

C3—C4—C14	114.2 (7)	O4—C18—O5	127.3 (9)	Ahmad, V. U., Farooqui, T. A., Sultana, A., Fizza, K. & Khatoon, R. (1992). <i>Phytochemistry</i> , 31 , 2888–2890.
C5—C4—C14	113.9 (6)	O4—C18—C19	109.1 (7)	Ahmad, V. U., Fizza, K. & Amber, A.-Ur-R. (1989). <i>J. Nat. Prod.</i> 52 , 861–863.
C4—C5—C6	113.9 (5)	O5—C18—C19	123.5 (7)	Gilmore, C. J. (1984). <i>J. Appl. Cryst.</i> 17 , 42–46,
C4—C5—C10	114.9 (6)	O6—C19—C18	112.7 (7)	Ivie, R. A., Watson, W. H. & Dominguez, X. A. (1974). <i>Acta Cryst. B</i> 30 , 2891–2893.
C6—C5—C10	111.6 (5)	O6—C19—C20	61.5 (6)	Johnson, C. K. (1976). <i>ORTEPII</i> . Report ORNL-5138. Oak Ridge National Laboratory, Tennessee, USA.
C5—C6—C7	113.0 (6)	O6—C19—C21	113.6 (7)	Molecular Structure Corporation (1985). <i>TEXSAN. TEXRAY Structure Analysis Package</i> . MSC, 3200 Research Forest Drive, The Woodlands, TX 77381, USA.
C6—C7—C8	114.4 (7)	C18—C19—C20	115.1 (8)	Molecular Structure Corporation (1988). <i>MSC/AFC Diffractometer Control Software</i> . MSC, 3200 Research Forest Drive, The Woodlands, TX 77381, USA.
C6—C7—C11	124.0 (7)	C18—C19—C21	117.1 (7)	Motherwell, S. & Clegg, W. (1978). <i>PLUTO. Program for Plotting Molecular and Crystal Structures</i> . University of Cambridge, England.
C8—C7—C11	121.5 (7)	C20—C19—C21	123 (1)	Mukhopadhyay, S., Cordell, G. A., Ruangrungsi, N., Rodkird, S., Tantivatana, P. & Hylands, P. J. (1983). <i>J. Nat. Prod.</i> 46 , 671–674.
O1—C8—C7	122.5 (8)	O6—C20—C19	59.6 (5)	Nakanishi, K., Crouch, R., Miura, I., Dominguez, X. A., Zamudio, A. & Villarreal, R. (1974). <i>J. Am. Chem. Soc.</i> 96 , 609–611.
O1—C8—C9	122.3 (7)	O6—C20—C22	114 (1)	North, A. C. T., Phillips, D. C. & Mathews, F. S. (1968). <i>Acta Cryst. A24</i> , 351–359.
C7—C8—C9	115.2 (6)	C19—C20—C22	127 (1)	Spek, A. L. (1990). <i>Acta Cryst. A46</i> , C-34
C8—C9—C10	114.4 (6)			Uchiyama, T., Miyase, T., Ueno, A. & Usmanhani, K. (1991). <i>Phytochemistry</i> , 30 , 655–657.
O1—C8—C7—C6	139.2 (9)	C3—C2—C1—C10	-56.7 (9)	
O1—C8—C7—C11	-39 (1)	C3—C4—O2—C16	-58.5 (8)	
O1—C8—C9—C10	-131.2 (8)	C3—C4—C5—C6	179.6 (6)	
O2—C4—C3—O4	-37.5 (7)	C3—C4—C5—C10	49.1 (8)	
O2—C4—C3—C2	-159.6 (6)	C4—O2—C16—C17	-176.2 (7)	
O2—C4—C5—C6	-63.0 (6)	C4—C3—O4—C18	151.4 (6)	
O2—C4—C5—C10	166.4 (5)	C4—C5—C6—C7	172.9 (6)	
O3—C16—O2—C4	6 (1)	C4—C5—C10—C9	-169.2 (6)	
O4—C3—C2—C1	-66.9 (9)	C4—C5—C10—C15	71.5 (8)	
O4—C3—C4—C5	71.3 (6)	C5—C4—O2—C16	-175.2 (6)	
O4—C3—C4—C14	-158.3 (5)	C5—C6—C7—C8	43.9 (9)	
O4—C18—C19—O6	-179.3 (6)	C5—C6—C7—C11	-138.1 (8)	
O4—C18—C19—C20	-111.3 (8)	C5—C10—C9—C8	-56.7 (8)	
O4—C18—C19—C21	46 (1)	C6—C5—C4—C14	49.1 (8)	
O5—C18—O4—C3	7 (1)	C6—C5—C10—C9	59.1 (8)	
O5—C18—C19—O6	3 (1)	C6—C5—C10—C15	-60.2 (9)	
O5—C18—C19—C20	71 (1)	C6—C7—C8—C9	-41 (1)	
O5—C18—C19—C21	-132 (1)	C6—C7—C11—C12	180.0 (7)	
O6—C19—C20—C22	99 (1)	C6—C7—C11—C13	3 (1)	
O6—C20—C19—C18	-103.3 (9)	C7—C6—C5—C10	-54.9 (9)	
O6—C20—C19—C21	101 (1)	C7—C8—C9—C10	49 (1)	
C1—C2—C3—C4	55 (1)	C8—C7—C11—C12	-2 (1)	
C1—C10—C5—C4	-50.4 (8)	C8—C7—C11—C13	-179.4 (8)	
C1—C10—C5—C6	177.9 (6)	C8—C9—C10—C15	66.8 (8)	
C1—C10—C9—C8	-174.6 (6)	C9—C8—C7—C11	140.9 (8)	
C2—C1—C10—C5	54.5 (9)	C10—C5—C4—C14	-81.5 (7)	
C2—C1—C10—C9	170.9 (6)	C14—C4—O2—C16	66.7 (8)	
C2—C1—C10—C15	-70.9 (8)	C18—C19—O6—C20	107.3 (9)	
C2—C3—O4—C18	-83.0 (7)	C18—C19—C20—C22	-5 (1)	
C2—C3—C4—C5	-50.8 (9)	C19—O6—C20—C22	-121 (1)	
C2—C3—C4—C14	79.6 (8)	C20—O6—C19—C21	-117 (1)	
C3—O4—C18—C19	-170.5 (6)	C21—C19—C20—C22	-161 (1)	

All non-H atoms were refined with anisotropic displacement parameters. H atoms were included in the difference Fourier map locations or at calculated positions with isotropic displacement parameters estimated from the displacement parameter of the adjacent C atom.

