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2,5-Dibenzoyl-1,4-phenylenediamine

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Key indicators

Single-crystal X-ray study T = 296 KMean $\sigma(\text{C-C}) = 0.005 \text{ Å}$ R factor = 0.054 wR factor = 0.151Data-to-parameter ratio = 12.9

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

The title compound, $C_{20}H_{16}N_2O_2$, was synthesized from the reaction of 2,5-dibenzoylterephthalamide and sodium hypochlorite solution. The asymmetric unit contains one half-molecule, the molecule being centrosymmetric. Intra- and intermolecular $N-H\cdots O$ hydrogen bonds are highly effective in forming a two-dimensional layer structure.

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Comment

2,5-Dibenzoyl-1,4-phenylenediamine, (I), is a significant material in the synthesis of extended lattice compounds with a centrosymmetric system. It is also an important compound in preparation of electron-transport materials (Tonzola *et al.*, 2003). The synthesis of 2,5-dibenzoyl-1,4-phenylenediamine has been reported (Imai *et al.*, 1975).

$$\begin{array}{c|c}
O & NH_2 \\
C & C \\
H_2N & O
\end{array}$$

The molecular structure of (I) is shown in Fig. 1 and selected bond lengths and angles are given in Table 1. The asymmetric unit contains one half molecule, the whole molecule being centrosymmetric.

The crystal packing is stabilized by intra- and intermolecular $N-H\cdots O$ hydrogen bonds (Table 2), forming a two-dimensional layer structure (Fig. 2).

Experimental

Sodium hypochlorite solution (10 ml, 5.25%) was added with stirring to a mixture of 2,5-dibenzoylterephthalamide (1 g, 2.7 mmol) and potassium hydroxide solution (30 ml, 10.45%) cooled in an ice-water bath for half an hour. The mixture was stirred for an additional hour at 343–353 K and the precipitate began to separate. The resulting precipitate was filtered off, washed with hot water and dried under reduced pressure. The crude product was obtained by slow evaporation of a solution in benzene (yield: 0.6 g, 71%; m.p. 492 K).

Crystal data

 $C_{20}H_{16}N_2O_2$ $M_r = 316.35$ Orthorhombic, Pcab a = 7.4651 (15) Å b = 13.0034 (16) Å c = 15.9759 (18) Å V = 1550.8 (4) Å³ Z = 4 $D_x = 1.355$ Mg m⁻³ Mo $K\alpha$ radiation Cell parameters from 25 reflections $\theta = 9-12^{\circ}$ $\mu = 0.09 \text{ mm}^{-1}$ T = 296 (2) K Prism, brown $0.3 \times 0.3 \times 0.1 \text{ mm}$

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Data collection

Enraf–Nonius CAD-4 diffractometer $\omega/2\theta$ scans Absorption correction: ψ scan (XPREP in SHELXTL; Bruker, 2000) $T_{\min} = 0.974, T_{\max} = 0.991$ 1519 measured reflections 1519 independent reflections

614 reflections with $I > 2\sigma(I)$ $\theta_{\rm max} = 26.0^{\circ}$ $h = 0 \rightarrow 9$ $k = 0 \rightarrow 16$ $l = 0 \rightarrow 19$ 3 standard reflections every 200 reflections intensity decay: none

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.054$ $wR(F^2) = 0.151$ S = 1.021519 reflections 118 parameters H atoms treated by a mixture of independent and constrained refinement

$$\begin{split} w &= 1/[\sigma^2(F_{\rm o}^2) + (0.0432P)^2] \\ \text{where } P &= (F_{\rm o}^2 + 2F_{\rm c}^2)/3 \\ (\Delta/\sigma)_{\rm max} &= 0.001 \\ \Delta\rho_{\rm max} &= 0.19 \ {\rm e} \ {\rm \mathring{A}}^{-3} \\ \Delta\rho_{\rm min} &= -0.18 \ {\rm e} \ {\rm \mathring{A}}^{-3} \\ \text{Extinction correction: } SHELXL97 \\ \text{Extinction coefficient: } 0.0047 \ (19) \end{split}$$

Table 1 Selected geometric parameters (Å, °).

O-C4	1.231 (4)	N-C1	1.393 (4)
C10-C4	1.478 (5)	C3-C4	1.491 (4)
C2-C1-N	120.1 (3)	C1 ⁱ -C3-C4	120.7 (3)
C1-C2-C3	122.4 (3)	C10-C4-C3	120.9 (3)

Symmetry code: (i) -x, -y, -z.

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

$D-H\cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	D $ H$ $\cdot \cdot \cdot A$
$N-H1\cdots O^{i}$ $N-H1\cdots O^{ii}$	0.87 (4)	2.25 (4)	2.855 (5)	127 (3)
	0.87 (4)	2.61 (4)	3.220 (5)	128 (3)

Symmetry codes: (i) -x, -y, -z; (ii) $\frac{1}{2} - x, -\frac{1}{2} + y, -z$.

Atoms H1 and H3 were located in a difference synthesis and refined freely [N-H = 0.87 (4)-0.95 (5) Å]. The remaining H atoms were positioned geometrically (C-H = 0.93 Å) and refined as riding, with $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm parent\ atom})$.

Data collection: *CAD-4 Software* (Enraf–Nonius,1989); cell refinement: *CAD-4 Software*; data reduction: *XCAD4* (Harms, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 2001); software used to prepare material for publication: *SHELXTL*.

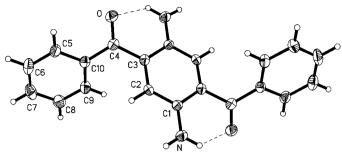


Figure 1

A drawing of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. Dashed lines represent hydrogen bonds. [Symmetry code for unlabelled atoms: -x, -y, -z.]

Figure 2

The two-dimensional layer structure of (I). Dashed lines indicate hydrogen bonds.

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