

**N-(2-Methoxyphenyl)-2-nitrobenzamide****Aamer Saeed,<sup>a,\*</sup> Shahid Hussain<sup>a</sup> and Michael Bolte<sup>b</sup>**<sup>a</sup>Department of Chemistry, Quaid-i-Azam University, Islamabad, Pakistan, and<sup>b</sup>Institut für Anorganische Chemie, J. W. Goethe-Universität Frankfurt, Max-von-Laue-Strasse 7, 60438 Frankfurt/Main, Germany

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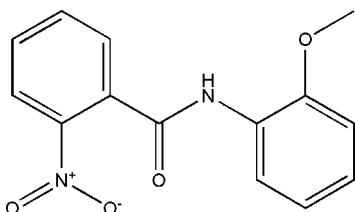
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Key indicators: single-crystal X-ray study;  $T = 173\text{ K}$ ; mean  $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$ ;  $R$  factor = 0.045;  $wR$  factor = 0.111; data-to-parameter ratio = 7.3.

Geometric parameters of the title compound,  $\text{C}_{14}\text{H}_{12}\text{N}_2\text{O}_4$ , are in the usual ranges. The dihedral angle between the two aromatic rings is  $28.9(1)^\circ$ . The nitro group is twisted by  $40.2(1)^\circ$  out of the plane of the aromatic ring to which it is attached. The crystal structure is stabilized by an  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bond.

**Related literature**

For related literature, see: Capdeville *et al.* (2002); Ho *et al.* (2002); Igawa *et al.* (1999); Jackson *et al.* (1994); Makino *et al.* (2001, 2003); Manley *et al.* (2002); Zhichkin *et al.* (2007).

**Experimental***Crystal data* $M_r = 272.26$ Orthorhombic,  $P2_12_12_1$  $a = 7.6467(11)\text{ \AA}$  $b = 9.9272(8)\text{ \AA}$  $c = 16.5032(14)\text{ \AA}$  $V = 1252.8(2)\text{ \AA}^3$  $Z = 4$ Mo  $K\alpha$  radiation $\mu = 0.11\text{ mm}^{-1}$  $T = 173(2)\text{ K}$  $0.37 \times 0.33 \times 0.21\text{ mm}$ **Data collection**

Stoe IPDSII two-circle diffractometer  
Absorption correction: none  
8342 measured reflections

1368 independent reflections  
1275 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.042$

**Refinement**

$R[F^2 > 2\sigma(F^2)] = 0.044$   
 $wR(F^2) = 0.110$   
 $S = 1.05$   
1368 reflections  
187 parameters

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\text{max}} = 0.28\text{ e \AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.21\text{ e \AA}^{-3}$

**Table 1**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}-\text{H1}\cdots\text{O2}^{\text{i}}$	0.82 (4)	2.57 (4)	3.352 (3)	159 (3)

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

Data collection: *X-AREA* (Stoe & Cie, 2001); cell refinement: *X-AREA*; data reduction: *X-AREA*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: EZ2111).

**References**

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# supporting information

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## N-(2-Methoxyphenyl)-2-nitrobenzamide

Aamer Saeed, Shahid Hussain and Michael Bolte

### S1. Comment

The benzamilide core is present in compounds with such a wide range of biological activities that it has been called a privileged structure. Benzamilides serve as intermediates to benzothiadiazin-4-ones (Makino *et al.*, 2003), quinazoline-2,4-diones (Makino *et al.*, 2001), benzodiazepine-2,5-diones (Ho *et al.*, 2002) and 2,3-disubstituted 3*H*-quinazoline-4-ones (Zhichkin *et al.*, 2007). Benzamilides have established their efficacy as central elements of ligands that bind to a wide variety of receptor types. Thus a benzamilide containing aminoalkyl groups originally designed as a peptidomimetic has been incorporated in an Arg-Gly-Asp cyclic peptide yielding a high affinity GPIIb/IIIa ligand (Jackson *et al.*, 1994). Imatinib is an ATP-site binding kinase inhibitor and platelet-derived growth factor receptor kinases (Capdeville *et al.*, 2002). Pyridylmethyl containing benzamilides are vascular endothelial growth factor receptors and tyrosine kinase inhibitors (Manley *et al.*, 2002). Furthermore, benzamides have been reported to have activities as acetyl-CoA carboxylase and farnesyl transferase inhibitors (Igawa *et al.*, 1999).

