

2-Nitro-N-(8-quinolyl)benzamide**Gang Lei,* Lin-Hai Jing and Li Zhou**

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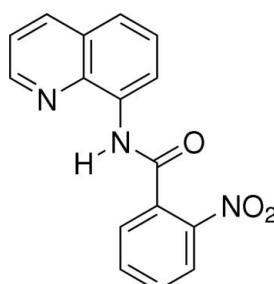
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Key indicators: single-crystal X-ray study; $T = 93\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.051; wR factor = 0.145; data-to-parameter ratio = 14.8.

In the title compound, $\text{C}_{16}\text{H}_{11}\text{N}_3\text{O}_3$, the amide group is twisted away from the quinoline ring system and nitrobenzene ring by $8.02(1)^\circ$ and $54.92(1)^\circ$, respectively. The crystal packing is stabilized by intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds, and $\pi-\pi$ interactions between the quinoline ring systems of inversion-related molecules, with a centroid–centroid distance of $3.4802(13)\text{ \AA}$.

Related literature

For the biological activities of quinoline derivatives, see: Oku *et al.* (1998, 1999).

**Experimental***Crystal data*

$\text{C}_{16}\text{H}_{11}\text{N}_3\text{O}_3$	$V = 1301.4(6)\text{ \AA}^3$
$M_r = 293.28$	$Z = 4$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 12.430(3)\text{ \AA}$	$\mu = 0.11\text{ mm}^{-1}$
$b = 10.144(3)\text{ \AA}$	$T = 93(2)\text{ K}$
$c = 11.528(3)\text{ \AA}$	$0.50 \times 0.40 \times 0.15\text{ mm}$
$\beta = 116.449(3)^\circ$	

Data collection

Rigaku R-AXIS RAPID diffractometer
Absorption correction: none
10410 measured reflections

2949 independent reflections
2679 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.029$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.051$
 $wR(F^2) = 0.145$
 $S = 1.00$
2949 reflections

199 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.22\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.28\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C3—H3 \cdots O3 ⁱ	0.95	2.55	3.209 (2)	127
C4—H4 \cdots O2 ⁱⁱ	0.95	2.48	3.319 (2)	147
C17—H17 \cdots O1 ⁱⁱⁱ	0.95	2.42	3.160 (2)	135

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

Data collection: *RAPID-AUTO* (Rigaku, 2004); cell refinement: *RAPID-AUTO*; data reduction: *RAPID-AUTO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *XP* in *SHELXTL* (Sheldrick, 2008); 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: CI2716).

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- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.

supporting information

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2-Nitro-*N*-(8-quinolyl)benzamide

Gang Lei, Lin-Hai Jing and Li Zhou

S1. Comment

Quinoline derivatives are a class of important compound for the treatment of bone metabolic disorders (Oku *et al.*, 1998) and as H⁺-ATPases inhibitors (Oku *et al.*, 1999). We report here the crystal structure of the title compound.

Bond lengths and angles in title molecule (Fig. 1) are normal. The quinoline ring system is planar, with a maximum deviation of 0.033 (2) Å for atom C3. As a result of steric effects, the amide group is twisted away from the planes of the quinoline ring system and the nitrobenzene ring. The C5-C10 and C12-C17 planes form dihedral angles of 8.02 (1) and 54.92 (1)°, respectively, with the O1/N2/C8/C11 plane. The dihedral angle between the C12-C17 and O2/O3/N3/C13 planes is 36.83 (1)°.