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1988). Cell refinement: *MSC/AFC Diffractometer Control Software*. Data reduction: *PROCESS TEXSAN* (Molecular Structure Corporation, 1985). Program(s) used to solve structure: *TEXSAN*; *MITHRIL* (Gilmore, 1984). Program(s) used to refine structure: *LS TEXSAN*. Molecular graphics: *ORTEPII* (Johnson, 1976); *PLUTO* (Motherwell & Clegg, 1978). Software used to prepare material for publication: *PLATON* (Spek, 1990).

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: BK1143). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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Structural Aspects of the Nucleophilic Attack of Ketone Enolate on Dehydronaphthalene

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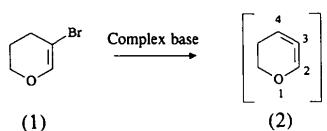
Abstract

The results of the crystal structure analyses of (6*RS*,7*SR*,8*RS*)-7-methyl-8-phenyl-3-oxabicyclo[4.2.0]oct-1-en-8-ol ($C_{14}H_{16}O_2$), (1*SR*,2*RS*,8*RS*)-8-methylthio-3-oxatricyclo[6.4.0.0^{2,7}]dodec-6-en-1-ol ($C_{12}H_{18}O_2S$) and (1*RS*,2*RS*,6*SR*,7*SR*,8*RS*)-8-methyl-3-oxatricyclo[6.4.0.0^{2,7}]dodeca-1,6,7-triol ($C_{12}H_{20}O_4$) allow interpretation of the nucleophilic attack reactions of ketone enolates on the short-lived dehydronaphthalene intermediate, showing

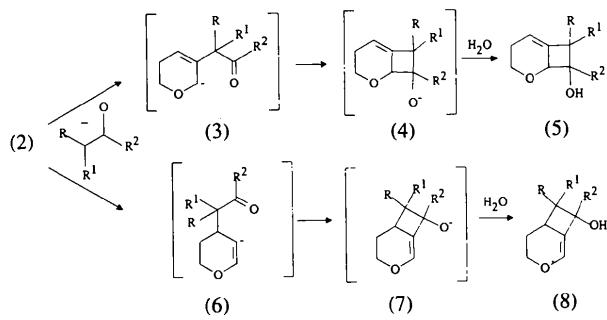
that the presence of the O atom influences the reactivity of the dienic system. The structures and conformations of the above compounds are discussed.

Comment

We have shown previously (Jamart-Grégoire, Grand, Ianelli, Nardelli & Caubère, 1990) that dehydrodihydro-pyran (2) can easily be generated from 3-bromodihydro-pyran (1) and a complex base (Caubère, 1978, 1991). In addition, we showed that, under the reaction conditions defined, nucleophiles such as ketone enolates condensed easily with the short-lived intermediate (2).



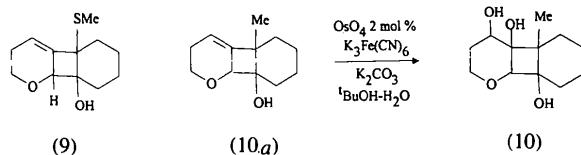
Taking into account the nature of (2), the same type of behaviour as observed for 1,2-cyclohexadienes (Caubère & Brunet, 1972; Brunet, Fixari & Caubère, 1974) could be expected, with nucleophilic attack of the ketone enolate at the C3 position followed by a cyclization of the anionic intermediate. However, the presence of the O atom in the ring could cause some perturbation leading to alternative attacks from the diene system. In reality, depending on the nature of the ketone enolate, attack at the C3 and/or C4 positions was observed, as shown below. Moreover, it was also observed that (4) and (7) could evolve differently to give ketone derivatives (Jamart-Grégoire, Grand, Ianelli, Nardelli & Caubère, 1990).



With respect to the reactions above, the following questions must be answered: What is the stereochemistry of (5) and in particular does it come from a *syn* addition of (2) on the ketone enolate? And, in the absence of information about the reactivity of (2), were the structures attributed to the completely unexpected compound (8) correct and what was their stereochemistry?

The answer to the first question was easily obtained through the usual spectroscopic data as far as the unsubstituted ketone enolates were used (R or $R^1 = H$). Unfortunately, no such information could be obtained

with α -substituted ketones (R and $R^1 \neq H$), so the X-ray crystal structure analysis of compounds (9) and (10) was performed, the latter compound being studied to infer the stereochemistry of the liquid compound (10a).



Regarding the second question, we were lucky enough to obtain single crystals of (8) ($R = \text{CH}_3$, $\bar{R}^2 = \text{Ph}$) suitable for X-ray crystal structure analysis.

The results of these structure analyses, illustrated in the present paper, are of interest as they allow definition of the structures of analogous compounds obtained according to the elimination-addition reaction of the scheme above, through comparison of their spectroscopic data. Moreover, they show that the *syn*-addition of ketone enolates with the short-lived 1,2-dienes seems to be the rule. More importantly, they definitively establish that nucleophilic attacks on the C4-position of (2) may take place, showing the dramatic influence of the presence of the O atom on the reactivity of the dienic system.

Fig. 1 shows the *ORTEP* (Johnson, 1965) drawings of the molecules of the three compounds. It should be noted that, while both enantiomeric species are present in the crystals of compounds (9) and (10) as their space groups are centrosymmetric, in the case of compound (8), the two enantiomers do not co-crystallize and only one is present in the crystal. For the crystals of (8), the anomalous-scattering effects were not large enough to allow assignment of the absolute configuration [the value of Flack's (1983) parameter was $x = -0.3(4)$], so that assigned in Fig. 1 is one of the two possible.

The geometry of the three molecules is described in Tables 2, 4 and 6, respectively, while Table 7 gives the relevant parameters describing the conformation of the three ring systems in the three molecules. The presence of the double bond in the dihydropyran ring of compounds (8) and (9) reduces the tilting of the plane through ring A with respect to that through the central cyclobutane ring B.

The conformation of compound (8) can be deduced from the torsion angles in Table 2, which show that the C1—C8 bond is (-)anticlinal to C2—C3, (+)-synperiplanar to C7—H7 [C8—C1—C7—H7 = 18(2) $^{\circ}$] and (+)anticlinal to C7—C14, which is (+)synperiplanar to C1—O1.

