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Key indicators

Single-crystal X-ray study T = 292 KMean σ (C–C) = 0.003 Å R factor = 0.059 wR factor = 0.160 Data-to-parameter ratio = 16.3

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

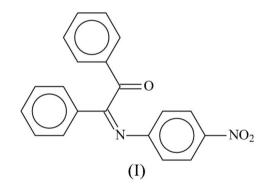
2-(4-Nitrophenylimino)-1,2-diphenylethanone

The O=C-C=N sequence of atoms in the title molecule [alternative name: 2,3-diphenyl-4-(4-nitrophenyl)-1-oxa-4-azabutadiene], $C_{20}H_{14}N_2O_3$, adopts a *gauche* conformation, with an O=C-C=N torsion angle of 88.9 (1)°.

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Comment

Dibenzoyl condenses with a primary amine to form a monoimino Schiff base such as, for example, 2-(phenylimino)-1,2diphenylethanone, $(C_6H_5)C(=O)C(C_6H_5)(=NC_6H_5)$; this compound has a molecular structure which contributes to the understanding of *peri*-selectivity in cycloadditions. In this compound, the ==C==O and ==C==N – functional groups are almost perpendicular to each other [O==C-C==N torsion angle = 84.5 (2)°; Guner *et al.*, 2000].



In the title molecular structure, (I) (Fig. 1), the addition of an electron-withdrawing nitro substituent to the phenylimino unit does not lead to a change either in the C—N distance [1.277 (2) Å, compared with 1.278 (2) Å found in the parent molecule] or in the bond lengths and angles in other parts of the molecule, although the O—C–C—N torsion angle is slightly different [O1–C7–C8–N1 torsion angle = 88.9 (1)°].

Experimental

4-Nitroaniline (2.00 g, 14.5 mmol) and dibenzoyl (2.99 g, 14.2 mmol) were dissolved in ethanol (35 ml) along with formic acid (1 ml). The solution was refluxed for 6 h. The solvent was then removed and the pure product obtained upon recrystallization from a 1:1 ethanol-chloroform mixture (35 ml) in 80% yield. Crystals of (I) were grown from ethanol as solvent. CHN elemental analysis: calculated for C₂₀H₁₄O₃N₂: C 72.72, H 4.27, N 8.48%; found: C 72.68, H 4.19, N 8.60%.

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Crystal data

 $C_{20}H_{14}N_2O_3$ $M_r = 330.33$ Monoclinic, $P2_1/c$ a = 8.8265 (7) Å b = 11.5590(9) Å c = 16.365 (1) Å $\beta = 101.323 (1)^{\circ}$ V = 1637.1 (2) Å³

Data collection

Bruker APEX CCD area-detector diffractometer φ and ω scans Absorption correction: none 8569 measured reflections

Refinement

II store consectors con
H-atom parameters con
$w = 1/[\sigma^2(F_o^2) + (0.086H)]$
where $P = (F_0^2 + 2F_c^2)$
$(\Delta/\sigma)_{\rm max} = 0.001$
$\Delta \rho_{\rm max} = 0.22 \ {\rm e} \ {\rm \AA}^{-3}$
$\Delta \rho_{\rm min} = -0.18 \text{ e} \text{ Å}^{-3}$

H atoms were placed in calculated positions, with C-H 0.93 Å and $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm C})$, and were included in the refinement in the riding-model approximation.

Data collection: SMART (Bruker, 2003); cell refinement: SAINT (Bruker, 2003); 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.

Z = 4 $D_x = 1.340 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation $\mu = 0.09 \text{ mm}^{-1}$ T = 292 (2) K Block, yellow $0.36 \times 0.23 \times 0.19 \text{ mm}$

3684 independent reflections 2574 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.059$ $\theta_{\rm max} = 27.5^{\circ}$

nstrained $(P)^{2}]_{c}^{2}/3$

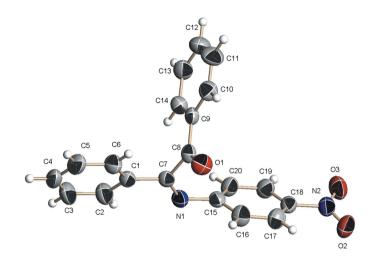


Figure 1

The molecular structure of (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as spheres of arbitrary radii.

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