

Sebacic acid

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Key indicators

Single-crystal X-ray study
 $T = 180$ K
 Mean $\sigma(\text{C}-\text{C}) = 0.002$ Å
 R factor = 0.051
 wR factor = 0.139
 Data-to-parameter ratio = 18.2

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The crystal structure of octane-1,8-dicarboxylic acid (sebacic acid), $\text{C}_{10}\text{H}_{18}\text{O}_4$, has been redetermined at 180 K. The molecular units are centrosymmetric and linked *via* the ubiquitous *syn-syn* carboxylic acid dimer to form infinite chains running along the $[\bar{1}01]$ vector.

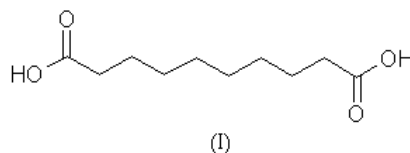
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Comment

Crystal data for sebacic acid, (I), have been reported on four previous occasions, by Caspari (1928), Morrison & Robertson (1949) (SEBAAC01), Housty & Hospital (1966) (SEBAAC) and Haget *et al.* (1980) (SEBAAC02 in the Cambridge Structural Database; Allen & Kennard, 1993). All four studies essentially report the same structure in space group $P2_1/c$ with unit-cell dimensions *ca* $a = 15.1$, $b = 5.0$, $c = 10.1$ Å, $\beta = 133.1^\circ$. [Morrison & Robertson (1949) describe the structure in the non-standard setting $P2_1/a$.] The most precise cell determination (SEBAAC02) to date is derived from powder X-ray diffraction data, and atomic coordinates were not determined. We have re-examined sebacic acid at 180 K and report the structure to significantly greater precision in $P2_1/c$ with β *ca* 92° . The previously reported structures in $P2_1/c$ with β *ca* 133.1° may be transformed to our structure by the matrix (101, 010, 001) followed by an origin shift of $(\frac{1}{2}, \frac{1}{2}, \frac{1}{2})$ to bring an entire molecule into the unit cell.



Experimental

Sebacic acid was obtained from Aldrich and recrystallized from ethanol.

Crystal data

$\text{C}_{10}\text{H}_{18}\text{O}_4$
 $M_r = 202.24$
 Monoclinic, $P2_1/c$
 $a = 10.9197$ (7) Å
 $b = 4.9876$ (6) Å
 $c = 9.964$ (1) Å
 $\beta = 92.273$ (6) $^\circ$
 $V = 542.27$ (9) Å 3
 $Z = 2$

$D_x = 1.239$ Mg m $^{-3}$
 Mo $K\alpha$ radiation
 Cell parameters from 2040 reflections
 $\theta = 1.0$ – 27.5°
 $\mu = 0.10$ mm $^{-1}$
 $T = 180$ (2) K
 Plate, colourless
 $0.23 \times 0.18 \times 0.05$ mm

Data collection

Nonius KappaCCD diffractometer
Thin-slice ω and φ scans
Absorption correction: multi-scan
(*SORTAV*; Blessing, 1995)
 $T_{\min} = 0.912$, $T_{\max} = 0.995$
3745 measured reflections
1241 independent reflections

796 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.054$
 $\theta_{\text{max}} = 27.5^\circ$
 $h = -11 \rightarrow 14$
 $k = -5 \rightarrow 6$
 $l = -12 \rightarrow 9$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.051$
 $wR(F^2) = 0.139$
 $S = 1.05$
1241 reflections
68 parameters

H atoms treated by a mixture of
independent and constrained
refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0705P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.006$
 $\Delta\rho_{\text{max}} = 0.22 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.18 \text{ e } \text{\AA}^{-3}$

Table 1

Hydrogen-bonding geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$O2-H1\cdots O1^i$	1.06 (3)	1.58 (3)	2.6406 (15)	175 (2)

Symmetry code: (i) $-x, 1-y, 2-z$.

All H atoms (apart from H1) were placed geometrically and allowed to ride during subsequent refinement with an isotropic displacement parameter fixed at 1.2 times that for the C atom to which they are attached. H1 was refined without restraint.

Data collection: *COLLECT* (Nonius, 1998); cell refinement: *HKL SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *HKL DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* (Sheldrick, 1993) and *CAMERON* (Watkin *et al.*, 1996); software used to prepare material for publication: *SHELXL97*.

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References

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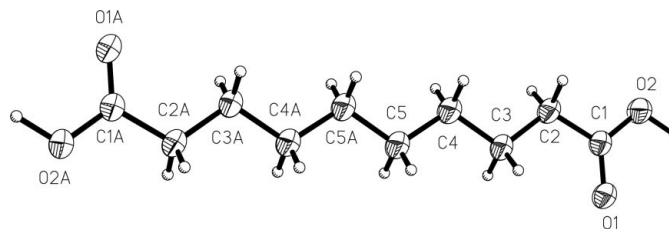


Figure 1

The molecular unit of the title compound showing displacement ellipsoids at the 50% probability level.

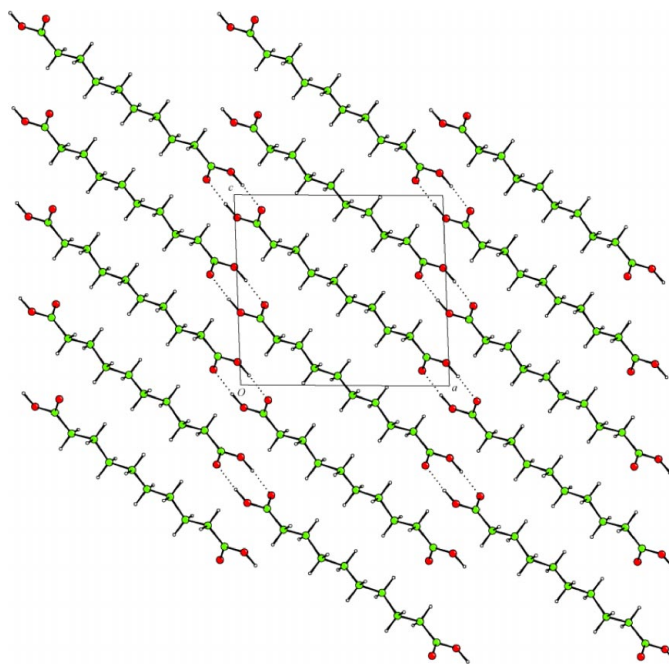


Figure 2

Projection onto (010) showing chains of sebacic acid running along the $[101]$ vector.

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