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Key indicators

Single-crystal X-ray study T = 150 KMean σ (C–C) = 0.005 Å R factor = 0.040 wR factor = 0.098 Data-to-parameter ratio = 9.1

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

(2S)-[(2S)-2-Hydroxy-2-phenylethanoyloxy]phenylacetic acid

The novel title compound, $C_{16}H_{14}O_5$, forms four intermolecular hydrogen bonds and takes part in two chains running along the shortest crystallographic *b* axis.

Comment

During structural studies of S-(+)-mandelic acid [S-(+)-MA], needle-like crystals of an unknown compound were isolated. The compound was shown to be the unusual 'ester' with the assumed stereochemistry (2S)-[(2S)-2-hydroxy-2-phenyl-ethanoyloxy]phenylacetic acid, (I). The molecular structure of (I) is shown in Fig. 1 and the unit-cell contents in Fig. 2.



Each molecule is involved in four intermolecular hydrogen bonds to four symmetry-related molecules, as shown in Fig. 3. These hydrogen bonds form a network comprising two distinct $C_1^1(5)$ and $C_1^1(4)$ catemeric chains along the shortest crystallographic *b* axis. The $C_1^1(5)$ chain connects the 'ester' carbonyl of one molecule to the alcohol hydroxyl group of the next molecule. The $C_1^1(4)$ chain arises from $-C=O\cdots H-O$ hydrogen bonding through adjacent carboxylic acid groups. Fig. 4 shows the hydrogen-bonding scheme giving rise to the two chains, and details are given in Table 1.

Experimental

Saturated solutions of S-(+)-MA (supplied by Sigma Aldrich, 99+%) in distilled water were prepared and rhombic crystals of S-(+)-MA were obtained. The mother liquors were allowed to evaporate slowly under ambient conditions. After several weeks, needle-like crystals of the title compound were isolated as the minor product.



Figure 1

© 2004 International Union of Crystallography Printed in Great Britain – all rights reserved The molecular structure of the title compound, (I), showing the labelling scheme, with displacement ellipsoids drawn at the 30% probability level.

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 $D_x = 1.317 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation Cell parameters from 4075

reflections $\theta = 1.0-27.5^{\circ}$

 $\mu = 0.10~\mathrm{mm}^{-1}$

Needle, colourless $0.30 \times 0.15 \times 0.10$ mm

1413 reflections with $I > 2\sigma(I)$

 $w = 1/[\sigma^2(F_o^2) + (0.0442P)^2]$

Extinction correction: SHELXL97

Extinction coefficient: 0.115 (14)

+ 0.1103P] where $P = (F_o^2 + 2F_c^2)/3$

 $(\Delta/\sigma)_{\rm max} < 0.001$

 $\Delta \rho_{\rm max} = 0.20 \ {\rm e} \ {\rm \AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.21 \text{ e } \text{\AA}^{-3}$

T = 150 K

 $R_{\rm int}=0.040$

 $\theta_{\rm max} = 27.3^{\circ}$

 $\begin{array}{l} h = -11 \rightarrow 13 \\ k = -7 \rightarrow 8 \end{array}$

 $l = -12 \rightarrow 15$

Figure 2

The unit-cell contents of (I), viewed along b.

Crystal data

 $\begin{array}{l} C_{16}H_{14}O_5\\ M_r = 286.27\\ \text{Monoclinic, } P2_1\\ a = 10.5196 \ (4) \ \text{\AA}\\ b = 6.4129 \ (3) \ \text{\AA}\\ c = 11.7201 \ (7) \ \text{\AA}\\ \beta = 114.107 \ (2)^\circ\\ V = 721.69 \ (6) \ \text{\AA}^3\\ Z = 2 \end{array}$

Data collection

Nonius KappaCCD diffractometer Thick-slice ω scans Absorption correction: multi-scan (Blessing, 1995) $T_{\min} = 0.916, T_{\max} = 0.990$ 4184 measured reflections 1760 independent reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.040$ $wR(F^2) = 0.098$ S = 1.031760 reflections 193 parameters H-atom parameters constrained

Table 1

Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$\begin{array}{c} O1 - H1 \cdots O2^{i} \\ O13 - H13 \cdots O12^{ii} \end{array}$	0.82	1.94	2.715 (3)	157
	0.82	2.04	2.817 (3)	159

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

The Friedel pairs were merged and the $\Delta f''$ term was set to zero. [The calculated absolute structure parameter (Flack, 1983) of 0.23 (13) was meaningless in this analysis.] All H atoms attached to C atoms were positioned geometrically and refined as riding, with C–H = 0.93–0.98 Å and $U_{\rm iso}(\rm H) = 1.2U_{eq}(\rm C)$. The OH H atoms were positioned geometrically with torsion angles selected so as to maximise electron density and were refined as rotating groups, with O–H = 0.82 Å, and $U_{\rm iso}(\rm H) = 1.5U_{eq}(\rm O)$.

Data collection: COLLECT (Nonius, 1997–2000); cell refinement: HKL SCALEPACK (Otwinowski & Minor, 1997); data reduction:





The hydrogen bonds (dashed lines) formed by each independent molecule with four neighbouring molecules.



Figure 4

The $C_1^1(5)$ (blue) and $C_1^1(4)$ (pink) hydrogen-bonded chains of the acid, viewed along *c*. Hydrogen bonds are indicated by dashed lines.

HKL DENZO (Otwinowski & Minor, 1997) and *SCALEPACK*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *CERIUS*² (Accelrys, 2001); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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