

Luiz Everson da Silva,<sup>a,b</sup>  
Antonio Carlos Joussef,<sup>a</sup> Sabine  
Foro<sup>b</sup> and Boris Schmidt<sup>b\*</sup>

<sup>a</sup>Departamento de Química—UFSC, 88040-900 Florianópolis, SC, Brazil, and <sup>b</sup>Clemens Schöpf-Institut für Organische Chemie und Biochemie, Technische Universität Darmstadt, Petersenstrasse 22, D-64287 Darmstadt, Germany

Correspondence e-mail: foro@tu-darmstadt.de

#### Key indicators

Single-crystal X-ray study  
 $T = 100$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å  
 $R$  factor = 0.039  
 $wR$  factor = 0.102  
Data-to-parameter ratio = 16.0

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

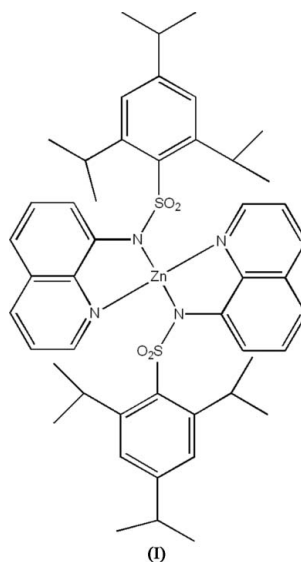
## Bis[2,4,6-triisopropyl-*N*-(quinolin-8-yl)benzene-sulfonamidato- $\kappa^2\text{N},\text{N}'$ ]zinc(II)

In the title compound,  $[\text{Zn}(\text{C}_{24}\text{H}_{29}\text{N}_2\text{O}_2\text{S})_2]$ , the Zn atom is four-coordinated by the N atoms of the sulfonamide and quinoline groups. The crystal packing is stabilized by intermolecular  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds.

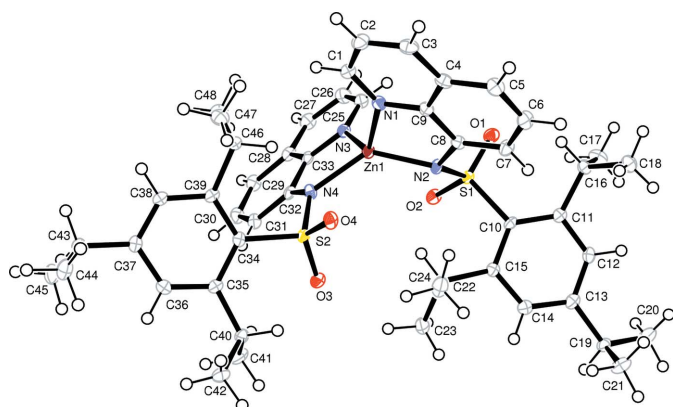
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#### Comment

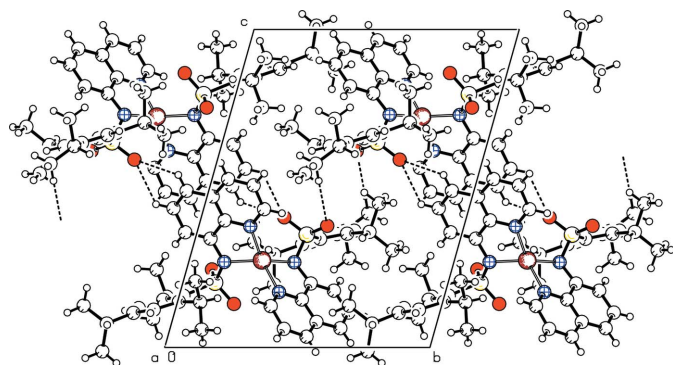
Metal-specific fluorescence probes are of increasing importance in understanding the neurobiology and general cell biology of zinc (Fahrni & O'Halloran, 1999). Several quinoline-based compounds, such as TSQ [6-methoxy-8-(4-tolyl-sulfonamido)quinoline] and zinquin, have been employed to detect zinc in living systems (Nasir *et al.*, 1999; Hendrickson *et al.*, 2003). As part of our study of the chemistry of quinoline-based fluorescence probes for the biological chemistry of zinc (da Silva *et al.*, 2005*a,b,c,d*), the structure of the title compound, (I), has been determined. As reported by Macías *et al.* (2003), the Zn atom is coordinated by four N atoms in a highly distorted tetrahedral geometry. The Zn—N(quinoline) bond lengths [2.074 (2) and 2.049 (2) Å] are slightly larger than the Zn—N(sulfonamide) bonds [1.972 (2) and 1.974 (2) Å], which are in the usual range. Table 1 shows selected bond distances and angles around the central Zn atom.