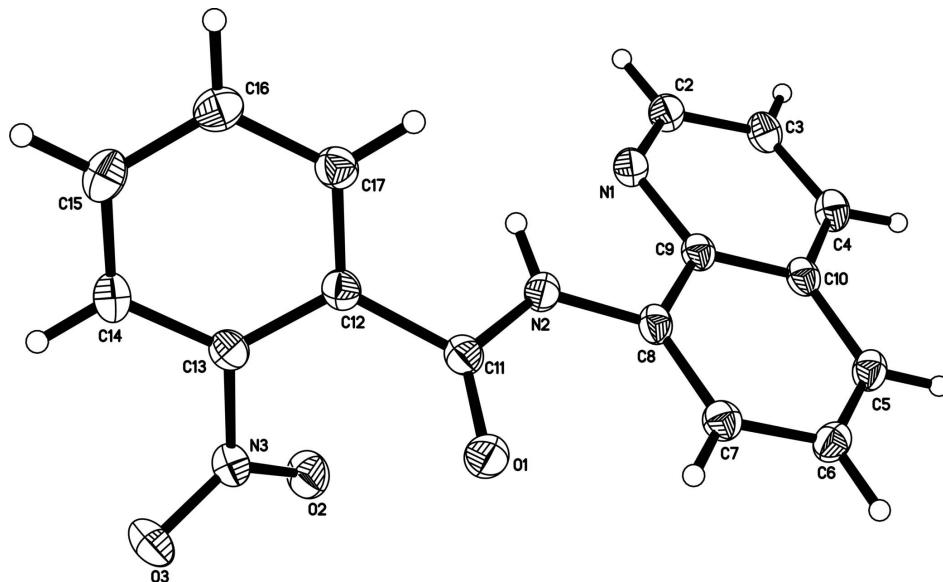
The crystal packing is stabilized by C—H···O hydrogen bonds (Table 1), and π-π interactions between the benzene rings of the inversion-related molecules at (x, y, z) and (-x, 1 - y, 1 - z), with a centroid-centroid distance of 3.4802 (13) Å.

S2. Experimental

O-Nitrobenzoic acid (2 mmol) and an excess of thionyl chloride (3 mmol) in dioxane (20 ml) were boiled under reflux for 6 h. The solution was distilled under reduced pressure and a yellow solid was obtained. 8-Aminoquinoline (2 mmol) in tetrahydrofuran (20 ml) was added to the yellow solid and boiled under reflux for 6 h. The solution was then cooled to ambient temperature and filtered to remove the tetrahydrofuran. The precipitate was dissolved in the dimethyl sulfoxide and allowed to stand for one month at ambient temperature, after which time white single crystals of the title compound suitable for X-ray diffraction were obtained.

S3. Refinement

All H atoms were placed in calculated positions, with C-H = 0.95 Å and N-H = 0.88 Å, and refined using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$.

**Figure 1**

The molecular structure of the title compound, showing 50% probability displacement ellipsoids and the atomic numbering.

2-Nitro-N-(8-quinolyl)benzamide

Crystal data

$C_{16}H_{11}N_3O_3$
 $M_r = 293.28$
Monoclinic, $P2_1/c$
Hall symbol: -P 2ybc
 $a = 12.430 (3)$ Å
 $b = 10.144 (3)$ Å
 $c = 11.528 (3)$ Å
 $\beta = 116.449 (3)^\circ$
 $V = 1301.4 (6)$ Å³
 $Z = 4$

$F(000) = 608$
 $D_x = 1.497 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 3984 reflections
 $\theta = 3.2\text{--}27.5^\circ$
 $\mu = 0.11 \text{ mm}^{-1}$
 $T = 93$ K
Block, white
 $0.50 \times 0.40 \times 0.15$ mm

Data collection

Rigaku R-AXIS RAPID
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
10410 measured reflections
2949 independent reflections

2679 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.029$
 $\theta_{\text{max}} = 27.5^\circ, \theta_{\text{min}} = 3.5^\circ$
 $h = -16 \rightarrow 16$
 $k = -12 \rightarrow 13$
 $l = -14 \rightarrow 11$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.051$
 $wR(F^2) = 0.145$
 $S = 1.00$
2949 reflections
199 parameters

0 restraints
Primary atom site location: structure-invariant
direct methods
Secondary atom site location: difference Fourier
map
Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.089P)^2 + 0.36P]$
 where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.005$
 $\Delta\rho_{\max} = 0.22 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.28 \text{ e } \text{\AA}^{-3}$

