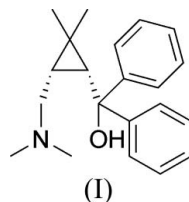


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Key indicators

Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.005$ Å
 R factor = 0.037
 wR factor = 0.076
Data-to-parameter ratio = 8.0For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.**{(1*R*,3*S*)-3-[(Dimethylamino)methyl]-2,2-dimethyl-
cyclopropyl}diphenylmethanol**The title compound, $\text{C}_{21}\text{H}_{27}\text{NO}$, crystallizes with two rather
similar molecules in the asymmetric unit. Its molecular
conformation is stabilized by an intramolecular $\text{O}-\text{H}\cdots\text{N}$
hydrogen bond.Received 26 January 2007
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Comment

Cyclopropane occurs as a basic structural motif in a wide
range of naturally occurring compounds (Patai & Rappoport,
1987). Recently, Molander *et al.* (2004) used chiral cyclo-
propane-based ligands in the palladium-catalyzed allylic
alkylation of 1,3-diphenylpropenyl acetate with dimethyl
malonate. In the light of the above report and our interest in
devising new chiral cyclopropane-based ligands, the crystal
structure of the title compound, (I), has been determined.There are two rather similar molecules in the asymmetric
unit of (I), which differ slightly in the orientation of the phenyl
rings. The molecular conformation is stabilized by an intra-
molecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bond.

Experimental

Compound (I) was prepared according to the method of Zhong *et al.*
(2006). Single crystals of (I) suitable for X-ray data collection were
obtained by recrystallization from *n*-hexane.

Crystal data

$\text{C}_{21}\text{H}_{27}\text{NO}$	$Z = 4$
$M_r = 309.44$	$D_x = 1.114$ Mg m ⁻³
Monoclinic, $P2_1$	Mo $K\alpha$ radiation
$a = 6.2697$ (13) Å	$\mu = 0.07$ mm ⁻¹
$b = 16.168$ (3) Å	$T = 293$ (2) K
$c = 18.260$ (4) Å	Plate, colourless
$\beta = 94.60$ (3)°	$0.50 \times 0.45 \times 0.12$ mm
$V = 1845.0$ (6) Å ³	

Data collection

Rigaku R-Axis RAPID image- plate diffractometer	10713 measured reflections
ω scans	3389 independent reflections
Absorption correction: multi-scan (<i>ABSCOR</i> ; Higashi, 1995)	2089 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.967$, $T_{\max} = 0.992$	$R_{\text{int}} = 0.045$
	$\theta_{\text{max}} = 25.0^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.076$
 $S = 0.99$
 3389 reflections
 424 parameters
 H atoms treated by a mixture of
 independent and constrained
 refinement

$w = 1/[\sigma^2(F_o^2) + (0.027P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.16 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -0.13 \text{ e } \text{Å}^{-3}$
 Extinction correction: *SHELXL97*
 (Sheldrick, 1997)
 Extinction coefficient: 0.074 (2)

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1—HO1 \cdots N1	0.98 (4)	1.88 (4)	2.782 (4)	152 (3)
O2—HO2 \cdots N2	0.90 (4)	1.86 (4)	2.716 (4)	157 (3)

H atoms bonded to O were refined freely. The remaining H atoms were positioned geometrically, with $C-H = 0.93\text{--}0.98 \text{ Å}$, and refined using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. In the absence of significant anomalous scattering effects, Friedel pairs were merged.

Data collection: *RAPID-AUTO* (Rigaku, 2000); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku, 2000); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* in *SHELXTL* (Sheldrick, 1991); software used to prepare material for publication: *SHELXL97*.

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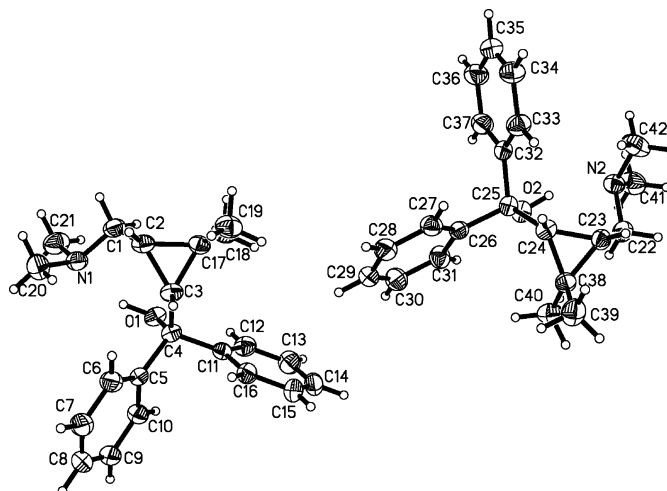


Figure 1

The asymmetric unit of (I), showing the atom-labelling scheme and with displacement ellipsoids drawn at the 30% probability level.

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