Financial support from the New Zealand Lottery Scientific Research Committee for the purchase of a diffractometer is gratefully acknowledged. MMT is also grateful to the Chemistry Department at the University of Canterbury, New Zealand, for providing facilities during his sabbatical stay.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: FR1092). Services for accessing these data are described at the back of the journal.

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Comment

Keto carboxylic acids offer options for varying the pattern of dimeric hydrogen bonding that dominates functionally unadorned acids. Most often, the ketone fails to participate, resulting in typical carboxyl dimers, but less commonly, intermolecular carboxyl-ketone hydrogen bonds occur, yielding a catemer. A third, rare, arrangement has an internal hydrogen bond, while two instances are known of acid-ketone dimerization, and several cases are known of hydrates with more complex hydrogen-bonding patterns. In addition, we have recently reported an instance of carboxyl catemerization, not previously observed in keto acids (Lalancette *et al.*, 1998), and we have cited and discussed numerous examples of these hydrogen-bonding modes (Thompson *et al.*, 1992; Coté *et al.*, 1996).

We report here the structure and hydrogen-bonding behavior of the title δ -keto acid, (I). This category of keto acids embraces examples of dimeric and internal hydrogen bonds, as well as both anhydrous and hydrated catemers. Compound (I) was of particular interest to us as the next-highest homolog of (\pm) -2,3-dihydro-3-oxo-1*H*-indenecarboxylic acid. The crystal structures for both the anhydrous and hydrated forms of this lower homolog, the former of which displays a homochiral (screw-related) catemeric hydrogen-bonding pattern, have been reported recently (Lalancette *et al.*, 1997).



Acta Cryst. (1998). C54, 1180-1182

(±)-4-Oxo-1,2,3,4-tetrahydronaphthalene-1-carboxylic Acid: Hydrogen Bonding and Carboxyl Disordering in a δ -Keto Acid

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(Received 18 December 1997; accepted 13 February 1998)

Abstract

The crystal structure of the title compound $(C_{11}H_{10}O_3)$ contains hydrogen-bonded centrosymmetric carboxyl dimers. Significant carboxyl disorder was found, and was modeled by two rotational conformers, yielding two $O \cdots O$ distances of 2.649 (14) and 2.666 (17) Å.

Our analysis of (I) reveals that the carboxyl group is disordered, both rotationally and with respect to bond lengths and angles, and is turned approximately orthogonal to the plane of the aromatic ring, with the mean carboxyl plane producing a dihedral angle of $77.6 (4)^{\circ}$ with the aromatic ring. Fig. 1 shows a centrosymmetric dimer pair and depicts the modeled rotational disorder of the acid group. The two $O \cdots O$ distances, one for each rotational conformer, are: $O3 \cdot O2^{i} = 2.649(14)$ and $O3' \cdots O2'^{i} = 2.666 (17) \text{ Å}$ [symmetry code: (i) -x, 1-y, -z]. A single molecule of this pair constitutes the asymmetric unit, and its numbering is shown. The flexible ketone ring adopts a conformation resembling a folded envelope, which allows the ketone to lie in the plane of the aromatic ring [torsion angle O1-C4—C4a—C5 = $0.8(3)^{\circ}$], and which positions the carboxyl group pseudo-equatorially [torsion angle C9- $C1-C2-C3 = -176.4(2)^{\circ}$].

For highly ordered dimeric carboxyls, typical values cited for the C—O bond lengths and C—C—O angles are 1.21, 1.31 Å and 123, 112° (Borthwick, 1980).



Fig. 1. ORTEPII (Johnson, 1976) plot of (I), showing a disordered centrosymmetric dimer pair. The major carboxyl conformer [54 (2)%] is shown with filled bonds and segmented ellipsoids; the minor conformer is shown with open bonds and open ellipsoids. The numbering scheme is also shown. Ellipsoids are set at the 20% probability level.