In compounds (9) and (10), the system of fused tricycles differs mainly in the presence of the double bond in ring A of compound (9), which causes bond C1—C2 to be (-)synperiplanar to C7—C8 (Table 4), while in compound (10) C1—C2 is (-)antiperiplanar to C7—C8 (Table 6). Another relevant difference in

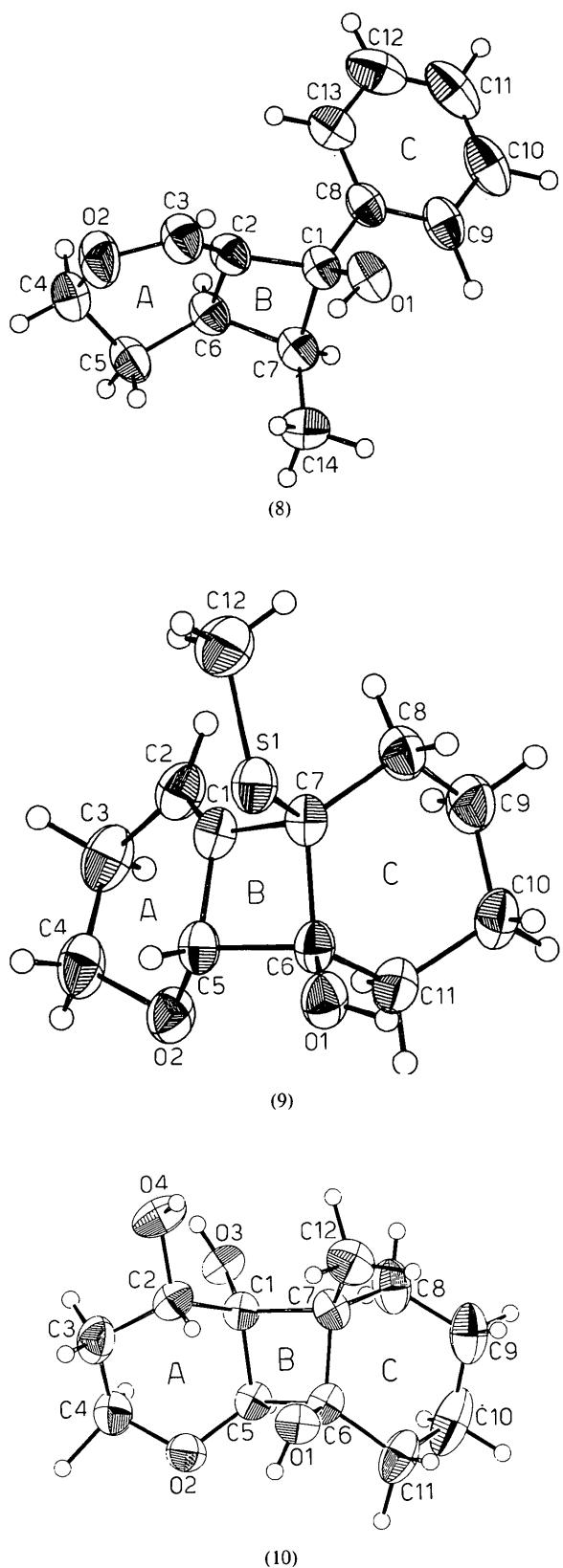


Fig. 1. ORTEP drawings of the molecules of compounds (8), (9) and (10). Ellipsoids are plotted at the 50% probability level.

the conformation of these two systems of fused rings is observed for bond C5—O2 which is (+)synperiplanar to C6—C11 in compound (9) (Table 4) and (+)anticlinal in compound (10) (Table 6).

The substituent-carrying side of the cyclobutane ring [C1—C7 for (8), C6—C7 for (9) and (10)] is significantly longer than the other three sides in all three compounds, in agreement with the results of the molecular-mechanics calculations (Tables 2, 4 and 6).

To check the reliability of the structural parameters describing the geometry of the three molecules, particularly for compound (10) whose analysis gave a quite unsatisfactory *R* value (see *Experimental*), and to see how packing forces in the crystal influence the molecular conformation, molecular-mechanics calculations were carried out using both MMX and MM+ force fields of the PCMODEL (Serena Software, 1989) and HYPERCHEM (Autodesk, Inc., 1992) systems of programs, respectively. In these calculations, isolated molecules were considered and program default parameters were used, starting the energy-minimization process from the molecular structures found experimentally by X-ray diffraction analysis. The results in terms of bond distances, bond angles and torsion angles are compared with those from the experimental analysis in Tables 2, 4 and 6. From this comparison, the following general comments can be made: (i) a rough general evaluation of the disagreement between the experimental and calculated models is given by the averages of the absolute values of the differences $|\Delta|$ (Table 9); (ii) concerning bond distances, negative differences can be justified by the fact that the experimental bond distances are not corrected for thermal motion; (iii) some relevant differences in bond angles are observed when atoms involved in hydrogen bonding or belonging to the cyclobutane ring are considered; (iv) the torsion angles are the parameters showing the largest discrepancies, which can be justified by the packing interactions not considered in the calculated models; (v) it is surprising to find that the lowest differences are observed with compound (10) whose diffraction results, in terms of the *R* index, are the least accurate.

The orientation of the phenyl ring about the C1—C8 bond in compound (8) is defined by the torsion angles about this bond; the differences between the observed and calculated values found for these angles in the isolated molecule (Table 2) are indicative that there is some influence of the packing forces. This finding is in agreement with the non-bonded energy profile, calculated by rotating the phenyl ring about C1—C8 in the isolated molecule, which shows a minimum shifted by *ca* 7° with respect to the position found for the molecule in the crystal. A similar situation is found for the orientation of the methylthio group in the molecule of compound (9), which is defined by the torsion angles about the C7—S1 bond (Table 4). Also in this case, the differences are only a few degrees and the non-bonded

energy profile shows a wide minimum shifted by *ca* 10° with respect to the position found in the crystal.

The orientation of the hydroxyl group in compounds (8) and (9) is determined by the hydrogen bond it forms as a donor towards the pyran O atom and the methylthio S atom of an adjacent molecule, respectively (Table 8). These bonds, together with normal van der Waals interactions, are responsible for the packing in the crystals. In compound (10), the hydroxyls O1—H1O and O3—H3O are both involved, as donor, in intra- and intermolecular bifurcated hydrogen bonds of the type O—H···O, while the O4—H4O hydroxyl group forms only an intermolecular hydrogen bond of this type. Intra- and intermolecular interactions of the type C—H···O, which can be considered as weak hydrogen bonds (Taylor & Kennard, 1982; Berkovitch-Yellin & Leiserowitz, 1984), are also observed (Table 8).

The analysis of 'thermal' motion of the molecules in the crystals of the three compounds, carried out in terms of the Schomaker & Trueblood (1968) TLS rigid-body approximation using the THMV program (Trueblood, 1984), shows that there are no cases of possible disorder, giving quite satisfactory agreements between the observed and calculated atomic displacement parameters, particularly for compound (9). The overall residual disagreement index wR_U is 0.085, 0.055 and 0.094 for (8), (9) and (10), respectively, and these improve to 0.057, 0.039 and 0.067, respectively, if internal motions are considered according to Dunitz & White (1973). These findings for compound (10) are also indicative of the reliability of the results of the diffraction analysis in spite of the inferior value of the *R* index.