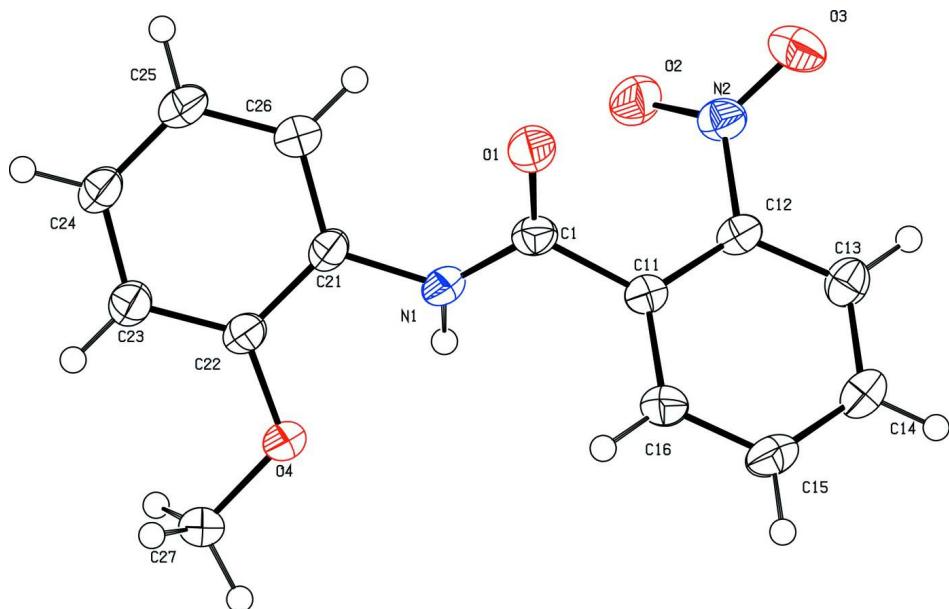
Geometric parameters of the title compound, C<sub>14</sub>H<sub>12</sub>N<sub>2</sub>O<sub>4</sub>, are in the usual ranges. The dihedral angle between the two aromatic rings is 28.9 (1) $^{\circ}$ . The nitrogroup is twisted by 40.2 (1) $^{\circ}$  out of the plane of the phenyl ring to which it is attached. The crystal structure is stabilized by an N—H···O hydrogen bond.

### S2. Experimental

A mixture of 2-methoxyaniline (10.0 g, 65.7 mmol), 2-nitrobenzoyl chloride (10 ml, 86.9 mmol), and pyridine (20 ml) was left at 25 °C for 15 h. Water (100 ml) was then added, and the resulting precipitate was collected. Recrystallization of the precipitate from benzene gave 12.6 g (75%) of the title compound as yellow blocks: mp 95–96 °C 1H NMR (CDCl<sub>3</sub>)  $\delta$  7.23–8.30 (m, 8H, Ar—Hs), 11.36 (br s, 1H, NH).

### S3. Refinement

In the absence of anomalous scatterers, Friedel pairs had been merged and the absolute structure was arbitrarily assigned. All H atoms were found in a difference map, but those bonded to C were geometrically positioned and refined with fixed individual displacement parameters [U(H) = 1.2 U<sub>eq</sub>(C) or U(H) = 1.5 U<sub>eq</sub>(C<sub>methyl</sub>)] using a riding model with C—H = 0.95 Å or C—H = 0.98 Å for aromatic and methyl C, respectively. The methyl group was allowed to rotate, but not to tip. The amino H atom was freely refined.

**Figure 1**

Molecular structure of the title compound.

### *N*-(2-Methoxyphenyl)-2-nitrobenzamide

#### Crystal data

$C_{14}H_{12}N_2O_4$   
 $M_r = 272.26$   
Orthorhombic,  $P2_12_12_1$   
 $a = 7.6467(11)$  Å  
 $b = 9.9272(8)$  Å  
 $c = 16.5032(14)$  Å  
 $V = 1252.8(2)$  Å<sup>3</sup>  
 $Z = 4$   
 $F(000) = 568$

#### Data collection

Stoe IPDSII two-circle diffractometer  
Radiation source: fine-focus sealed tube  
Graphite monochromator  
 $\omega$  scans  
8342 measured reflections  
1368 independent reflections