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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.33297 (10)	0.27884 (13)	0.71053 (11)	0.0283 (3)
O2	0.42529 (10)	0.55360 (12)	0.75700 (12)	0.0290 (3)
O3	0.60516 (11)	0.50766 (12)	0.90418 (11)	0.0291 (3)
N1	0.07416 (11)	0.46239 (13)	0.28513 (13)	0.0202 (3)
N2	0.23340 (11)	0.35402 (13)	0.50283 (13)	0.0201 (3)
H2N	0.2460	0.3835	0.4381	0.024*
N3	0.52329 (11)	0.49852 (13)	0.79419 (13)	0.0213 (3)
C2	-0.00265 (14)	0.52243 (15)	0.17885 (15)	0.0212 (3)
H2	0.0284	0.5671	0.1278	0.025*
C3	-0.12782 (14)	0.52428 (15)	0.13644 (16)	0.0223 (4)
H3	-0.1791	0.5691	0.0590	0.027*
C4	-0.17449 (14)	0.46062 (15)	0.20824 (16)	0.0221 (4)
H4	-0.2588	0.4591	0.1803	0.027*
C5	-0.13518 (14)	0.33419 (15)	0.40878 (16)	0.0210 (3)
H5	-0.2184	0.3298	0.3865	0.025*
C6	-0.05278 (14)	0.27986 (15)	0.52209 (16)	0.0215 (3)
H6	-0.0800	0.2388	0.5783	0.026*
C7	0.07210 (14)	0.28306 (15)	0.55819 (16)	0.0205 (3)
H7	0.1275	0.2441	0.6373	0.025*
C8	0.11263 (13)	0.34276 (14)	0.47822 (15)	0.0187 (3)
C9	0.02839 (13)	0.40129 (14)	0.35897 (15)	0.0181 (3)
C10	-0.09602 (13)	0.39701 (15)	0.32465 (15)	0.0194 (3)
C11	0.33334 (13)	0.32539 (15)	0.61325 (15)	0.0197 (3)
C12	0.44935 (13)	0.34895 (15)	0.60488 (14)	0.0184 (3)
C13	0.54399 (13)	0.41992 (15)	0.69958 (15)	0.0189 (3)
C14	0.65666 (14)	0.42618 (15)	0.70338 (16)	0.0220 (3)
H14	0.7198	0.4741	0.7698	0.026*
C15	0.67563 (14)	0.36117 (15)	0.60838 (17)	0.0243 (4)
H15	0.7527	0.3629	0.6100	0.029*
C16	0.58228 (15)	0.29374 (16)	0.51112 (16)	0.0239 (4)
H16	0.5951	0.2518	0.4447	0.029*
C17	0.47000 (14)	0.28663 (15)	0.50942 (15)	0.0216 (3)

