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Key indicators

Single-crystal X-ray study T = 295 KMean σ (C–C) = 0.003 Å R factor = 0.048 wR factor = 0.144 Data-to-parameter ratio = 13.5

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

© 2006 International Union of Crystallography All rights reserved In the crystal structure of the title compound, $C_{20}H_{25}NO$, all bond lengths and angles show normal values. Non-classical hydrogen bonds are responsible for the deviation from a planar arrangement.

Comment

In the past few years, α -diimines and iminoketones have attracted interest as ligands in the synthesis of complexes used as catalysts in olefin polymerization reactions (Koten & Vrieze, 1980; Johnson *et al.*, 1995; Binotti *et al.*, 2004). Crystallographic studies of free iminoketones are relatively scarce compared to α -diimines. The present communication describes the chief characteristics of 2-(2,6-diisopropylphenylimino)-1-phenylpropan-1-one, (I).



The structure of (I) can be discussed in terms of its hydrogen-bonding scheme. The individual molecules show intramolecular non-classical hydrogen bonds including a trifurcated acceptor geometry (C6–H6, C16–H16, C19–H19)…N1 (Jeffrey *et al.*, 1985), with atom N1 located 0.27 (1) Å above the plane of the three H atoms (Fig. 1 and Table 2).

The non-classical hydrogen bonds $C19-H19\cdots N1$ and $C16-H16\cdots N1$ give rise to two five-membered rings sharing an edge (N1-C7 bond). The N1/C7/C8/C19/H19 ring is almost planar (r.m.s. deviation = 0.0764°), while the N1/C7/ C12/C16/H16 ring shows a small deviation from planarity (r.m.s. deviation = 0.1121 Å). An interplanar angle of 9.0 (1)° is observed between the rings. If we take into account the third non-classical hydrogen bond, *viz*. C6-H6···N1, additional consequences can also be inferred. The first is the formation of a six-membered ring (N1/C14/C13/C1/C6/H6), with a half-boat conformation. The second feature is the O1-C13-C14-N1 torsion angle [-150.7 (2)°], which is equivalent to an angle of 27.1 (1)° between the planes N1/C14/C15 and O1/C13/C1. Finally, we believe that this hydrogen bond is responsible for the $68.49 (3)^{\circ}$ deviation from coplanarity of the phenyl ring and the ring carrying the isopropyl groups.

The values found for the C–O and C–N distances (Table 1) agree well with what is expected for this bonding mode as described in the literature (*International Tables for Crystallography*, 1985, Vol. III, p. 270).

Experimental

2,6-Diisopropylaniline (2.85 g, 3.7 mmol) was added slowly at room temperature to a methanol solution (20 ml) of 1-phenylpropane-1,2dione (0.55 g, 3.7 mmol) in the presence of a catalytic amount of formic acid. After 6 h of magnetic stirring, the crude yellow product was isolated by filtration, washed with cold methanol (3×20 ml) and dried in vacuum. Yellow block-shaped crystals suitable for X-ray diffraction were obtained by recrystallization from a methanol solution (yield 0.79 g, 70%; m.p. 368 K).

Z = 4

 $D_x = 1.114 \text{ Mg m}^{-3}$ Mo K\alpha radiation $\mu = 0.07 \text{ mm}^{-1}$ T = 295 (2) K Block, yellow 0.26 \times 0.24 \times 0.15 mm

3408 independent reflections

 $R_{\rm int} = 0.027$

 $\theta_{\rm max} = 25.5^{\circ}$

2508 reflections with $I > 2\sigma(I)$

Crystal data

$C_{21}H_{25}NO$
$M_r = 307.42$
Monoclinic, $P2_1/n$
a = 9.9291 (4) Å
b = 18.5953 (6) Å
c = 10.4394 (4) Å
$\beta = 108.015 \ (2)^{\circ}$
$V = 1832.98 (12) \text{ Å}^3$

Data collection

Bruker APEX-II CCD areadetector diffractometer φ and ω scans Absorption correction: none 16097 measured reflections

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0598P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.048$	+ 0.2624P]
$wR(F^2) = 0.144$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.07	$(\Delta/\sigma)_{\rm max} = 0.001$
3408 reflections	$\Delta \rho_{\rm max} = 0.18 \ {\rm e} \ {\rm A}^{-3}$
253 parameters	$\Delta \rho_{\rm min} = -0.17 \text{ e } \text{\AA}^{-3}$
H atoms treated by a mixture of	
independent and constrained	
refinement	

Table 1

Selected geometric parameters (Å, °).

C13-O1 C13-C14	1.2147 (19) 1.519 (2)	C14-N1	1.2735 (19)
O1-C13-C1 O1-C13-C14 N1-C14-C15 N1-C14-C13	120.54 (14) 117.38 (15) 126.89 (14) 116.59 (14)	C15-C14-C13 C14-N1-C7 C14-N1-H6 C7-N1-H6	116.38 (13) 121.32 (13) 91.4 (4) 128.0 (4)



Figure 1

Perspective view of (I). Displacement ellipsoids are drawn at the 50% probability level.

Table 2			
Hydrogen-bond	geometry	(Å,	°).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C6-H6\cdots N1$	0.95 (2)	2.47 (2)	2.967 (2)	112.3 (15)
$C16-H16\cdots N1$	0.96 (2)	2.56 (2)	2.943 (2)	103.9 (13)
$C19-H19\cdots N1$	0.97 (2)	2.37 (2)	2.856 (2)	110.3 (14)

Phenyl and methine H atoms were located in a Fourier difference map and refined freely. Methyl H atoms were positioned geometrically, the C—H distances fixed (0.96 Å for Csp^3 H atoms), and the atoms refined as riding on their respective C atoms, with an isotropic displacement parameter 1.5 times U_{eq} for the attached Csp^3 atom. The methyl groups were refined as rigid groups, with rotation around the respective C—C bonds as a free variable.

Data collection: *COSMO* and *APEX2* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); 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: *WinGX* (Farrugia, 1999).

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