#### Refinement

Refinement on  $F^2$   
Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.044$   
 $wR(F^2) = 0.110$   
 $S = 1.05$   
1368 reflections  
187 parameters  
0 restraints

$D_x = 1.444$  Mg m<sup>-3</sup>  
Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å  
Cell parameters from 7856 reflections  
 $\theta = 3.8\text{--}25.6^\circ$   
 $\mu = 0.11$  mm<sup>-1</sup>  
 $T = 173$  K  
Block, light yellow  
0.37 × 0.33 × 0.21 mm

1275 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.042$   
 $\theta_{\text{max}} = 25.6^\circ$ ,  $\theta_{\text{min}} = 3.6^\circ$   
 $h = -7 \rightarrow 9$   
 $k = -12 \rightarrow 12$   
 $l = -20 \rightarrow 18$

Primary atom site location: structure-invariant direct methods  
Secondary atom site location: difference Fourier map  
Hydrogen site location: inferred from neighbouring sites  
H atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0824P)^2]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} < 0.001$$

$$\Delta\rho_{\max} = 0.28 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$$

Extinction correction: *SHELXL97* (Sheldrick, 2008),  $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.052 (7)

### Special details

**Geometry.** All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

**Refinement.** Refinement of  $F^2$  against ALL reflections. The weighted  $R$ -factor  $wR$  and goodness of fit  $S$  are based on  $F^2$ , conventional  $R$ -factors  $R$  are based on  $F$ , with  $F$  set to zero for negative  $F^2$ . The threshold expression of  $F^2 > \sigma(F^2)$  is used only for calculating  $R$ -factors(gt) etc. and is not relevant to the choice of reflections for refinement.  $R$ -factors based on  $F^2$  are statistically about twice as large as those based on  $F$ , and  $R$ -factors based on ALL data will be even larger.

### Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.7040 (3)	0.4189 (2)	0.58905 (15)	0.0271 (5)
N1	0.6935 (3)	0.46807 (19)	0.51339 (14)	0.0287 (5)
H1	0.618 (5)	0.449 (3)	0.4799 (19)	0.040 (9)*
N2	0.7600 (3)	0.14453 (18)	0.64422 (13)	0.0299 (5)
O1	0.8235 (3)	0.44489 (17)	0.63779 (11)	0.0370 (5)
O2	0.8450 (3)	0.16188 (18)	0.58190 (13)	0.0410 (5)
O3	0.8114 (3)	0.07915 (18)	0.70333 (13)	0.0419 (5)
O4	0.5752 (2)	0.62521 (16)	0.39737 (11)	0.0305 (5)
C11	0.5537 (3)	0.3307 (2)	0.61508 (14)	0.0263 (5)
C12	0.5833 (3)	0.2030 (2)	0.64870 (16)	0.0264 (5)
C13	0.4498 (4)	0.1261 (2)	0.68256 (15)	0.0307 (6)
H13	0.4744	0.0404	0.7055	0.037*
C14	0.2795 (4)	0.1764 (2)	0.68236 (17)	0.0338 (6)
H14	0.1869	0.1248	0.7049	0.041*
C15	0.2456 (4)	0.3028 (3)	0.64891 (17)	0.0360 (6)
H15	0.1294	0.3368	0.6483	0.043*
C16	0.3821 (4)	0.3800 (2)	0.61615 (17)	0.0324 (6)
H16	0.3579	0.4667	0.5945	0.039*
C21	0.8150 (3)	0.5560 (2)	0.47494 (14)	0.0260 (5)
C22	0.7505 (3)	0.6385 (2)	0.41263 (15)	0.0267 (5)
C23	0.8635 (4)	0.7254 (2)	0.37131 (16)	0.0293 (6)
H23	0.8199	0.7825	0.3297	0.035*
C24	1.0410 (4)	0.7277 (2)	0.39163 (16)	0.0314 (6)
H24	1.1177	0.7872	0.3638	0.038*
C25	1.1068 (4)	0.6440 (2)	0.45211 (17)	0.0323 (6)
H25	1.2280	0.6452	0.4649	0.039*
C26	0.9933 (4)	0.5579 (2)	0.49399 (15)	0.0299 (5)
H26	1.0374	0.5007	0.5354	0.036*
C27	0.5074 (4)	0.6982 (2)	0.32917 (16)	0.0323 (6)
H27A	0.5211	0.7951	0.3385	0.048*
H27B	0.3831	0.6769	0.3222	0.048*

