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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.002 Å R factor = 0.045 wR factor = 0.112 Data-to-parameter ratio = 34.8

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

2-(2-Thienyl)quinoxaline

The crystal structure of the title compound, $C_{12}H_8N_2S$, was determined at 293 K. The molecule is planar and packs in a herring-bone pattern.

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Comment

We have previously investigated the crystal structures of 2,3dithienylquinoxalines (Crundwell, Sayers *et al.*, 2003) and bromo-substituted 2,3-dithienylquinoxalines (Crundwell *et al.*, 2004). Quinoxalines, in general, have shown versatility in binding a wide variety of metals. We have now expanded our studies to the monosubstituted thienylquinoxalines.



The title compound, (I), sits on a general position. All bond lengths and angles are in agreement with those of other published quinoxalines and thienyl-containing compounds. Thienyl ring geometries and difference maps show no evidence of thienyl ring flip disorders, which are common in molecules with unsubstituted thienyl rings (Crundwell, Sullivan *et al.*, 2003). Like 2-phenylquinoxaline (Yeng & Li, 1989) and 2-(2'-pyridyl)quinoxaline (Kasselouri *et al.*, 1994), the molecule is planar and packs in a herring-bone motif. The interplanar spacing for layers in (I) is 3.537 Å, whereas the interplanar spacings for 2-phenyl and 2-(2'-pyridiyl)quinoxalines are 3.428 and 3.490 Å, respectively.



A view of (I) (Farrugia, 1997). Displacement ellipsoids are drawn at the

50% probability level. H atoms have been omitted for clarity.

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Experimental

Equimolar amounts of thiophen-2-ylglyoxal and 1,2-phenylenediamine were refluxed in 95% ethanol for 2 h. The resulting mixture was filtered and washed with ice-cold ethanol. The precipitate was recrystallized from a boiling ethanol-toluene mixture and crystals were grown from slowly evaporated solutions of ethanol. A light-brown plate that displayed homogeneous birefringence was mounted for diffraction studies. ¹H NMR and UV-vis spectroscopies, as well as melting points of the title compound, agree with literature values (Peter *et al.*, 2004).

> $D_x = 1.359 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation Cell parameters from 1179

> > reflections

 $\theta = 3.8 - 29.7^{\circ}$

 $\begin{aligned} R_{\rm int} &= 0.032\\ \theta_{\rm max} &= 29.7^\circ \end{aligned}$

 $h = -6 \rightarrow 7$

 $k = -22 \rightarrow 22$

 $l = -16 \rightarrow 12$

 $(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta\rho_{\text{max}} = 0.58 \text{ e } \text{\AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.80 \text{ e } \text{\AA}^{-3}$

 $\mu = 0.28 \text{ mm}^{-1}$ T = 293 K Plate, light brown

 $0.32 \times 0.16 \times 0.09 \text{ mm}$

4739 independent reflections

2529 reflections with $I > 3\sigma(I)$

Weighting scheme: Prince modified

1994, Prince, 1982) [weight] =

 $1.0/[3.05T_0(x) + 3.88T_1(x)]$

Chebychev polynomial (Watkin,

+ 0.969 $T_3(x)$], where $x = F_{calc}/F_{max}$

Crystal data

$C_{12}H_8N_2S$
$M_r = 212.27$
Monoclinic, $P2_1/c$
a = 5.5062 (13) Å
<i>b</i> = 16.119 (3) Å
c = 11.939 (3) Å
$\beta = 101.77 \ (2)^{\circ}$
V = 1037.4 (4) Å ³
Z = 4

Data collection

Oxford Diffraction Sapphire3			
diffractometer			
ω scans			
Absorption correction: multi-scan			
(SADABS; Sheldrick, 2003)			
$T_{\min} = 0.891, T_{\max} = 0.989$			
8510 measured reflections			

Refinement

Refinement on F $R[F^2 > 2\sigma(F^2)] = 0.045$ $wR(F^2) = 0.112$ S = 1.004739 reflections 136 parameters Only H-atom coordinates refined

Table 1

Selected geometric parameters (Å, °).

C1-N1	1.3361 (17)	C5-C6	1.401 (3)
C1-C2	1.436 (2)	C6-C7	1.372 (3)
C1-C9	1.457 (2)	C7-C8	1.426 (2)
N1-C8	1.3658 (19)	S1-C9	1.7226 (15)
C2-N2	1.301 (2)	S1-C12	1.7091 (16)
N2-C3	1.381 (2)	C9-C10	1.3893 (19)
C3-C4	1.413 (2)	C10-C11	1.403 (2)
C3-C8	1.414 (2)	C11-C12	1.361 (2)
C4-C5	1.361 (3)		
N1-C1-C2	120.36 (14)	C6-C7-C8	119.95 (16)
N1-C1-C9	118.09 (13)	C7-C8-C3	119.08 (15)
C2-C1-C9	121.55 (12)	C7-C8-N1	119.43 (14)
C1-N1-C8	117.23 (13)	C3-C8-N1	121.50 (13)
C1-C2-N2	123.57 (14)	C9-S1-C12	91.56 (8)
C2-N2-C3	116.69 (14)	C1-C9-S1	119.14 (9)
N2-C3-C4	120.04 (15)	C1-C9-C10	129.99 (13)
N2-C3-C8	120.63 (15)	S1-C9-C10	110.84 (12)
C4-C3-C8	119.30 (15)	C9-C10-C11	112.52 (14)
C3-C4-C5	120.13 (17)	C10-C11-C12	112.66 (14)
C4-C5-C6	121.30 (17)	S1-C12-C11	112.42 (13)
C5-C6-C7	120.21 (16)		

Figure 2

Packing diagram for (I) (Bruno *et al.*, 2002), showing the herringbone motif as viewed along the [001] direction.

H atoms were found in difference maps and their positional parameters were refined. The $U_{iso}(H)$ values were set at 0.05 Å².

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2005); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2005); data reduction: *CrysAlis RED*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *CRYSTALS* (Betteridge *et al.*, 2003); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *MERCURY* (Bruno *et al.*, 2002); software used to prepare material for publication: *CRYSTALS*.

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