Experimental

Compound (8)

Crystal data

$C_{14}H_{16}O_2$
 $M_r = 216.28$
Orthorhombic
 $P2_12_12_1$
 $a = 16.305 (7) \text{ \AA}$
 $b = 10.366 (4) \text{ \AA}$
 $c = 6.900 (4) \text{ \AA}$
 $V = 1166.2 (9) \text{ \AA}^3$
 $Z = 4$
 $D_x = 1.232 \text{ Mg m}^{-3}$

Data collection

Siemens-AED diffractometer
 θ/θ scans
Absorption correction:
none
2508 measured reflections
2209 independent reflections
1821 observed reflections [$I > 2\sigma(I)$]
 $R_{\text{int}} = 0.0525$

$\text{Cu } K\alpha$ radiation
 $\lambda = 1.54178 \text{ \AA}$
Cell parameters from 30 reflections
 $\theta = 23\text{--}40^\circ$
 $\mu = 0.644 \text{ mm}^{-1}$
 $T = 293 (2) \text{ K}$
Tablets
 $0.36 \times 0.29 \times 0.21 \text{ mm}$
Colourless

Refinement

Refinement on F^2
 $R(F) = 0.0429$
 $wR(F^2) = 0.0975$
 $S = 1.107$
2203 reflections
210 parameters
All H-atom parameters refined
 $w = 1/[\sigma^2(F_o^2) + (0.0539P)^2 + 0.1849P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.214 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.163 \text{ e \AA}^{-3}$

Extinction correction:
SHELXL93 (Sheldrick, 1993)

Extinction coefficient:
0.107 (5)
Atomic scattering factors from *International Tables for Crystallography* (1992, Vol. C, Tables 4.2.6.8, 6.1.1.4)

Absolute configuration:
Flack (1983)
Flack parameter = -0.3 (4)

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (\AA^2) for (8)

	x	y	z	U_{eq}
O1	0.1200 (1)	0.7085 (2)	0.1529 (3)	0.0564 (6)
O2	0.3118 (1)	0.5375 (2)	-0.2144 (4)	0.0750 (8)
C1	0.1687 (1)	0.7806 (2)	0.0209 (3)	0.0440 (6)
C2	0.2267 (1)	0.7074 (2)	-0.1091 (4)	0.0476 (6)
C3	0.2376 (2)	0.5852 (2)	-0.1503 (5)	0.0617 (9)
C4	0.3746 (2)	0.6342 (3)	-0.2406 (5)	0.0642 (9)
C5	0.3767 (2)	0.7320 (3)	-0.0774 (5)	0.0579 (8)
C6	0.2952 (1)	0.8034 (2)	-0.0807 (4)	0.0475 (7)
C7	0.2480 (1)	0.8508 (2)	0.1023 (4)	0.0481 (7)
C8	0.1105 (1)	0.8727 (2)	-0.0799 (4)	0.0462 (7)
C9	0.0708 (1)	0.9679 (2)	0.0295 (5)	0.0587 (8)
C10	0.0156 (2)	1.0522 (3)	-0.0601 (6)	0.0725 (12)
C11	0.0006 (2)	1.0429 (3)	-0.2547 (7)	0.0802 (12)
C12	0.0396 (2)	0.9510 (4)	-0.3618 (6)	0.0802 (13)
C13	0.0949 (2)	0.8654 (3)	-0.2763 (4)	0.0603 (9)
C14	0.2775 (2)	0.7986 (3)	0.2933 (4)	0.0625 (9)

Table 2. Comparison of bond distances (\AA), bond angles ($^\circ$) and selected torsion angles ($^\circ$) in the molecule of compound (8)

E.s.d.'s in parentheses. MMX = force field of PCMODEL; MM+ = force field of HYPERCHEM; Δ = obs. - calc.

	X-ray obs.	MMX calc.	MM+ Δ	MM+ calc.	MM+ Δ
O1—C1	1.421 (3)	1.408	0.013	1.431	-0.010
O2—C3	1.380 (3)	1.357	0.023	1.365	0.015
O2—C4	1.443 (4)	1.424	0.019	1.416	0.027
C1—C2	1.508 (3)	1.503	0.005	1.454	0.054
C1—C7	1.586 (3)	1.578	0.008	1.582	0.004
C1—C8	1.515 (3)	1.509	0.006	1.478	0.037
C2—C3	1.311 (3)	1.331	-0.020	1.337	-0.026
C2—C6	1.509 (3)	1.500	0.009	1.445	0.064
C4—C5	1.516 (4)	1.546	-0.030	1.541	-0.025
C5—C6	1.521 (3)	1.530	-0.009	1.545	-0.024
C6—C7	1.559 (4)	1.557	0.002	1.571	-0.012
C7—C14	1.504 (4)	1.530	-0.026	1.548	-0.044
C8—C9	1.401 (3)	1.345	0.056	1.399	0.002
C8—C13	1.381 (4)	1.345	0.036	1.400	-0.019
C9—C10	1.399 (4)	1.342	0.057	1.396	0.003
C10—C11	1.368 (6)	1.341	0.027	1.395	-0.027
C11—C12	1.363 (5)	1.341	0.022	1.395	-0.032
C12—C13	1.396 (5)	1.342	0.054	1.396	0.000
C3—O2—C4	114.4 (2)	121.0	-6.6	116.5	-2.1
O1—C1—C8	106.0 (2)	113.0	-7.0	112.2	-6.2
O1—C1—C7	118.0 (2)	113.2	4.8	114.5	3.5