H27C	0.5717	0.6724	0.2802	0.048*
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*Atomic displacement parameters ( $\text{\AA}^2$ )*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C1	0.0227 (13)	0.0256 (10)	0.0331 (13)	0.0022 (9)	-0.0009 (10)	-0.0013 (9)
N1	0.0195 (11)	0.0312 (10)	0.0354 (12)	-0.0040 (8)	-0.0015 (10)	0.0024 (8)
N2	0.0261 (12)	0.0246 (9)	0.0390 (12)	0.0014 (9)	-0.0020 (9)	-0.0034 (8)
O1	0.0324 (11)	0.0415 (9)	0.0372 (10)	-0.0095 (9)	-0.0085 (8)	0.0057 (8)
O2	0.0308 (11)	0.0429 (9)	0.0491 (11)	0.0067 (8)	0.0110 (9)	0.0029 (9)
O3	0.0413 (12)	0.0373 (8)	0.0470 (12)	0.0104 (9)	-0.0093 (9)	0.0037 (8)
O4	0.0209 (9)	0.0330 (8)	0.0374 (10)	-0.0019 (7)	-0.0012 (8)	0.0062 (7)
C11	0.0217 (12)	0.0282 (11)	0.0289 (12)	0.0003 (10)	-0.0005 (11)	0.0009 (9)
C12	0.0208 (14)	0.0269 (10)	0.0315 (12)	-0.0008 (9)	0.0007 (10)	-0.0038 (9)
C13	0.0312 (15)	0.0271 (10)	0.0338 (13)	-0.0040 (10)	0.0025 (12)	-0.0029 (9)
C14	0.0268 (15)	0.0368 (12)	0.0377 (14)	-0.0069 (11)	0.0047 (12)	-0.0027 (10)
C15	0.0216 (14)	0.0457 (13)	0.0408 (15)	0.0011 (11)	0.0024 (12)	-0.0023 (11)
C16	0.0255 (14)	0.0355 (12)	0.0362 (13)	0.0055 (10)	0.0001 (12)	0.0043 (10)
C21	0.0216 (12)	0.0247 (10)	0.0315 (12)	-0.0026 (10)	0.0041 (10)	0.0000 (9)
C22	0.0210 (12)	0.0261 (10)	0.0328 (12)	-0.0010 (9)	0.0016 (10)	-0.0011 (9)
C23	0.0270 (14)	0.0264 (10)	0.0345 (13)	-0.0007 (10)	0.0036 (11)	0.0005 (9)
C24	0.0247 (14)	0.0304 (11)	0.0390 (14)	-0.0047 (10)	0.0044 (12)	-0.0004 (10)
C25	0.0225 (13)	0.0345 (12)	0.0398 (13)	-0.0043 (11)	-0.0005 (11)	-0.0034 (10)
C26	0.0243 (13)	0.0328 (12)	0.0326 (12)	0.0005 (11)	-0.0010 (10)	-0.0008 (9)
C27	0.0240 (14)	0.0362 (12)	0.0367 (14)	0.0010 (10)	-0.0036 (11)	0.0074 (9)

*Geometric parameters ( $\text{\AA}$ ,  $^\circ$ )*

C1—O1	1.244 (3)	C15—C16	1.404 (4)
C1—N1	1.343 (3)	C15—H15	0.9500
C1—C11	1.508 (3)	C16—H16	0.9500
N1—C21	1.424 (3)	C21—C26	1.399 (4)
N1—H1	0.82 (4)	C21—C22	1.404 (3)
N2—O2	1.229 (3)	C22—C23	1.399 (3)
N2—O3	1.236 (3)	C23—C24	1.399 (4)
N2—C12	1.473 (3)	C23—H23	0.9500
O4—C22	1.371 (3)	C24—C25	1.393 (4)
O4—C27	1.435 (3)	C24—H24	0.9500
C11—C16	1.400 (4)	C25—C26	1.401 (4)
C11—C12	1.402 (3)	C25—H25	0.9500
C12—C13	1.392 (4)	C26—H26	0.9500
C13—C14	1.395 (4)	C27—H27A	0.9800
C13—H13	0.9500	C27—H27B	0.9800
C14—C15	1.395 (4)	C27—H27C	0.9800
C14—H14	0.9500		
O1—C1—N1	124.7 (2)	C11—C16—H16	119.6
O1—C1—C11	119.7 (2)	C15—C16—H16	119.6

