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Key indicators

Single-crystal X-ray study T = 295 K Mean σ (C–C) = 0.004 Å R factor = 0.043 wR factor = 0.130 Data-to-parameter ratio = 9.7

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

2-(2,6-Diisopropylphenylamino)cyclohex-1-enyl phenyl ketone

The two independent molecules of the title compound, $C_{25}H_{31}NO$, are linked by a pair of weak hydrogen bonds across a non-crystallographic inversion center in the monoclinic space group *Cc*.

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Comment

A previous report (Gao *et al.*, 2005) describes the structure of a sterically crowded Schiff base; the compound features diisopropyl substituents in the 2,6-positions of the aromatic ring that is bonded to the imino N atom. The condensation of 2,6-diisopropylphenylamine with 2-benzoylcyclohexanone probably yielded the expected Schiff base; however, the imine had isomerized to an amine, (I) (Fig. 1). The same amine was obtained even if the ratio of the amine reagent was raised to 3:1. The two independent molecules are linked by a pair of weak hydrogen bonds across a non-crystallographic inversion center (Table 1 and Fig. 2).



Experimental

2-Benzoylcyclohexanone (3.10 g, 0.015 mol), 2,6-diisopropylphenylamine (3.00 ml, 0.015 mol) and a catalytic amount of *p*-toluene-



Figure 1

© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved *ORTEPII* plot (Johnson, 1976) of one of the two independent molecules of (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as spheres of arbitrary radii.



Figure 2

ORTEPII plot (Johnson, 1976) illustrating the arrangement of the two independent molecules across a non-crystallographic center of inversion.

sulfonic acid were heated in toluene (50 ml) in a Dean–Stark water separator. When the water was removed after about a day, the solvent was evaporated to give a product that was recrystallized from hexane in about 80% yield (m.p. 388 K). Elemental analysis calculated for $C_{25}H_{31}NO$: C 82.98, H 8.64, N 3.87%; found: C 82.86, H 8.76, N 3.75%.

Crystal data

C ₂₅ H ₃₁ NO	$D_x = 1.114 \text{ Mg m}^{-3}$
$M_r = 361.51$	Mo $K\alpha$ radiation
Monoclinic, Cc	Cell parameters from 946
a = 9.815(1) Å	reflections
b = 23.404 (3) Å	$\theta = 1.7 - 27.1^{\circ}$
c = 18.777 (2) Å	$\mu = 0.07 \text{ mm}^{-1}$
$\beta = 92.002 \ (2)^{\circ}$	T = 295 (2) K
$V = 4310.5 (9) \text{ Å}^3$	Block, pale yellow
Z = 8	$0.50 \times 0.48 \times 0.45 \text{ mm}$
Data collection	
Bruker SMART area-detector diffractometer	4021 reflections with $I > 2\sigma(I)$ $R_{int} = 0.015$

 φ and ω scans Absorption correction: none 13505 measured reflections 4703 independent reflections 4021 reflections with $I > 2\sigma$ $R_{\text{int}} = 0.015$ $\theta_{\text{max}} = 27.1^{\circ}$ $h = -12 \rightarrow 12$ $k = -29 \rightarrow 30$

 $l = -23 \rightarrow 12$

n	C .
RO	tinomont
ne	111101110111

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0842P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.043$	+ 0.6097P]
$wR(F^2) = 0.130$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.03	$(\Delta/\sigma)_{\rm max} = 0.001$
4703 reflections	$\Delta \rho_{\rm max} = 0.27 \ {\rm e} \ {\rm \AA}^{-3}$
487 parameters	$\Delta \rho_{\rm min} = -0.14 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Table 1Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	H···A	$D \cdots A$	$D - H \cdots A$
$\overline{N1-H1n\cdots O1}$	0.85	1.93	2.598 (2)	135
$N1-H1n\cdots O1'$	0.85	2.51	3.204 (3)	140
$N1'-H1'n\cdots O1'$ $N1'-H1'n\cdots O1$	0.85 0.85	1.96 2.51	2.614 (3) 3.206 (3)	133 140

In the absence of significant anomalous scattering effects, Friedel pairs were merged and the calculation of the Flack (1983) parameter was suppressed. The carbon-bound H atoms were placed at calculated positions (C-H = 0.93 Å aromatic H atoms, 0.97 Å for methylene H atoms and 0.96 Å for methyl H atoms) and were included in the refinement in the riding-model approximation, with $U_{\rm iso}({\rm H})$ values set at 1.2 times $U_{\rm eq}$ of the parent C atoms, except for the methyl H atoms, for which this was set at 1.5 times $U_{\rm eq}$. The nitrogen-bound H atoms were similarly treated [N-H = 0.85 Å and $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm N})$]

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: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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