O1—C1—C2	117.8 (2)	112.1	5.7	113.1	4.7	<i>Refinement</i>
C7—C1—C8	112.6 (2)	114.7	-2.1	114.1	-1.5	Refinement on F^2
C2—C1—C8	115.9 (2)	117.0	-1.1	114.9	1.0	$R(F) = 0.0322$
C2—C1—C7	86.0 (2)	83.9	2.1	85.5	0.5	$wR(F^2) = 0.0899$
C1—C2—C6	93.2 (2)	94.5	-1.3	99.2	-6.0	$S = 1.091$
C1—C2—C3	134.4 (2)	139.0	-4.6	128.3	6.1	2212 reflections
C3—C2—C6	124.4 (2)	126.3	-1.9	123.7	0.7	209 parameters
O2—C3—C2	122.3 (2)	119.2	3.1	123.5	-1.2	All H-atom parameters
O2—C4—C5	112.8 (2)	111.5	1.3	111.9	0.9	refined
C4—C5—C6	107.1 (2)	108.1	-1.0	107.0	0.1	$w = 1/[\sigma^2(F_o^2) + (0.0626P)^2]$
C2—C6—C5	109.2 (2)	104.5	4.7	107.0	2.2	where $P = (F_o^2 + 2F_c^2)/3$
C5—C6—C7	125.0 (2)	125.9	-0.9	121.9	3.1	$(\Delta/\sigma)_{\max} < 0.001$
C2—C6—C7	86.9 (2)	84.8	2.1	86.2	0.7	$\Delta\rho_{\max} = 0.268 \text{ e } \text{\AA}^{-3}$
C1—C7—C6	88.4 (2)	89.5	-1.1	88.8	-0.4	$\Delta\rho_{\min} = -0.323 \text{ e } \text{\AA}^{-3}$
C6—C7—C14	116.0 (2)	115.3	0.7	116.5	-0.5	
C1—C7—C14	114.0 (2)	116.1	-2.1	115.8	-1.8	
C1—C8—C13	122.1 (2)	122.2	-0.1	121.5	0.6	
C1—C8—C9	119.1 (2)	119.5	-0.4	119.5	-0.4	
C9—C8—C13	118.8 (2)	118.2	0.6	118.9	-0.1	
C8—C9—C10	120.0 (2)	121.2	-1.2	120.6	-0.6	
C9—C10—C11	120.3 (3)	119.9	0.4	120.0	0.3	
C10—C11—C12	119.9 (4)	119.6	0.3	119.7	0.2	
C11—C12—C13	121.1 (3)	120.1	1.0	120.1	1.0	
C8—C13—C12	119.9 (3)	121.0	-1.1	120.5	-0.6	
O1—C1—C8—C13	-115.5 (2)	-110.8	-4.7	-103.9	-11.6	<i>Table 3. Fractional atom</i>
O1—C1—C7—C14	19.0 (3)	13.0	6.0	-1.8	20.8	<i>isotropic displacement</i>
O1—C1—C2—C3	10.3 (4)	43.3	-33.0	28.6	-18.3	$U_{\text{eq}} = (1/3)\Sigma$
C7—C1—C8—C9	-66.3 (3)	-63.2	-3.1	-60.9	-5.4	<i>x</i>
C2—C1—C8—C9	-163.1 (2)	-159.1	-4.0	-157.4	-5.7	S1 0.31535 (3) 0.104
C7—C1—C8—C13	114.1 (2)	117.5	-3.4	123.7	-9.6	O1 0.3930 (1) -0.050
C2—C1—C7—C14	-100.6 (2)	-98.5	-2.1	-115.2	14.6	O2 0.09865 (9) -0.267
C8—C1—C2—C3	-116.8 (3)	-89.4	-27.4	-102.1	-14.7	C1 0.1441 (1) -0.173
C8—C1—C7—C14	143.0 (2)	144.6	-1.6	129.5	13.5	C2 0.0430 (2) -0.263
C3—C2—C6—C7	-134.5 (3)	-156.3	21.8	-145.4	10.9	C3 -0.0586 (2) -0.320
C5—C6—C7—C14	-12.6 (3)	-5.1	-7.5	6.8	-19.4	C4 -0.0281 (2) -0.266
C2—C6—C7—C14	98.8 (2)	99.1	-0.3	114.6	-15.8	C5 0.1712 (1) -0.148
						C6 0.3060 (1) -0.165
						C7 0.2783 (1) -0.123

Distances: $|\Delta|_{\max}$ MMX = 0.057 Å, MM+ = 0.064 Å; angles: $|\Delta|_{\max}$ MMX = 7.0°, MM+ = 6.2°; torsions: $|\Delta|_{\max}$ MMX = 33.0°, MM+ = 20.8°.

Compound (9)

Crystal data

$$\text{C}_{12}\text{H}_{18}\text{O}_2\text{S}$$

$$M_r = 226.33$$

Triclinic

P1

$$a = 11.0$$

$$b = 8.30$$

$$c = 6.625 (1) \text{ \AA}$$

$$\alpha = 1$$

$$\beta = 93.66^\circ$$

$$V = 583.4 \text{ (2)} \text{ \AA}^3$$

Data collection

Data collection

Siemens A
A12A scans

0/20 scans

Response
none

2231 measured reflections

2218 independent reflections

1913 observed reflections

$$[I > 2\sigma(I)]$$

$$R_{\text{int}} = 0.0090$$

Cu K α radiation
 $\lambda = 1.54178 \text{ \AA}$
 Cell parameters from 30
 reflections
 $\theta = 25\text{--}39^\circ$
 $\mu = 2.240 \text{ mm}^{-1}$
 $T = 293$ (2) K
 Prisms
 $0.43 \times 0.33 \times 0.28 \text{ mm}$
 Colourless

Table 4. Comparison of bond distances (\AA), bond angles ($^\circ$) and selected torsion angles ($^\circ$) in the molecule of compound (9)

E.s.d.'s in parentheses. MMX = force field of PCMODEL; MM+ = force field of HYPERCHEM; Δ = obs. - calc.

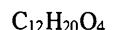
		X-ray obs.	MMX	MM+		
		calc.	Δ	calc.	Δ	
$\theta = 25\text{--}39^\circ$						
$\mu = 2.240 \text{ mm}^{-1}$	S1—C7	1.839 (2)	1.829	0.010	1.799	0.040
$T = 293 \text{ (2) K}$	S1—C12	1.796 (3)	1.812	-0.016	1.813	-0.017
Prisms	O1—C6	1.416 (2)	1.412	0.004	1.432	-0.016
$0.43 \times 0.33 \times 0.28 \text{ mm}$	O2—C4	1.443 (2)	1.432	0.011	1.415	0.028
Colourless	O2—C5	1.419 (2)	1.415	0.004	1.436	-0.017
	C1—C2	1.319 (2)	1.330	-0.011	1.337	-0.018
	C1—C5	1.492 (2)	1.500	-0.008	1.445	0.047
	C1—C7	1.513 (2)	1.504	0.009	1.454	0.059
	C2—C3	1.514 (3)	1.507	0.007	1.507	0.007
	C3—C4	1.518 (3)	1.550	-0.032	1.540	-0.022
	C5—C6	1.553 (2)	1.560	-0.007	1.576	-0.023
	C6—C7	1.585 (2)	1.581	0.004	1.592	-0.007
	C6—C11	1.524 (2)	1.536	-0.012	1.555	-0.031
	C7—C8	1.528 (2)	1.531	-0.003	1.553	-0.025
	C8—C9	1.528 (3)	1.532	-0.004	1.532	-0.004
	C9—C10	1.517 (3)	1.533	-0.016	1.531	-0.014
	C10—C11	1.528 (3)	1.531	-0.003	1.529	-0.001
$\theta_{\max} = 70.24^\circ$	C7—S1—C12	99.7 (1)	100.8	-1.1	98.4	1.3
$h = -13 \rightarrow 13$	C4—O2—C5	106.8 (1)	108.3	-1.5	108.9	-2.1
$k = -10 \rightarrow 9$	C5—C1—C7	92.3 (1)	94.3	-2.0	99.1	-6.8
$l = -1 \rightarrow 8$	C2—C1—C7	139.3 (1)	139.2	0.1	130.5	8.8
1 standard reflection	C2—C1—C5	124.1 (1)	126.3	-2.2	124.4	-0.3
monitored every 50	C1—C2—C3	117.6 (1)	116.3	1.3	118.4	-0.8
reflections	C2—C3—C4	112.8 (2)	113.5	-0.7	112.2	0.6
intensity variation: within						
statistical fluctuation						

