

Figure 1
ORTEP (Johnson, 1976) drawing of (3), with displacement ellipsoids plotted at the 50% probability level.

1.5056 (19) and C=O 1.2212 (15) Å. Both phenyl rings are essentially planar, as expected, and their mean planes are inclined at right angles, 89.41 (5)°, to each other. The mean plane of the nitro group is oriented at 21.40 (10)° to the plane of the aromatic ring (C1–C6) to which it is attached. The atoms lying in between the two aromatic rings, *i.e.* atoms O1/N2/C1/C7/C8 are almost coplanar, with a maximum deviation of 0.103 (2) Å for C7, and the mean planes of the aromatic rings C1–C6 and C8–C13 form dihedral angles of 58.44 (6) and 31.94 (6)°, respectively, with the mean plane of these atoms.

The structure is stabilized by a strong hydrogen bond between the amido H and carbonyl O atoms [H2···O1 1.99, N2···O1 2.8364 (13) Å and N2–H2···O1 167°], thus linking the molecules into chains along the *b* axis (Fig. 2). A search of the Cambridge Structural Database (Allen & Kennard, 1993) for similar structures revealed a dozen or so phenylbenzamide derivatives closely related to the structure of (3).

Experimental

2-Nitrobenzoic acid, (1) (10 g), was added to benzene (10 ml) and thionyl chloride (10 ml). The mixture was refluxed on a steam bath for 1 h, after which more thionyl chloride (10 ml) was added. The mixture was further refluxed for 2 h. The benzene was removed under reduced pressure. The resulting oil was treated with benzene (10 ml) in order to remove excess thionyl chloride. Benzene was removed

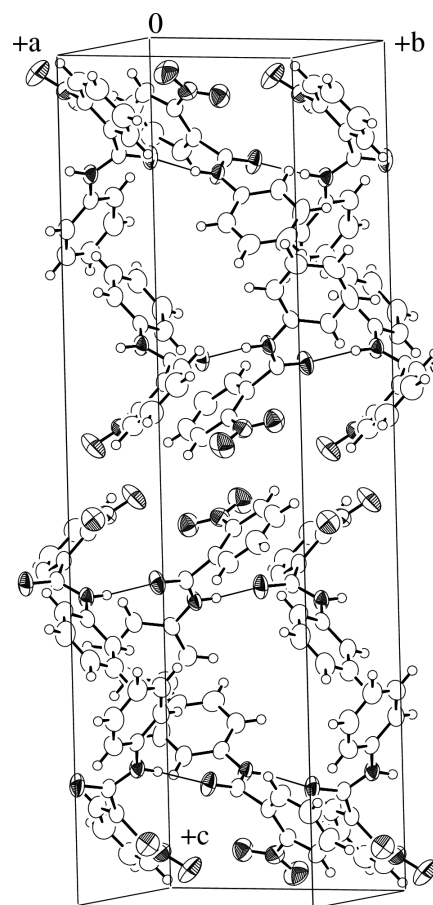


Figure 2
Hydrogen-bonding pattern in (3), showing a hydrogen-bonded polymeric chain along the *b* axis.

under reduced pressure and the residue was cooled, leaving a dark brown oil, 2-nitrobenzoyl chloride, (2) (10.5 g). Aniline (10 ml) was added dropwise to (2) and the reaction mixture was left at room temperature for 1 h. Cold water (150 ml) was added and the mixture allowed to stand for half an hour. The solid was filtered, washed with water, dried and crystallized from ethanol to give *N*-phenyl-2-nitrobenzamide, (3) (12.04 g, 87.69%), in the form of colourless prisms suitable for X-ray diffraction analysis.

Crystal data

C₁₃H₁₀N₂O₃
M_r = 242.23
 Orthorhombic, *Pbca*
a = 7.9406 (1) Å
b = 9.4695 (2) Å
c = 31.1671 (5) Å
V = 2343.56 (7) Å³
Z = 8
D_x = 1.373 Mg m⁻³

Mo *K*α radiation
 Cell parameters from 2949 reflections
 θ = 1.0–27.5°
 μ = 0.10 mm⁻¹
T = 293 (2) K
 Prismatic, colorless
 0.32 × 0.17 × 0.15 mm

Data collection

Nonius KappaCCD diffractometer
 ω and φ scans
 Absorption correction: multi-scan
 (SORTAV; Blessing, 1995, 1997)
 T_{\min} = 0.97, T_{\max} = 0.99
 4857 measured reflections
 2639 independent reflections

1945 reflections with $I > 2\sigma(I)$
 R_{int} = 0.022
 θ_{max} = 27.5°
 h = -10 → 10
 k = -12 → 12
 l = -40 → 40

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.065P)^2 + 0.31P]$
$R[F^2 > 2\sigma(F^2)] = 0.042$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.128$	$(\Delta/\sigma)_{\max} < 0.001$
$S = 1.06$	$\Delta\rho_{\max} = 0.14 \text{ e } \text{\AA}^{-3}$
2639 reflections	$\Delta\rho_{\min} = -0.16 \text{ e } \text{\AA}^{-3}$
164 parameters	Extinction correction: <i>SHELXL</i>
H-atom parameters constrained	Extinction coefficient: 0.035 (6)

Table 1
Selected geometric parameters (\AA , $^\circ$).

O1–C7	1.2212 (15)	N1–C6	1.470 (2)
O2–N1	1.2124 (16)	N2–C7	1.3436 (17)
O3–N1	1.2205 (17)	N2–C8	1.4142 (17)
O2–N1–O3	123.65 (14)	O3–N1–C6	118.05 (12)
O2–N1–C6	118.28 (13)	C7–N2–C8	126.45 (11)

Table 2
Hydrogen-bonding geometry (\AA , $^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N2–H2 \cdots O1 ⁱ	0.86	1.99	2.8364 (13)	167

Symmetry code: (i) $\frac{3}{2} - x, y - \frac{1}{2}, z$.

The H atoms were located from difference Fourier syntheses and were included in the refinement at geometrically idealized positions with C–H = 0.93 \AA and N–H = 0.86 \AA , utilizing a riding model.

Data collection: *COLLECT* (Hooft, 1998); cell refinement: *HKL DENZO* (Otwinowski & Minor, 1997); data reduction: *SCALEPACK* (Otwinowski & Minor, 1997); program(s) used to solve structure: *SAPI91* (Fan, 1991); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *TEXSAN* (Molecular Structure Corporation, 1994); software used to prepare material for publication: *SHELXL97* (Sheldrick, 1997).

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