Quinolinesulfonamidate ligates through the sulfonamide and quinoline N atoms, forming a five-membered ring with the Zn cation. Intermolecular  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds connect the molecules into a three-dimensional network, as shown in the packing diagram (Fig. 2). Details of these hydrogen bonds are given in Table 2.



**Figure 1**  
The molecular structure of (I), showing the atom labelling and with displacement ellipsoids drawn at the 50% probability level.



**Figure 2**  
The molecular packing of (I), with hydrogen bonds shown as dashed lines.

## Experimental

Compound (I) was prepared according to a literature procedure (Macías *et al.*, 2003). Single crystals of (I) suitable for X-ray data collection appeared after a few days from a methanol–dichloromethane (1:1) solution (m.p. 562 K).

### Crystal data

$[\text{Zn}(\text{C}_{24}\text{H}_{29}\text{N}_2\text{O}_2\text{S})_2]$	$Z = 2$
$M_r = 884.47$	$D_x = 1.358 \text{ Mg m}^{-3}$
Triclinic, $P\bar{1}$	Mo $K\alpha$ radiation
$a = 9.9732(6) \text{ \AA}$	Cell parameters from 7555 reflections
$b = 14.3281(8) \text{ \AA}$	$\theta = 2.2\text{--}27.6^\circ$
$c = 17.0425(8) \text{ \AA}$	$\mu = 0.71 \text{ mm}^{-1}$
$\alpha = 71.723(5)^\circ$	$T = 100(2) \text{ K}$
$\beta = 79.277(4)^\circ$	Prism, bright yellow
$\gamma = 69.875(5)^\circ$	$0.36 \times 0.34 \times 0.22 \text{ mm}$
$V = 2162.9(2) \text{ \AA}^3$	

### Data collection

Oxford Diffraction Xcalibur diffractometer with Sapphire CCD detector	15135 measured reflections
$\omega$ and $\varphi$ scans	8527 independent reflections
Absorption correction: analytical ( <i>CrysAlis RED</i> ; Oxford Diffraction, 2004)	7417 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.803$ , $T_{\max} = 0.900$	$R_{\text{int}} = 0.028$
	$\theta_{\text{max}} = 26.4^\circ$
	$h = -11 \rightarrow 12$
	$k = -13 \rightarrow 17$
	$l = -20 \rightarrow 21$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.039$   
 $wR(F^2) = 0.102$   
 $S = 1.08$   
 8527 reflections  
 532 parameters  
 H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0529P)^2 + 1.6914P]$$

where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.001$   
 $\Delta\rho_{\text{max}} = 0.70 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.64 \text{ e \AA}^{-3}$

**Table 1**

Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ).

N1—Zn1	2.0738 (18)	N3—Zn1	2.0487 (18)
N2—Zn1	1.9718 (17)	N4—Zn1	1.9742 (17)
N2—Zn1—N4	131.86 (7)	N2—Zn1—N1	82.95 (7)
N2—Zn1—N3	122.20 (7)	N4—Zn1—N1	119.23 (7)
N4—Zn1—N3	82.37 (7)	N3—Zn1—N1	123.93 (7)

**Table 2**

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

$D\text{—}H\cdots A$	$D\text{—}H$	$H\cdots A$	$D\cdots A$	$D\text{—}H\cdots A$
C20—H20B $\cdots$ O1 <sup>i</sup>	0.98	2.55	3.533 (3)	179
C29—H29 $\cdots$ O2 <sup>ii</sup>	0.95	2.55	3.158 (3)	123
C30—H30 $\cdots$ O2 <sup>ii</sup>	0.95	2.54	3.165 (3)	123

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

H atoms were positioned with idealized geometry, with C—H = 0.95 (aromatic), 0.98 (methyl) or 1.00  $\text{\AA}$  (methine), and were refined as riding, with isotropic displacement parameters set at 1.2 times  $U_{\text{eq}}$  of the parent atom.

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2004); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2004); data reduction: *CrysAlis RED*; 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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