O2—C4—C3	113.5 (1)	112.6	0.9	110.9	2.6	226 parameters	Extinction coefficient:
O2—C5—C1	113.8 (1)	108.0	5.8	110.8	3.0	All H-atom parameters	0.001 (0)
C1—C5—C6	87.3 (1)	83.9	3.4	85.5	1.8	refined	Atomic scattering factors
O2—C5—C6	119.6 (1)	120.2	-0.6	119.3	0.3	$w = 1/[\sigma^2(F_o^2) + (0.0598P)^2]$	from International Tables
O1—C6—C5	112.4 (1)	110.7	1.7	112.5	-0.1	+ 3.1057P]	for Crystallography (1992,
C5—C6—C11	113.3 (1)	114.6	-1.3	112.3	1.0	where $P = (F_o^2 + 2F_c^2)/3$	Vol. C, Tables 4.2.6.8,
C5—C6—C7	87.3 (1)	89.0	-1.7	88.3	-1.0	$(\Delta/\sigma)_{\text{max}} < 0.001$	6.1.1.4)
O1—C6—C11	112.6 (1)	110.9	1.7	111.2	1.4		
O1—C6—C7	116.1 (1)	112.9	3.2	113.6	2.5		
C7—C6—C11	112.8 (1)	117.1	-4.3	117.2	-4.4		
C1—C7—C6	85.5 (1)	83.0	2.5	84.6	0.9		
S1—C7—C6	107.4 (1)	114.0	-6.6	112.0	-4.6		
S1—C7—C1	109.8 (1)	110.2	-0.4	109.7	0.1		
C6—C7—C8	119.4 (1)	116.4	3.0	116.4	3.0		
C1—C7—C8	120.4 (1)	117.4	3.0	117.9	2.5		
S1—C7—C8	111.4 (1)	112.6	-1.2	113.1	-1.7		
C7—C8—C9	112.5 (1)	115.0	-2.5	114.7	-2.2		
C8—C9—C10	111.1 (2)	110.4	0.7	109.4	1.7	x	$U_{\text{eq}} = (1/3)\sum_i \sum_j U_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j$
C9—C10—C11	109.5 (2)	108.7	0.8	108.2	1.3	y	
C6—C11—C10	113.2 (1)	111.7	1.5	112.9	0.3	z	
C12—S1—C7—C1	-76.9 (1)	-73.0	-3.9	-70.8	-6.1	U_{eq}	
C12—S1—C7—C6	-168.4 (1)	-164.3	-4.1	-163.0	-5.4	O1	0.8452 (2)
C12—S1—C7—C8	59.0 (1)	60.2	-1.2	63.0	-4.0	O2	0.7566 (2)
C4—O2—C5—C6	-153.5 (1)	-154.4	0.9	-154.9	1.4	O3	0.7777 (2)
C2—C1—C7—S1	119.1 (2)	95.7	23.4	108.2	10.9	O4	0.7127 (2)
C2—C1—C7—C8	-12.3 (3)	-35.1	22.8	-23.2	10.9	C1	0.7935 (2)
C5—C1—C2—C3	8.3 (3)	-3.0	11.3	4.2	C2	0.7316 (2)	
O2—C5—C6—O1	-106.7 (2)	-115.6	8.9	-122.5	15.8	C3	0.6690 (3)
O2—C5—C6—C7	136.2 (1)	130.2	6.0	122.7	13.5	C4	0.6910 (3)
O2—C5—C6—C11	22.4 (2)	10.7	11.7	3.9	C5	0.8110 (2)	
O1—C6—C7—S1	-24.4 (2)	-25.8	1.4	-15.9	-8.5	C6	0.1319 (4)
O1—C6—C7—C8	103.8 (2)	107.9	-4.1	116.4	-12.6	C7	0.1927 (3)

Distances: $|\Delta|_{\text{max}}$ MMX = 0.032 Å, MM+ = 0.059 Å; angles: $|\Delta|_{\text{max}}$ MMX = 6.6°, MM+ = 8.8°; torsions: $|\Delta|_{\text{max}}$ MMX = 23.4°, MM+ = 18.5°.

Compound (10)

Crystal data



$M_r = 228.29$

Orthorhombic

Pcab (No. 61)

$a = 18.992 (7)$ Å

$b = 10.595 (3)$ Å

$c = 11.875 (5)$ Å

$V = 2389 (2)$ Å³

$Z = 8$

$D_x = 1.269$ Mg m⁻³

Cu K α radiation

$\lambda = 1.54178$ Å

Cell parameters from 30

reflections

$\theta = 24\text{--}40^\circ$

$\mu = 0.773$ mm⁻¹

$T = 293 (2)$ K

Thin tablets

0.41 × 0.33 × 0.16 mm

Colourless

Data collection

Siemens AED diffractometer

$\theta/2\theta$ scans

Absorption correction:

none

4462 measured reflections

2277 independent reflections

2001 observed reflections

[$I > 2\sigma(I)$]

$R_{\text{int}} = 0.0462$

$\theta_{\text{max}} = 70.20^\circ$

$h = -23 \rightarrow 23$

$k = -1 \rightarrow 12$

$l = 0 \rightarrow 14$

1 standard reflection

monitored every 50

reflections

intensity variation: within statistical fluctuation

Refinement

Refinement on F^2

$R(F) = 0.1071$

$wR(F^2) = 0.2100$

$S = 1.265$

2277 reflections

$\Delta\rho_{\text{max}} = 0.241$ e Å⁻³

$\Delta\rho_{\text{min}} = -0.211$ e Å⁻³

Extinction correction:

SHELXL93 (Sheldrick, 1993)

Table 6. Comparison of bond distances (Å), bond angles (°) and selected torsion angles (°) in the molecule of compound (10)

E.s.d.'s in parentheses. MMX = force field of PCMODEL; MM+ = force field of HYPERCHEM; Δ = obs. — calc.