H17	0.4070	0.2389	0.4426	0.026*
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Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0213 (6)	0.0419 (7)	0.0199 (6)	-0.0011 (5)	0.0074 (5)	0.0064 (5)
O2	0.0197 (6)	0.0347 (7)	0.0308 (7)	0.0053 (5)	0.0098 (5)	-0.0042 (5)
O3	0.0255 (6)	0.0327 (7)	0.0207 (6)	-0.0014 (5)	0.0026 (5)	-0.0051 (5)
N1	0.0183 (6)	0.0203 (6)	0.0204 (7)	-0.0013 (5)	0.0073 (5)	0.0005 (5)
N2	0.0168 (6)	0.0242 (7)	0.0187 (7)	-0.0022 (5)	0.0074 (5)	0.0018 (5)
N3	0.0191 (6)	0.0210 (7)	0.0213 (7)	-0.0014 (5)	0.0067 (6)	-0.0016 (5)
C2	0.0218 (8)	0.0205 (7)	0.0200 (8)	-0.0014 (6)	0.0081 (6)	0.0023 (6)
C3	0.0201 (7)	0.0195 (7)	0.0229 (8)	0.0002 (6)	0.0056 (6)	0.0004 (6)
C4	0.0174 (7)	0.0197 (7)	0.0267 (9)	0.0012 (6)	0.0076 (6)	-0.0010 (6)
C5	0.0181 (7)	0.0197 (7)	0.0271 (8)	-0.0013 (6)	0.0118 (6)	-0.0031 (6)
C6	0.0244 (8)	0.0186 (7)	0.0259 (9)	-0.0027 (6)	0.0151 (7)	-0.0014 (6)
C7	0.0213 (8)	0.0188 (7)	0.0211 (8)	-0.0005 (6)	0.0092 (6)	-0.0016 (6)
C8	0.0169 (7)	0.0167 (7)	0.0216 (8)	-0.0023 (5)	0.0079 (6)	-0.0034 (6)
C9	0.0184 (7)	0.0156 (7)	0.0196 (8)	-0.0014 (5)	0.0078 (6)	-0.0024 (6)
C10	0.0184 (7)	0.0165 (7)	0.0219 (8)	-0.0010 (5)	0.0077 (6)	-0.0039 (6)
C11	0.0193 (7)	0.0199 (7)	0.0188 (8)	-0.0008 (6)	0.0076 (6)	-0.0015 (6)
C12	0.0171 (7)	0.0189 (7)	0.0165 (7)	0.0014 (5)	0.0050 (6)	0.0040 (5)
C13	0.0188 (7)	0.0181 (7)	0.0180 (8)	0.0014 (5)	0.0066 (6)	0.0013 (6)
C14	0.0165 (7)	0.0205 (7)	0.0259 (8)	-0.0006 (6)	0.0065 (6)	0.0007 (6)
C15	0.0216 (8)	0.0207 (7)	0.0328 (9)	0.0030 (6)	0.0140 (7)	0.0049 (7)
C16	0.0281 (8)	0.0219 (8)	0.0249 (9)	0.0015 (6)	0.0148 (7)	0.0021 (6)
C17	0.0225 (8)	0.0223 (8)	0.0189 (8)	-0.0012 (6)	0.0081 (6)	0.0009 (6)

Geometric parameters (\AA , $^\circ$)

O1—C11	1.219 (2)	C6—C7	1.417 (2)
O2—N3	1.2307 (17)	C6—H6	0.95
O3—N3	1.2257 (17)	C7—C8	1.373 (2)
N1—C2	1.320 (2)	C7—H7	0.95
N1—C9	1.365 (2)	C8—C9	1.434 (2)
N2—C11	1.3562 (19)	C9—C10	1.416 (2)
N2—C8	1.4028 (19)	C11—C12	1.507 (2)
N2—H2N	0.88	C12—C17	1.389 (2)
N3—C13	1.463 (2)	C12—C13	1.396 (2)
C2—C3	1.408 (2)	C13—C14	1.383 (2)
C2—H2	0.95	C14—C15	1.386 (2)
C3—C4	1.366 (2)	C14—H14	0.95
C3—H3	0.95	C15—C16	1.383 (2)
C4—C10	1.415 (2)	C15—H15	0.95
C4—H4	0.95	C16—C17	1.389 (2)
C5—C6	1.366 (2)	C16—H16	0.95
C5—C10	1.416 (2)	C17—H17	0.95
C5—H5	0.95		

