01	0.0244 (5)	0.0408 (6)	0.0233 (5)	0.0023 (4)	0.0086 (4)	-0.0059 (4)
C1	0.0194 (6)	0.0190 (6)	0.0190 (6)	-0.0020 (5)	0.0018 (5)	-0.0008 (5)
O2	0.0236 (5)	0.0198 (5)	0.0219 (4)	0.0021 (3)	0.0031 (3)	0.0030 (4)
C2	0.0171 (6)	0.0179 (6)	0.0181 (6)	0.0011 (4)	0.0028 (4)	0.0012 (5)
O3	0.0348 (5)	0.0249 (5)	0.0264 (5)	0.0084 (4)	0.0165 (4)	0.0053 (4)
C3	0.0206 (6)	0.0218 (6)	0.0176 (6)	-0.0016 (5)	0.0040 (5)	-0.0038 (5)
C4	0.0192 (6)	0.0197 (6)	0.0214 (6)	-0.0029 (5)	0.0046 (5)	-0.0027 (5)
C4A	0.0185 (6)	0.0173 (6)	0.0164 (6)	0.0023 (5)	0.0000 (4)	0.0026 (5)
C5	0.0180 (6)	0.0232 (6)	0.0182 (6)	0.0004 (5)	0.0007 (5)	-0.0003 (5)
C6	0.0227 (6)	0.0231 (7)	0.0175 (6)	0.0046 (5)	0.0029 (5)	0.0028 (5)
C7	0.0266 (7)	0.0237 (7)	0.0167 (6)	0.0021 (5)	0.0016 (5)	-0.0036 (5)
C8	0.0235 (6)	0.0220 (6)	0.0201 (6)	-0.0013 (5)	0.0016 (5)	-0.0028 (5)
C8A	0.0177 (6)	0.0189 (6)	0.0175 (6)	0.0009 (5)	0.0004 (5)	0.0005 (5)
С9	0.0155 (6)	0.0183 (6)	0.0190 (6)	-0.0012 (4)	0.0008 (4)	-0.0016 (5)

Geometric parameters (Å, °)

1.2337 (15)	C4—H4A	0.9900
1.5319 (16)	C4—H4B	0.9900
1.5373 (16)	C4A—C5	1.3443 (17)
0.9900	C4A—C8A	1.5106 (16)
0.9900	C5—C6	1.4597 (17)
1.2114 (15)	С5—Н5	0.9500
1.5119 (16)	C6—C7	1.5006 (17)
1.5395 (16)	С7—С8	1.5290 (17)
1.0000	С7—Н7А	0.9900
1.3315 (15)	С7—Н7В	0.9900
0.888 (19)	C8—C8A	1.5287 (17)
1.5323 (16)	C8—H8A	0.9900
0.9900	C8—H8B	0.9900
	1.2337 (15) 1.5319 (16) 1.5373 (16) 0.9900 0.9900 1.2114 (15) 1.5119 (16) 1.5395 (16) 1.0000 1.3315 (15) 0.888 (19) 1.5323 (16) 0.9900	1.2337 (15) $C4$ —H4A $1.5319 (16)$ $C4$ —H4B $1.5373 (16)$ $C4A$ —C5 0.9900 $C4A$ —C8A 0.9900 $C5$ —C6 $1.2114 (15)$ $C5$ —H5 $1.5119 (16)$ $C6$ —C7 $1.5395 (16)$ $C7$ —C8 1.0000 $C7$ —H7A $1.3315 (15)$ $C7$ —H7B $0.888 (19)$ $C8$ —C8A $1.5323 (16)$ $C8$ —H8A 0.9900 $C8$ —H8B

