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Key indicators

Single-crystal X-ray study T = 291 KMean $\sigma(C-C) = 0.002 \text{ Å}$ Disorder in main residue R factor = 0.038 wR factor = 0.120 Data-to-parameter ratio = 11.1

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

Benzylideneacetophenone 2,4-dinitrophenylhydrazone

The title Schiff base compound, $C_{21}H_{16}N_4O_4$, features a crystallographically imposed planar -NH-N linkage of an *E* configuration.

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Comment

2,4-Dinitrophenylhydrazine is a reagent that is used for the formation of crystalline derivatives of aldehydes and ketones, as it usually affords sharp-melting solids (Furniss *et al.*, 1989), and the crystal structures of a large number of these Schiff base hydrazones have been determined, as noted from the Cambridge Structural Database (Version 5.27; Allen, 2002). The majority of the aryl-aryl and aryl-alkyl ketone derivatives whose structures have been reported have substituents on the phenyl ring. There are few examples of phenyl-alkyl ketone derivatives, and these include only the derivatives of acet-ophenone, *viz.* the derivatives of acetophenone itself (Shan *et al.*, 2002*a*), phenyl ethyl ketone (Shan *et al.*, 2001).



In the title compound, (I), the aromatic ring of the dinitrophenylhydrazine unit and benzylidene portion of the benzylideneacetophenone unit lie on a mirror plane (Fig. 1). The structure is disordered, as the two nitro groups of the phenylhydrazine portion as well as the phenyl ring that is next to the -NH-N= unit, lie off the mirror plane; the phenyl ring is perpendicular to the mirror plane [dihedral angle 90.0 (4)°].

Experimental

2,4-Dinitrophenylhydrazine (0.20 g, 1 mmol) was dissolved in anhydrous ethanol (10 ml) and to the solution was added concentrated sulfuric acid (0.5 ml). The solution was heated to the boiling point of ethanol and then benzylideneacetophenone (0.21 g, 1 mmol) dissolved in ethanol (6 ml) was added. The mixture was then refluxed for several hours. After removing the solvent by evaporation, the product was collected and recrystallized from dichloromethane to yield orange crystals of (I). The preparation was first reported in 1937 (Dippy *et al.*, 1937).

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Figure 1

A plot of (I). Displacement ellipsoids are drawn at the 50% probability level. Only one disorder component is shown.

Z = 4

Crystal data

 $\begin{array}{l} C_{21}H_{16}N_4O_4\\ M_r = 388.38\\ Orthorhombic, Pnma\\ a = 18.570 (1) Å\\ b = 6.9335 (4) Å\\ c = 14.6389 (9) Å\\ V = 1884.8 (2) Å^3 \end{array}$

Data collection

Bruker APEX area-detector diffractometer φ and ω scans Absorption correction: none 16257 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.038$ $wR(F^2) = 0.120$ S = 1.052328 reflections 209 parameters H-atom parameters constrained $D_x = 1.369 \text{ Mg m}^{-3}$ Mo K α radiation $\mu = 0.10 \text{ mm}^{-1}$ T = 291 (2) KPlate, brown $0.24 \times 0.21 \times 0.05 \text{ mm}$

2328 independent reflections 1654 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.025$ $\theta_{\text{max}} = 27.5^{\circ}$

 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0605P)^{2} + 0.176P]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3$ $(\Delta/\sigma)_{max} = 0.001$ $\Delta\rho_{max} = 0.20 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{min} = -0.13 \text{ e} \text{ Å}^{-3}$ Extinction correction: *SHELXL97* (Sheldrick, 1997) Extinction coefficient: 0.0030 (8)

The two nitro groups are disordered across a mirror plane, as is the phenyl ring that is near to the -NH-N= unit. These were allowed

to refine across the symmetry element. The ring was restrained to be nearly flat and the 1,2-related distance was restrained to 1.39 (1) Å, whereas the 1,4-related distance was restrained to 2.78 (1) Å. The two nitro groups were allowed to refine off the mirror plane but with distance restraints of N-O = 1.22 (1) Å and O···O = 2.11 (1) Å. Additionally, the four-atom unit C(NO₂) was restrained to be approximately flat. As the refinement led to a *y*-coordinate of 0.251 (2) for N4, the N atom of this nitro group is ordered. The refinement proceeded with this atom lying on the mirror plane. H atoms were placed in calculated positions (N-H = 0.86 Å) and C-H = 0.93 Å), and were included in the refinement in the riding-model approximation, with $U_{iso}(H) = 1.2U_{eq}(C)$.

Data collection: *SMART* (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: *SHELXTL* (Bruker, 1998); software used to prepare material for publication: *SHELXL97*.

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