The type of disorder most commonly encountered in carboxyl dimers, detected as an averaging of these values (Leiserowitz, 1976), is most plausibly attributed either to static positional disorder (180° rotation) in the crystal or to rapid tautomeric hydrogen exchange (Meier et al., 1982; Nagaoka et al., 1983; Wilson et al., 1996) within the dimer. For (I), without modeling the rotational disorder, these bond lengths and angles are not only substantially averaged [1.240(3), 1.238(3) Å, and 119.5 (2), 118.3 (2) $^{\circ}$], but the carboxyl O atoms are markedly prolated, suggesting that the averaging may be accompanied or caused by rotation of the dimer subunit. As a general measure of such prolation, we use the ratio of the length of the longest ellipsoidal axis to the sum of the remaining two. Significant electron density was also found within the O-H bond distance from both carboxyl O atoms. We have modeled this disorder most successfully as a carboxyl rotation of $156(1)^{\circ}$, as shown in Fig. 1 (see refinement details). The major conformer in this treatment refined to a site occupancy of 54 (2)%, while the minor component was 46(2)%. This refinement led to a partial ordering of the carboxyl bond lengths and angles [C9-O2 = 1.234(8)], C9-O3 = 1.263(8), C9-O2' = 1.237(9), C9-O3' =1.259(9) Å; O2---C9---C1 = 120.5(6), O3---C9---C1 = 116.7 (5), O2'—C9—C1 = 122.5 (6), O3'—C9—C1 = 115.1 (7)°]. With this treatment, the atomic displacement parameters for the carboxyl O atoms are substantially reduced.

A close contact of 2.59 Å is found between the ketone oxygen, O1, and H7A of a translationally related neighbor. A further close contact of 2.69 Å to a translationally related neighbor is found between H2B and O2'; details are in Table 2. These may represent polar attractions contributing materially to the packing forces (Jönsson, 1972; Leiserowitz, 1976; Berkovitch-Yellin & Leiserowitz, 1982).

The solid-state (KBr) infrared spectrum of (I) has a double peak (1688, 1691 cm⁻¹) amid several lesser peaks, for both C=O groups. In CHCl₃ solution, the

ketone absorption at 1687 cm^{-1} is accompanied by a less intense one at 1712 cm^{-1} , having a major shoulder at *ca* 1725 cm^{-1} .

Experimental

Compound (I) was synthesized by polyphosphoric acid cyclization (Lloyd & Horning, 1954) of 2-phenylglutaric anhydride, purchased from Acros Organics, Pittsburgh, PA, USA. Crystals of (I) (m.p. 368 K) were obtained by evaporation from a solution in a 3:1 cyclohexane/ethyl acetate mixture.

Crystal data

$M_r = 190.19$ $\lambda = 0.71073$ ÅMonoclinicCell parameters from 2 $P2_1/n$ reflections	23
Monoclinic Cell parameters from 2 $P2_1/n$ reflections	23
$P2_1/n$ reflections	
$a = 6.4046(9) \text{ Å}$ $\theta = 10.2 - 14.0^{\circ}$	
$b = 5.2550 (8) \text{ Å}$ $\mu = 0.099 \text{ mm}^{-1}$	
c = 27.634 (3) Å $T = 293$ (2) K	
$\beta = 92.111 (8)^{\circ}$ Parallelepiped	
$V = 929.4$ (2) Å ³ $0.56 \times 0.20 \times 0.10$ m	m
Z = 4 Colorless	
$D_x = 1.359 \text{ Mg m}^{-3}$	
$D_m = 1.36 (1) \text{ Mg m}^{-3}$	
D_m measured by flotation in	
hexane/CCl ₄	

Data collection

Siemens P4 diffractometer	$R_{\rm int} = 0.035$
$2\theta/\theta$ scans	$\theta_{\rm max} = 25^{\circ}$
Absorption correction:	$h = -1 \rightarrow 7$
face-indexed numerical	$k = -1 \rightarrow 6$
(Sheldrick, 1994)	$l = -32 \rightarrow 32$
$T_{\min} = 0.960, T_{\max} = 0.982$	3 standard reflections
2600 measured reflections	every 97 reflections
642 independent reflections	intensity decay: 0.10%
101 reflections with	
$F > 4\sigma(F)$	

Refinement

j	
Refinement on F^2	$(\Delta/\sigma)_{\rm max} = -0.001$
$R[F^2 > 2\sigma(F^2)] = 0.041$	$\Delta \rho_{\rm max} = 0.115 \ {\rm e} \ {\rm \AA}^{-3}$
$wR(F^2) = 0.108$	$\Delta \rho_{\rm min} = -0.128 \ {\rm e} \ {\rm \AA}^{-3}$
S = 1.03	Extinction correction:
1639 reflections	SHELXTL/PC
156 parameters	Extinction coefficient:
H atoms: see below	0.124 (10)
$w = 1/[\sigma^2(F_o^2) + (0.045P)^2]$	Scattering factors from
+ 0.058P]	International Tables for
where $P = (F_o^2 + 2F_c^2)/3$	Crystallography (Vol. C)

