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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.006 Å R factor = 0.062 wR factor = 0.126 Data-to-parameter ratio = 8.6

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

2,5-Dioxopyrrolidin-1-yl (1*R*,2*S*,5*R*)-2-isopropyl-5-methylcyclohexyl carbonate

The title compound, $C_{15}H_{23}NO_5$, was obtained as colourless crystals by the reaction of *N*,*N'*-disuccinimidyl carbonate (DSC) and L-(-)-menthol. The molecular packing is stabilized by weak C-H···O hydrogen-bonding interactions.

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Comment

The title compound, (I), is a key intermediate used in the synthesis of a series of biologically active polyfunctional molecules for enzyme active sites (Ghosh *et al.*, 1992). It can react with various chiral alcohols and amines to yield active esters of interest in medicinal chemistry. The absolute configuration of the three stereocentres remains unchanged during the synthetic procedure. The molecular packing is stabilized by weak $C-H \cdots O$ hydrogen-bonding interactions (Table 1).



Experimental

N,N'-Disuccinimidyl carbonate (DSC), was prepared according to the procedure of Pereira *et al.* (1998). To a solution of DSC (1.5 mmol, 0.384 g) and L-(-)-menthol (1.0 mmol, 0.156 g) in dried acetonitrile (10 ml) at room temperature was added triethylamine (3 mmol) dropwise. The resulting mixture was stirred at room temperature



Figure 1

The molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are represented by spheres of arbitrary radii.

© 2006 International Union of Crystallography All rights reserved until no starting L-(-)-menthol remained, as monitored by thin-layer chromatography (8 h). The mixture was concentrated under reduced pressure and the residue was diluted with a saturated aqueous NaHCO₃ solution (10 ml) and extracted thoroughly with ethyl acetate (3×20 ml). The combined extracts were washed with a saturated aqueous NaCl solution (8 ml) and dried over Na₂SO₄. Evaporation of the solvent provided the mixed carbonate (I) which was purified by column chromatography over silica gel (petroleum ether/ethyl acetate 5:1 ν/ν) to obtain a white solid. Single crystals were obtained by recrystallization from a mixture of petroleum ether/ ethyl acetate (5:1 ν/ν).

Crystal data

 $C_{15}H_{23}NO_5$ $M_r = 297.34$ Orthorhombic, $P2_12_12_1$ a = 6.5911 (14) Å b = 7.7815 (17) Å c = 31.125 (7) Å $V = 1596.4 (6) \text{ Å}^3$

Data collection

Bruker APEX area-detector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Bruker, 2001) $T_{\min} = 0.964, T_{\max} = 0.982$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.062$ $wR(F^2) = 0.127$ S = 1.391652 reflections 193 parameters H-atom parameters constrained Z = 4 $D_x = 1.237 \text{ Mg m}^{-3}$ Mo K\alpha radiation $\mu = 0.09 \text{ mm}^{-1}$ T = 293 (2) KBlock, colourless $0.40 \times 0.27 \times 0.20 \text{ mm}$

7845 measured reflections 1652 independent reflections 1621 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.028$ $\theta_{\text{max}} = 25.0^{\circ}$

$$\begin{split} w &= 1/[\sigma^2(F_{\rm o}{}^2) + (0.0276P)^2 \\ &+ 0.8697P] \\ \text{where } P &= (F_{\rm o}{}^2 + 2F_{\rm c}{}^2)/3 \\ (\Delta/\sigma)_{\rm max} &= 0.004 \\ \Delta\rho_{\rm max} &= 0.18 \text{ e } \text{\AA}{}^{-3} \\ \Delta\rho_{\rm min} &= -0.18 \text{ e } \text{\AA}{}^{-3} \end{split}$$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C1-H1\cdots O5^{i}$	0.98	2.51	3.460 (5)	164
$C10-H10A\cdotsO1^{ii}$	0.97	2.41	3.255 (5)	145
Symmetry codes: (i) x –	1, y, z; (ii) $x +$	$\frac{1}{2}, -y + \frac{5}{2}, -z - z$	+ 2.	

All H atoms were placed in geometrically idealized positions and treated as riding on their parent atoms, with C–H = 0.98 (CH), 0.97 (CH₂) or 0.96 Å (CH₃), and $U_{\rm iso}(\rm H) = 1.2~U_{eq}(\rm CH~and~CH_2)$ or 1.5 $U_{\rm eq}(\rm methyl~C)$. In the absence of significant anomalous scattering effects, Friedel pairs were merged; the absolute configuration was assumed from the synthesis.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*.

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