C2—N1—C9	117.28 (13)	N1—C9—C10	123.11 (14)
C11—N2—C8	128.52 (13)	N1—C9—C8	117.16 (13)
C11—N2—H2N	115.7	C10—C9—C8	119.70 (14)
C8—N2—H2N	115.7	C4—C10—C5	123.56 (14)
O3—N3—O2	124.26 (14)	C4—C10—C9	117.17 (14)
O3—N3—C13	118.14 (13)	C5—C10—C9	119.23 (14)
O2—N3—C13	117.59 (13)	O1—C11—N2	124.71 (14)
N1—C2—C3	123.99 (15)	O1—C11—C12	121.09 (14)
N1—C2—H2	118.0	N2—C11—C12	114.12 (13)
C3—C2—H2	118.0	C17—C12—C13	117.75 (14)
C4—C3—C2	119.02 (15)	C17—C12—C11	119.90 (13)
C4—C3—H3	120.5	C13—C12—C11	121.93 (14)
C2—C3—H3	120.5	C14—C13—C12	122.48 (15)
C3—C4—C10	119.39 (14)	C14—C13—N3	117.59 (13)
C3—C4—H4	120.3	C12—C13—N3	119.80 (13)
C10—C4—H4	120.3	C13—C14—C15	118.66 (14)
C6—C5—C10	119.76 (14)	C13—C14—H14	120.7
C6—C5—H5	120.1	C15—C14—H14	120.7
C10—C5—H5	120.1	C16—C15—C14	119.96 (15)
C5—C6—C7	121.88 (15)	C16—C15—H15	120.0
C5—C6—H6	119.1	C14—C15—H15	120.0
C7—C6—H6	119.1	C15—C16—C17	120.81 (15)
C8—C7—C6	119.64 (15)	C15—C16—H16	119.6
C8—C7—H7	120.2	C17—C16—H16	119.6
C6—C7—H7	120.2	C12—C17—C16	120.29 (15)
C7—C8—N2	125.45 (14)	C12—C17—H17	119.9
C7—C8—C9	119.78 (14)	C16—C17—H17	119.9
N2—C8—C9	114.76 (13)		
C9—N1—C2—C3	-1.5 (2)	C8—C9—C10—C5	-0.3 (2)
N1—C2—C3—C4	0.0 (2)	C8—N2—C11—O1	3.0 (3)
C2—C3—C4—C10	1.4 (2)	C8—N2—C11—C12	179.71 (14)
C10—C5—C6—C7	-0.7 (2)	O1—C11—C12—C17	120.18 (17)
C5—C6—C7—C8	0.4 (2)	N2—C11—C12—C17	-56.67 (19)
C6—C7—C8—N2	179.31 (14)	O1—C11—C12—C13	-52.2 (2)
C6—C7—C8—C9	0.0 (2)	N2—C11—C12—C13	130.96 (15)
C11—N2—C8—C7	-9.2 (3)	C17—C12—C13—C14	-2.1 (2)
C11—N2—C8—C9	170.17 (14)	C11—C12—C13—C14	170.39 (14)
C2—N1—C9—C10	1.7 (2)	C17—C12—C13—N3	173.65 (13)
C2—N1—C9—C8	-176.46 (14)	C11—C12—C13—N3	-13.8 (2)
C7—C8—C9—N1	178.25 (13)	O3—N3—C13—C14	-37.9 (2)
N2—C8—C9—N1	-1.14 (19)	O2—N3—C13—C14	140.59 (15)
C7—C8—C9—C10	0.0 (2)	O3—N3—C13—C12	146.16 (15)
N2—C8—C9—C10	-179.40 (13)	O2—N3—C13—C12	-35.4 (2)
C3—C4—C10—C5	176.77 (14)	C12—C13—C14—C15	1.0 (2)
C3—C4—C10—C9	-1.2 (2)	N3—C13—C14—C15	-174.87 (13)
C6—C5—C10—C4	-177.25 (15)	C13—C14—C15—C16	1.1 (2)

C6—C5—C10—C9	0.6 (2)	C14—C15—C16—C17	−2.0 (2)
N1—C9—C10—C4	−0.4 (2)	C13—C12—C17—C16	1.2 (2)
C8—C9—C10—C4	177.72 (13)	C11—C12—C17—C16	−171.51 (14)
N1—C9—C10—C5	−178.44 (13)	C15—C16—C17—C12	0.9 (2)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
C3—H3···O3 ⁱ	0.95	2.55	3.209 (2)	127
C4—H4···O2 ⁱⁱ	0.95	2.48	3.319 (2)	147
C17—H17···O1 ⁱⁱⁱ	0.95	2.42	3.160 (2)	135

Symmetry codes: (i) $x-1, y, z-1$; (ii) $-x, -y+1, -z+1$; (iii) $x, -y+1/2, z-1/2$.