Table 1. Selected geometric parameters (Å, °)

01—C4 02—C9 03—C9	1.219 (2) 1.234 (8) 1.263 (8)	O2'—C9 O3'—C9	1.237 (9) 1.259 (9)
02—C9—C1	120.5 (6)	O3'C9C1	115.1 (7)
02′—C9—C1	122.5 (6)	O3C9C1	116.7 (5)

Table 2. Hydrogen-bonding and close-contact geometry $(\overset{\circ}{A} \circ)$

	(, , ,		
D—H···A	D—H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - H \cdot \cdot \cdot A$
O3—H33· · ·O2'	0.82	1.842	2.649 (14)	168
O3'-H33'···O2''	0.82	1.858	2.666 (17)	168
C2—H2B· · · O2′ ^𝑥	0.97	2.69	3.393 (9)	130
C7—H7A· · ·O1 [™]	0.93	2.59	3.388 (3)	144
Symmetry codes: (i)	-x, 1 - y, -	-z; (ii) x, 1 +	- y, z; (iii) x –	-1, y - 1, z

All non-carboxyl H atoms were found in electron-density difference maps but replaced in calculated positions and allowed to refine as riding models on their appropriate C atoms, with their displacement parameters free to refine. Residual electron density near both the carboxyl O atoms, and averaging of carboxyl bond lengths and angles, was indicative of carboxyl disorder. The disordered carboxyl group was modeled as two rotational conformers, with carboxyl O-atom occupancies initially set to 50%. Each carboxyl group was restrained to be planar during the early cycles of refinement, but this restraint was removed once the refinement had converged. Carboxyl H atoms (with occupancies of 50% each) were added in calculated positions to the carboxyl O atom of each conformer having the longest C—O bond; their displacement parameters were fixed at 0.08\AA^2 . The site occupancies of the two carboxyl O atoms and the H atom for both conformers were then allowed to refine with total occupancy constrained to be unity. The disordered C=O and C=O' bond lengths were restrained to be equal to each other within 0.005 Å; the same restraint was imposed on the disordered C-O and C-O' bond lengths. The major carboxyl conformer refined to a site occupancy of 54 (2)%, and the minor component was 46 (2)%.

Data collection: XSCANS (Siemens, 1996). Cell refinement: XSCANS. Data reduction: XSCANS. Program(s) used to solve structure: SHELXTL/PC (Sheldrick, 1994). Program(s) used to refine structure: SHELXTL/PC. Molecular graphics: SHELXTL/PC. Software used to prepare material for publication: SHELXTL/PC.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: FG1433). Services for accessing these data are described at the back of the journal.

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Acta Cryst. (1998). C54, 1182-1184

β-Dihydroentandrophragmin–Ethyl Acetate (1/0.355)

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(Received 7 January 1998; accepted 27 April 1998)

Abstract

Although the title compound, $C_{43}H_{58}O_{17}.0.355C_4H_8O_2$, from *Entandrophragma cylindricum*, exhibits significant disorder in its ester side chains, it was possible to determine the absolute configurations of the (2S)-2-methylbutanoate and (2S)-2-hydroxy-2-methylbutanoate sidechain groups by comparison with the stereochemistry of the core secotetranortriterpenoid skeleton.

Comment

The commercially important mahogany timber trees of the genus Entandrophragma produce a small number of highly functionalized structurally complex secotetranortriterpenes, such as utilin, (I), and entandrophragmin, (II), which are characterized by the presence of an orthoester function (Arene et al., 1966; Taylor & Wragg, 1967; Harrison et al., 1970; Taylor, 1974; Halsall et al., 1977). Interest in natural products containing this relatively rare structural unit is stimulated by the toxic and antileukaemic activities of other orthoestercontaining natural products, such as the daphnetoxintype Thymelaeceae diterpenoids (Evans & Taylor, 1983; Baxter & Ziegler, 1994). β -Dihydroentandrophragmin, (III), was isolated previously by Halsall et al. (1977) and its structure deduced by comparison of its ¹³C and ¹H NMR spectra with those of entandrophragmin, the structure and stereochemistry of which were in turn based on the crystal structure of utilin (Harrison et al., 1970). However, the absolute stereochemistry of the 2-methylbutanoate and 2-hydroxy-2-methylbutanoate side chains