	X-ray obs.	MMX calc.	MM+ calc.	MM+ Δ
O1—C6	1.424 (5)	1.414	0.010	1.436
O2—C4	1.432 (6)	1.422	0.010	1.411
O2—C5	1.418 (5)	1.418	0.000	1.437
O3—C1	1.427 (5)	1.412	0.015	1.431
O4—C2	1.434 (6)	1.414	0.020	1.404
C1—C2	1.519 (6)	1.533	-0.014	1.547
C1—C5	1.524 (6)	1.553	-0.029	1.562
C1—C7	1.571 (6)	1.563	0.008	1.577
C2—C3	1.538 (7)	1.539	-0.001	1.536
C3—C4	1.519 (9)	1.542	-0.023	1.540
C5—C6	1.541 (6)	1.552	-0.011	1.561
C6—C7	1.565 (6)	1.562	0.003	1.574
C6—C11	1.531 (7)	1.531	0.000	1.547
C7—C8	1.546 (7)	1.545	0.001	1.561
C7—C12	1.520 (7)	1.533	-0.013	1.551
C8—C9	1.494 (8)	1.532	-0.038	1.531
C9—C10	1.501 (12)	1.534	-0.033	1.533
C10—C11	1.525 (10)	1.530	-0.005	1.530
C4—O2—C5	109.4 (3)	110.1	-0.7	109.8
O3—C1—C2	109.7 (3)	109.6	0.1	110.7
O3—C1—C5	112.6 (3)	110.4	2.2	112.7
O3—C1—C7	117.5 (3)	114.4	3.1	115.7
C2—C1—C5	109.6 (3)	110.5	-0.9	108.4
C2—C1—C7	117.6 (3)	121.6	-4.0	117.9
C5—C1—C7	87.8 (3)	88.0	-0.2	89.4
O4—C2—C1	110.3 (3)	110.3	0.0	111.3
O4—C2—C3	109.9 (4)	110.4	-0.5	109.6
C3—C2—C1	108.6 (4)	108.7	-0.1	108.9
C2—C3—C4	113.3 (5)	111.8	1.5	111.7
O2—C4—C3	110.7 (4)	110.8	-0.1	110.8
O2—C5—C1	116.5 (3)	113.1	3.4	113.6
O2—C5—C6	116.8 (3)	113.7	3.1	114.0
C1—C5—C6	90.2 (3)	88.3	1.9	89.2
O1—C6—C5	110.3 (3)	110.8	-0.5	109.2
O1—C6—C7	109.6 (3)	111.3	-1.7	111.7

O1—C6—C11	108.9 (3)	108.6	0.3	109.3	-0.4
C5—C6—C7	87.4 (3)	88.1	-0.7	89.6	-2.2
C5—C6—C11	119.6 (4)	117.9	1.7	117.2	2.4
C7—C6—C11	119.3 (4)	118.8	0.5	118.5	0.8
C1—C7—C6	87.6 (3)	87.6	0.0	88.2	-0.6
C1—C7—C8	111.2 (3)	113.5	-2.3	111.0	0.2
C1—C7—C12	119.2 (4)	115.4	3.8	116.9	2.3
C6—C7—C8	111.0 (4)	112.7	-1.7	114.0	-3.0
C6—C7—C12	116.4 (4)	115.2	1.2	114.4	2.0
C8—C7—C12	109.8 (4)	110.7	-0.9	110.7	-0.9
C7—C8—C9	113.0 (4)	113.1	-0.1	112.9	0.1
C8—C9—C10	109.6 (6)	108.7	0.9	108.4	1.2
C9—C10—C11	110.9 (6)	109.7	1.2	109.5	1.4
C6—C11—C10	113.2 (4)	115.1	-1.9	114.6	-1.4
C4—O2—C5—C1	40.8 (5)	52.5	-11.7	50.1	-9.3
C4—O2—C5—C6	145.6 (4)	148.9	-3.3	152.8	-7.2
O3—C1—C2—O4	-53.7 (4)	-55.8	2.1	-54.1	0.4
O3—C1—C5—O2	-100.8 (4)	-108.4	7.6	-112.1	11.3
O3—C1—C7—C8	-22.5 (5)	-18.8	-3.7	-14.0	-8.5
C2—C1—C7—C12	-27.8 (6)	-24.9	-2.9	-20.0	-7.8
O4—C2—C3—C4	153.3 (4)	160.2	-6.9	165.7	-12.4
O2—C5—C6—O1	-30.1 (5)	-16.9	-13.2	-23.7	-6.4
O2—C5—C6—C11	97.3 (5)	102.3	-5.0	108.0	-10.7
C1—C5—C6—O1	89.9 (3)	90.9	-1.0	98.7	-8.8
O1—C6—C7—C1	-91.2 (3)	-90.6	-0.6	-96.6	5.4
O1—C6—C7—C12	30.4 (5)	26.5	3.9	22.3	8.1
C5—C6—C7—C8	-92.3 (4)	-93.6	1.3	-98.3	6.0

Distances: $|\Delta|_{\max}$ MMX = 0.038 Å, MM+ = 0.038 Å; angles: $|\Delta|_{\max}$ MMX = 3.8°, MM+ = 3.0°; torsions: $|\Delta|_{\max}$ MMX = 9.3°, MM+ = 13.2°.

Table 7. Conformation of the rings and dihedral angles (°) of ring planes

Q_T = total puckering amplitude (Å) (Cremer & Pople, 1975), DAP = minimum displacement asymmetry parameter (Nardelli, 1983b), HC = half chair, S = sofa, T = twist.

Ring	Q_T	DAP	Conforma-	A/B	B/C	A/C
			tion			
Compound (8)						
A	0.522 (3)	$D_2(C_3—C_2) = 0.024$ (1)	HC			
B	-0.0615 (6)			151.8 (1)	110.6 (1)	107.3 (1)
C	0.007 (2)					
Compound (9)						
A	0.508 (2)	$D_2(C_1—C_2) = 0.0546$ (8)	HC			
B	-0.0720 (5)			154.4 (1)	118.4 (1)	92.8 (1)
C	0.523 (2)	$DS(C_7) = 0.040$ (2)	S/HC			
		$D_2(C_6—C_7) = 0.037$ (1)				
Compound (10)						
A	0.749 (4)	$D_2(C_4—O_2) = 0.042$ (2)	T			
		$D_2(C_3) = 0.032$ (1)				
B	0.070 (1)			135.2 (2)	117.7 (2)	149.7 (2)
C	0.524 (6)	$DS(C_6) = 0.013$ (3)	S/T			
		$D_2(C_6—C_11) = 0.073$ (2)				
		$D_2(C_7—C_6) = 0.055$ (2)				

Table 8. Hydrogen-bonding geometry (Å, °)

