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# 4-Nitro-N-(8-quinolyl)benzenesulfonamide

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#### **Key indicators**

Single-crystal X-ray study T = 299 KMean  $\sigma(\text{C}-\text{C}) = 0.005 \text{ Å}$  R factor = 0.055 wR factor = 0.150 Data-to-parameter ratio = 10.1

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. In the title compound,  $C_{15}H_{11}N_3O_4S$ , the dihedral angle between the quinoline ring system and the benzene ring is 79.9 (3)°. Intermolecular hydrogen bonds of types N-H···O, C-H···O and C-H···N are observed. Received 4 October 2005 Accepted 12 October 2005 Online 22 October 2005

### Comment

Zinc(II)-specific fluorophores are of substantial importance in the study of intracellular  $Zn^{2+}$ . Many zinquin-related fluorophore esters based on an 8-*p*-toluenesulfonamidoquinoline system have recently been developed and their properties investigated (Kimber *et al.*, 2003). We report here the structure of the title compound, (I), determined as a part of our studies to investigate potential new fluorophores to be used as ligands for coordination to  $Zn^{II}$  and  $Cu^{II}$  ions. The key feature of the molecular structure of (I) (Fig. 1) is the C1-N1-S1-C10dihedral angle of 79.9 (3)°, *i.e.* that formed between the phenyl ring and the quinoline group. The crystal packing of (I) is stabilized through a hydrogen-bonding network, as shown in Fig. 2 and detailed in Table 1.



#### Experimental

Compound (I) was prepared by the overnight reaction of one equivalent of 8-aminoquinoline and a 1.1 equivalent of *p*-nitrobenzenesulfonyl chloride in the presence of pyridine, according to the literature procedure of Kimber *et al.* (2000). Single crystals of (I) suitable for X-ray data collection were obtained by recrystallization from a methanol-dichloromethane (1:1) solution of (I).

Crystal data C15H11N3O4S  $D_r = 1.538 \text{ Mg m}^{-3}$  $M_r = 329.33$ Cu Ka radiation Monoclinic, P21 Cell parameters from 25 a = 5.0135 (6) Å reflections b = 20.636 (2) Å  $\theta = 4.3 - 20.3^{\circ}$  $\mu = 2.27 \text{ mm}^{-1}$ c = 6.966 (1) Å $\beta = 99.28 \ (1)^{\circ}$ T = 299 (2) K Needle, colorless  $V = 711.26 (15) \text{ Å}^3$  $0.60 \times 0.18 \times 0.10 \ \mathrm{mm}$ Z = 2

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#### Figure 1

Molecular structure of (I), showing the atom labeling and displacement ellipsoids drawn at the 50% probability level.



#### Figure 2 Molecular packing of (I), with hydrogen bonds shown as dashed lines.

#### Data collection

Nonius CAD-4 diffractometer	$R_{\rm int} = 0.021$
$\omega/2\theta$ scans	$\theta_{\rm max} = 67.9^{\circ}$
Absorption correction: $\psi$ scan	$h = -6 \rightarrow 0$
(North et al., 1968)	$k = -24 \rightarrow 11$
$T_{\min} = 0.282, T_{\max} = 0.792$	$l = -8 \rightarrow 8$
2347 measured reflections	3 standard reflections
2109 independent reflections	frequency: 120 min
2081 reflections with $I > 2\sigma(I)$	intensity decay: 1.5%

#### Refinement

Refinement on $F^2$	$(\Delta/\sigma)_{\rm max} < 0.001$
$R[F^2 > 2\sigma(F^2)] = 0.055$	$\Delta \rho_{\rm max} = 0.64 \ {\rm e} \ {\rm \AA}^{-3}$
$wR(F^2) = 0.150$	$\Delta \rho_{\rm min} = -0.35 \text{ e } \text{\AA}^{-3}$
S = 1.14	Extinction correction: SHELXL97
2109 reflections	Extinction coefficient: 0.024 (3)
209 parameters	Absolute structure: Flack (1983),
H-atom parameters constrained	778 Friedel pairs
$w = 1/[\sigma^2(F_o^2) + (0.1304P)^2]$	Flack parameter: 0.05 (2)
where $P = (F^2 + 2F^2)/3$	

Table 1		
Hydrogen-bond geometry	(Å.	°).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N1 - H1N \cdots O2^i$	0.86	2.16	2.935 (4)	150
C11−H11···N2 <sup>ii</sup>	0.93	2.50	3.407 (5)	1
C12−H12···O1 <sup>iii</sup>	0.93	2.45	3.360 (4)	166

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

The H atoms were included in the riding-model approximation, with N-H = 0.86 Å and C-H = 0.93 Å, and with  $U_{iso}(H)$  =  $1.2U_{eq}(N,C).$ 

Data collection: CAD-4/PC Software (Nonius, 1996); cell refinement: CAD-4/PC Software; data reduction: REDU4 (Stoe & Cie, 1987); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXL97.

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