

## 4,5-Dibromo-N-(8-quinolyl)thiophene-2-sulfonamide

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## Key indicators

Single-crystal X-ray study  
 $T = 299$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.011$  Å  
 $R$  factor = 0.045  
 $wR$  factor = 0.103  
Data-to-parameter ratio = 16.0For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.There are two independent molecules in the asymmetric unit of the title compound,  $\text{C}_{13}\text{H}_8\text{Br}_2\text{N}_2\text{O}_2\text{S}_2$ . Intermolecular  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds are observed, linking each non-planar molecule to a symmetry-equivalent molecule.

## Comment

Zinc fluorophores have recently been attracting much interest in biological and environmental applications. Following carbonic anhydrase-based biosensors with fluorescent aromatic sulfonamides, a chemosensor, Zinquin, is now extensively used to study the role of intracellular  $\text{Zn}^{2+}$  in cellular biology (Kimura & Koike, 1998). Our interest in such metal chelators as potential agents for neuroprotection in Alzheimer's disease (Zheng *et al.*, 2005) led to the X-ray study of the title compound, (I).

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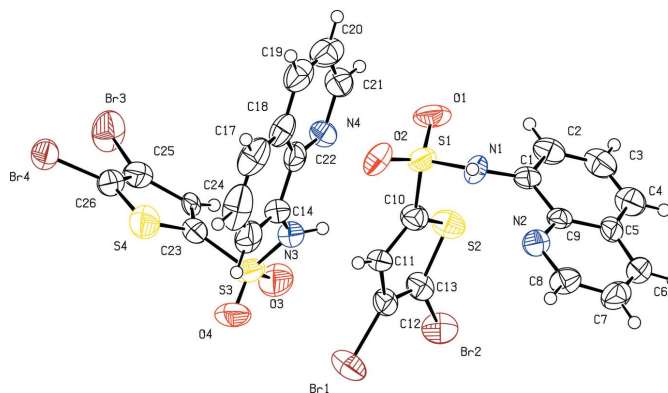
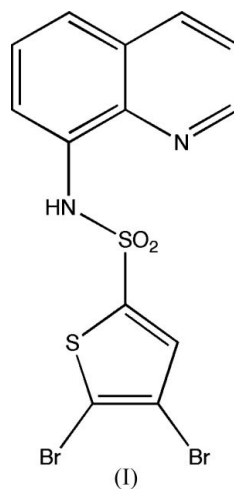


Figure 1

The asymmetric unit of (I), showing the atom labelling and with displacement ellipsoids drawn at the 50% probability level.

The quinoline ring system of each of the two independent molecules in the asymmetric unit (Fig. 1) is nearly planar. The torsion angles C1–N1–S1–C10 and C14–N3–S3–C23 are 59.7 (6) and 60.3 (7)°, respectively. Molecules of the title compound are linked by intermolecular C–H···O hydrogen bonds to form a chain, as shown in Fig. 2 and detailed in Table 1. The contributions of the two inversion twin components refined to 0.23 (1) and 0.77 (1).

### Experimental

The title compound, (I), was prepared according to the literature procedure of Xue *et al.* (2000). Suitable crystals were obtained by recrystallization from methanol–dichloromethane (1:1 *v/v*).

#### Crystal data

C <sub>13</sub> H <sub>8</sub> Br <sub>2</sub> N <sub>2</sub> O <sub>2</sub> S <sub>2</sub>	Mo K $\alpha$ radiation
<i>M<sub>r</sub></i> = 448.15	Cell parameters from 2740 reflections
Orthorhombic, <i>Pna</i> 2 <sub>1</sub>	$\theta$ = 2.8–18.0°
<i>a</i> = 28.318 (2) Å	$\mu$ = 5.66 mm <sup>−1</sup>
<i>b</i> = 7.0471 (5) Å	<i>T</i> = 299 (2) K
<i>c</i> = 15.094 (1) Å	Rod, pink
<i>V</i> = 3012.2 (4) Å <sup>3</sup>	0.28 × 0.10 × 0.08 mm
<i>Z</i> = 8	
<i>D<sub>x</sub></i> = 1.976 Mg m <sup>−3</sup>	

#### Data collection

Oxford Diffraction Xcalibur diffractometer with Sapphire CCD detector	20485 measured reflections
$\omega$ and $\varphi$ scans	6068 independent reflections
Absorption correction: analytical ( <i>CrysAlis RED</i> ; Oxford Diffraction, 2004)	3494 reflections with <i>I</i> > 2 $\sigma$ ( <i>I</i> )
<i>T<sub>min</sub></i> = 0.344, <i>T<sub>max</sub></i> = 0.760	<i>R<sub>int</sub></i> = 0.063
	$\theta_{\max}$ = 26.4°
	<i>h</i> = −35 → 34
	<i>k</i> = −5 → 8
	<i>l</i> = −18 → 18

#### Refinement

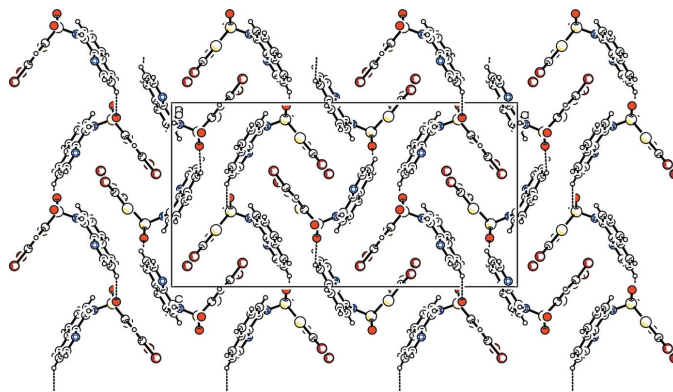
Refinement on <i>F</i> <sup>2</sup>	$w = 1/[\sigma^2(F_o^2) + (0.0499P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.045$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.103$	( $\Delta/\sigma$ ) <sub>max</sub> = 0.022
<i>S</i> = 0.88	$\Delta\rho_{\max} = 0.69 \text{ e } \text{Å}^{-3}$
6068 reflections	$\Delta\rho_{\min} = -0.54 \text{ e } \text{Å}^{-3}$
380 parameters	Absolute structure: Flack (1983)
H-atom parameters constrained	Flack parameter: 0.23 (1), with 2891 Friedel pairs

**Table 1**

Hydrogen-bond geometry (Å, °).

<i>D</i> –H··· <i>A</i>	<i>D</i> –H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> –H··· <i>A</i>
C6–H6···O1 <sup>i</sup>	0.93	2.51	3.215 (9)	133
C20–H20···O3 <sup>ii</sup>	0.93	2.60	3.253 (10)	127

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



**Figure 2**

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

H atoms were positioned with idealized geometry using a riding model (C–H = 0.93 Å and N–H = 0.86 Å) and were refined with isotropic displacement parameters (set to 1.2 times *U<sub>eq</sub>* of the parent atom).

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2003); 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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