N1—C1—C11	115.5 (2)	C26—C21—C22	119.9 (2)
C1—N1—C21	126.7 (2)	C26—C21—N1	122.9 (2)
C1—N1—H1	126 (2)	C22—C21—N1	117.1 (2)
C21—N1—H1	107 (2)	O4—C22—C23	125.0 (2)
O2—N2—O3	124.5 (2)	O4—C22—C21	115.0 (2)
O2—N2—C12	118.2 (2)	C23—C22—C21	120.0 (2)
O3—N2—C12	117.3 (2)	C22—C23—C24	119.5 (2)
C22—O4—C27	116.69 (19)	C22—C23—H23	120.2
C16—C11—C12	117.5 (2)	C24—C23—H23	120.2
C16—C11—C1	121.0 (2)	C25—C24—C23	120.8 (2)
C12—C11—C1	121.0 (2)	C25—C24—H24	119.6
C13—C12—C11	122.5 (2)	C23—C24—H24	119.6
C13—C12—N2	118.5 (2)	C24—C25—C26	119.6 (3)
C11—C12—N2	119.0 (2)	C24—C25—H25	120.2
C12—C13—C14	119.1 (2)	C26—C25—H25	120.2
C12—C13—H13	120.4	C21—C26—C25	120.1 (2)
C14—C13—H13	120.4	C21—C26—H26	120.0
C13—C14—C15	119.8 (2)	C25—C26—H26	120.0
C13—C14—H14	120.1	O4—C27—H27A	109.5
C15—C14—H14	120.1	O4—C27—H27B	109.5
C14—C15—C16	120.3 (3)	H27A—C27—H27B	109.5
C14—C15—H15	119.8	O4—C27—H27C	109.5
C16—C15—H15	119.8	H27A—C27—H27C	109.5
C11—C16—C15	120.7 (2)	H27B—C27—H27C	109.5
O1—C1—N1—C21	0.8 (4)	C12—C11—C16—C15	-0.8 (4)
C11—C1—N1—C21	178.3 (2)	C1—C11—C16—C15	-172.8 (2)
O1—C1—C11—C16	118.2 (3)	C14—C15—C16—C11	1.2 (4)
N1—C1—C11—C16	-59.5 (3)	C1—N1—C21—C26	29.1 (4)
O1—C1—C11—C12	-53.6 (3)	C1—N1—C21—C22	-154.1 (2)
N1—C1—C11—C12	128.8 (2)	C27—O4—C22—C23	5.6 (3)
C16—C11—C12—C13	-0.2 (4)	C27—O4—C22—C21	-174.3 (2)
C1—C11—C12—C13	171.8 (2)	C26—C21—C22—O4	177.9 (2)
C16—C11—C12—N2	176.4 (2)	N1—C21—C22—O4	1.0 (3)
C1—C11—C12—N2	-11.6 (3)	C26—C21—C22—C23	-2.0 (3)
O2—N2—C12—C13	138.0 (3)	N1—C21—C22—C23	-178.8 (2)
O3—N2—C12—C13	-40.8 (3)	O4—C22—C23—C24	-178.8 (2)
O2—N2—C12—C11	-38.8 (3)	C21—C22—C23—C24	1.1 (4)
O3—N2—C12—C11	142.4 (2)	C22—C23—C24—C25	0.4 (4)
C11—C12—C13—C14	0.8 (4)	C23—C24—C25—C26	-1.0 (4)
N2—C12—C13—C14	-175.8 (2)	C22—C21—C26—C25	1.3 (4)
C12—C13—C14—C15	-0.4 (4)	N1—C21—C26—C25	178.0 (2)
C13—C14—C15—C16	-0.6 (4)	C24—C25—C26—C21	0.1 (4)

Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ )

D—H···A	D—H	H···A	D···A	D—H···A
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## supporting information

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N1—H1···O2 <sup>i</sup>	0.82 (4)	2.57 (4)	3.352 (3)	159 (3)
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Symmetry code: (i)  $x-1/2, -y+1/2, -z+1$ .