С3—Н3В	0.9900	C8A—H8A1	C8A—H8A1 1.0000	
C4—C4A	1.5038 (17)			
C2—C1—C8A	113.85 (10)	C4A—C5—C6		122.56 (11)
C2—C1—H1A	108.8	C4A—C5—H5		118.7
C8A—C1—H1A	108.8	С6—С5—Н5		118.7
C2—C1—H1B	108.8	O1—C6—C5		120.66 (11)
C8A—C1—H1B	108.8	O1—C6—C7		122.28 (11)
H1A—C1—H1B	107.7	C5—C6—C7		117.00 (10)
C9—C2—C1	110.17 (10)	C6—C7—C8		111.03 (10)
C9—C2—C3	108.79 (9)	С6—С7—Н7А		109.4
C1—C2—C3	109.67 (9)	С8—С7—Н7А		109.4
С9—С2—Н2	109.4	С6—С7—Н7В		109.4
C1—C2—H2	109.4	С8—С7—Н7В		109.4
С3—С2—Н2	109.4	H7A—C7—H7B		108.0
С9—О3—Н3	110.2 (12)	C8A—C8—C7		111.88 (10)
C4—C3—C2	110.47 (9)	C8A—C8—H8A		109.2
С4—С3—Н3А	109.6	C7—C8—H8A		109.2
С2—С3—НЗА	109.6	C8A—C8—H8B		109.2
С4—С3—Н3В	109.6	C7—C8—H8B		109.2
С2—С3—Н3В	109.6	H8A—C8—H8B		107.9
НЗА—СЗ—НЗВ	108.1	C4A—C8A—C8		111.88 (10)
C4A—C4—C3	112.89 (10)	C4A—C8A—C1		111.57 (9)
C4A—C4—H4A	109.0	C8—C8A—C1		109.47 (10)
C3—C4—H4A	109.0	C4A—C8A—H8A1		107.9
C4A—C4—H4B	109.0	C8—C8A—H8A1		107.9
C3—C4—H4B	109.0	C1—C8A—H8A1		107.9
H4A—C4—H4B	107.8	O2—C9—O3		122.57 (11)
C5—C4A—C4	121.19 (11)	O2—C9—C2		123.90 (11)
C5—C4A—C8A	122.59 (11)	O3—C9—C2		113.51 (10)
C4—C4A—C8A	116.14 (10)			
C8A—C1—C2—C9	175.69 (9)	C6—C7—C8—C8A		-55.49 (14)
C8A—C1—C2—C3	55.97 (13)	C5—C4A—C8A—C8		-15.56 (16)
C9—C2—C3—C4	-177.89 (10)	C4—C4A—C8A—C8		167.65 (10)
C1—C2—C3—C4	-57.33 (13)	C5—C4A—C8A—C1		-138.58 (12)
C2—C3—C4—C4A	53.90 (13)	C4—C4A—C8A—C1		44.63 (14)
C3—C4—C4A—C5	134.81 (12)	C7—C8—C8A—C4A		45.26 (13)
C3—C4—C4A—C8A	-48.36 (14)	C7—C8—C8A—C1		169.47 (10)
C4—C4A—C5—C6	171.69 (11)	C2—C1—C8A—C4A		-48.74 (13)
C8A—C4A—C5—C6	-4.94 (18)	C2—C1—C8A—C8		-173.13 (9)
C4A—C5—C6—O1	177.19 (12)	C1—C2—C9—O2		-37.96 (15)
C4A—C5—C6—C7	-5.68 (17)	C3—C2—C9—O2		82.29 (14)
O1—C6—C7—C8	-147.35 (12)	C1—C2—C9—O3		143.64 (10)
C5—C6—C7—C8	35.57 (15)	C3—C2—C9—O3		-96.11 (12)
	. /			~ /
Hydrogen-bond geometry (Å, °)				
D—H···A	<i>D</i> —Н	H···A	$D \cdots A$	D—H··· A
O3—H3···O1 ⁱ	0.888	(19) 1.79 (2)	2.6797 (13)	174.8 (17)

C2—H2···O2 ⁱⁱ	1.00	2.40	3.3191 (15)	152
C7—H7A···O2 ⁱⁱⁱ	0.99	2.47	3.3708 (15)	151
(1)	1/0 1/0 (11)	+ 2/2 + 1/2		

Symmetry codes: (i) x+1, -y+3/2, z-1/2; (ii) -x+1, y-1/2, -z+1/2; (iii) x, -y+3/2, z+1/2.