$D—H···A$	$D—H$	$H···A$	$D···A$	$D—H···A$
Compound (8)				
O1—H1O···O2 ⁱ	0.88 (3)	2.11 (3)	2.929 (3)	154 (3)
Compound (9)				
O1—H1O···S1 ⁱⁱ	0.83 (3)	2.51 (3)	3.321 (1)	164 (2)
Compound (10)				
O1—H1O···O2	0.86 (6)	2.45 (5)	2.759 (4)	101 (4)
O3—H3O···O4 ⁱⁱⁱ	0.81 (6)	2.38 (6)	2.778 (4)	111 (5)
C2—H2···O1	0.99 (4)	2.33 (4)	3.037 (6)	128 (3)
O1—H1O···O4 ⁱⁱⁱ	0.86 (6)	2.04 (6)	2.838 (5)	153 (5)
O3—H3O···O2 ^{iv}	0.81 (6)	2.04 (6)	2.713 (4)	140 (6)
O4—H4O···O3 ^v	0.83 (9)	2.02 (9)	2.817 (4)	160 (9)
C5—H5···O1 ^{vi}	0.92 (5)	2.35 (5)	3.209 (5)	155 (4)

Symmetry codes: (i) $\frac{1}{2} - x, 1 - y, \frac{1}{2} + z$; (ii) $1 - x, -y, 1 - z$; (iii) $\frac{3}{2} - x, y, \frac{1}{2} + z$; (iv) $\frac{3}{2} - x, y, z - \frac{1}{2}$; (v) $\frac{3}{2} - x, \frac{1}{2} + y, -z$; (vi) $x, y - \frac{1}{2}, \frac{1}{2} - z$.

Table 9. Averages of the absolute differences $|\Delta|$

Distances (Å)	$ \Delta _{MMX}$	(8)		(9)		(10)	
		Ave*	Ave*	Ave*	Ave*	Ave*	Ave*
$ \Delta _{MM+}$	0.024	0.023	0.031	0.009	0.017	0.019	0.011
$ \Delta _{MM+}$	2.2	1.8	2.1	1.6	1.2	0.9	
$ \Delta _{MM+}$	1.7	2.1	2.1	1.3			
$ \Delta _{MMX}$	14.2	10.8	5.1	6.5	6.4	4.8	
$ \Delta _{MM+}$	10.3	7.8	7.8				

* Ave = $\Delta(|MMX| - |MM+|)$.

The integrated intensities were obtained by a modified version (Belletti, Uguzzoli, Cantoni & Pasquinelli, 1979) of the Lehmann & Larsen (1974) peak-profile analysis procedure. All reflections of the three compounds were corrected for Lorentz and polarization effects but not for absorption.

The three structures were solved by direct methods and refined by anisotropic full-matrix least squares. In all three cases, the H atoms were found from $\Delta\rho$ maps and refined isotropically. The crystals of compound (10) were of poor quality so the R index is not as good as for the other two compounds. Nevertheless, refinement proceeded smoothly for (10) (comprehensive of H atoms) giving a quite acceptable molecular geometry and anisotropic displacement parameters (see Tables 5–8 and *Comment*). Calculations were carried out on the ENCORE91 and GOULD-POWERNODE 6040 computers of the Centro di Studio per la Strutturistica Diffattometrica del CNR (Parma), and on a COMPAQ-486c portable computer.

For all compounds, data collection: local programs; cell refinement: LOPARM (Nardelli & Mangia, 1984); data reduction: local programs. Program(s) used to solve structures: SHELLXS86 (Sheldrick, 1990) for (8) and (9); SIR92 (Altomare *et al.*, 1994) for (10). For all compounds, program(s) used to refine structures: SHELLXL93 (Sheldrick, 1993); molecular graphics: ORTEP (Johnson, 1965); software used to prepare material for publication: PARST (Nardelli, 1983a); PARSTCIF (Nardelli, 1991).

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Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: SZ1044). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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(\pm)-trans-2-Benzoylcyclohexanecarboxylic Acid. Structure and Hydrogen-Bonding Pattern of a γ -Keto Acid

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Abstract

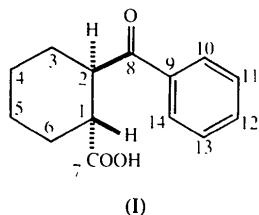
In the title compound, $C_{14}H_{16}O_3$, enantiomeric pairs of molecules form dimers by mutual hydrogen bonding of carboxyl groups, with the ketone group not involved in the hydrogen bonding. The components of the dimer differ in conformation, so that the dimer is the asymmetric unit.

Comment

Our interest in simple keto carboxylic acids arises from their four known crystalline hydrogen-bonding states. The commonest retains the dimer pattern characteristic of almost all functionally unelaborated acids in which the ketone is not involved. Less commonly, intermolec-

ular carboxyl-to-ketone hydrogen bonds repeat infinitely along one cell axis to yield a chain or catemer. A third rare arrangement is an internal hydrogen bond and one instance is known of acid-to-ketone dimerization. We have previously referenced and discussed many examples of these (Thompson, Lalancette & Vanderhoff, 1992; Coté, Thompson & Lalancette, 1996).

γ -Keto acids constitute a class particularly rich in hydrogen-bonding types, with examples of dimers, internal hydrogen bonds and catemers of both the helical and translational type. Despite the expectation of an intramolecular hydrogen bond for the *cis* counterpart of the title compound (Pourahmady & Eisenbraun, 1982), it was found to crystallize as a centrosymmetric dimer (Choney, Holt, Pourahmady & Eisenbraun, 1983). We report here that the *trans* epimer, (I), also adopts



(I)

the common carboxylic acid dimer motif, but that the dimer components are crystallographically unrelated, leading to an unusually long cell dimension [40.748 (6) Å] for such a small molecule (17 non-H atoms).

Friedel–Crafts acylation of benzene with *cis*-1,2-cyclohexanedicarboxylic anhydride yields, as the immediate product, the less stable *cis*-keto acid (m.p. 413 K). Aqueous base epimerizes this to the more stable *trans* diastereomer (m.p. 427 K) (Scribner & Miller, 1965).

Fig. 1 presents a view of the asymmetric unit with its numbering scheme. The cell contains four asymmetric units ($Z = 8$), each consisting of a pair of conformationally distinct molecules of opposite chirality, reciprocally hydrogen bonded through their carboxyl groups, with no ketone participation in the hydrogen bonding. Each molecule, as expected, has its substituents equatorial to a chair cyclohexane and the two molecules in the asymmetric unit differ almost entirely in the rotational orientation of the benzene ring. The carboxyl group adopts a conformation relative to its ring H atom whose torsion angle is 31.9° for molecule A and 30.1° for molecule B. The torsion angle for the benzoyl C=O relative to its axial ring H atom is 138.9° for molecule A and 141.0° for molecule B. Hence, the dihedral angles between the ketone and carboxyl planes are nearly identical in the two molecules, 73.0° in A and 70.2° in B. In molecule A, however, the dihedral angle of the aromatic ring relative to the ketone is 31.9°, whereas this angle is 9.7° in molecule B.

The carboxyl C—O bond lengths and C—C—O angles are found to be only slightly disordered, although