

1,4-Bis[[5-(pyridin-4-yl)-1,3,4-oxadiazol-2-yl]sulfanyl]butane

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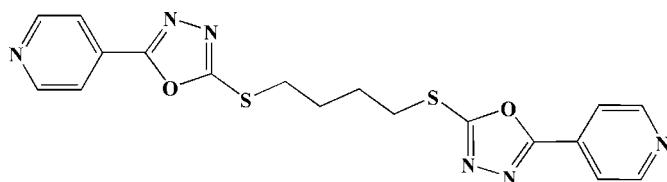
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Key indicators: single-crystal X-ray study; $T = 113\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$; R factor = 0.032; wR factor = 0.089; data-to-parameter ratio = 16.8.

In the centrosymmetric title compound, $\text{C}_{18}\text{H}_{16}\text{N}_6\text{O}_2\text{S}_2$, the 1,3,4-oxadiazole and the attached pyridinyl ring are twisted by $5.3(3)^\circ$.

Related literature

For applications of heterocyclic derivatives, see: Al-Talib *et al.* (1990); Nakagawa *et al.* (1996); Zhang *et al.* (2007). For related structures, see: Wang *et al.* (2010, 2011); Zhao *et al.* (2010).



Experimental

Crystal data

$\text{C}_{18}\text{H}_{16}\text{N}_6\text{O}_2\text{S}_2$
 $M_r = 412.49$

Monoclinic, $P2_1/c$
 $a = 4.9780(6)\text{ \AA}$

$b = 5.7933(7)\text{ \AA}$
 $c = 31.003(4)\text{ \AA}$
 $\beta = 92.588(5)^\circ$
 $V = 893.20(18)\text{ \AA}^3$
 $Z = 2$

Mo $K\alpha$ radiation
 $\mu = 0.33\text{ mm}^{-1}$
 $T = 113\text{ K}$
 $0.20 \times 0.18 \times 0.10\text{ mm}$

Data collection

Rigaku Saturn CCD area-detector diffractometer
Absorption correction: multi-scan (*CrystalClear*; Rigaku/MSC, 2005)
 $T_{\min} = 0.937$, $T_{\max} = 0.962$

8437 measured reflections
2128 independent reflections
1811 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.035$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.032$
 $wR(F^2) = 0.089$
 $S = 1.10$
2128 reflections

127 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.40\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.19\text{ e \AA}^{-3}$

Data collection: *CrystalClear* (Rigaku/MSC, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: KP2310).

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supporting information

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1,4-Bis{[5-(pyridin-4-yl)-1,3,4-oxadiazol-2-yl]sulfanyl}butane

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S1. Comment

Heterocycle derivatives containing N, O and S atoms are under intensive studies due to their wide applications in medicine, industry and coordination chemistry (Al-Talib *et al.*, 1990; Nakagawa *et al.*, 1996; Zhang *et al.*, 2007). We are focusing the synthetic and structural studies on the novel thio-based ligands (Wang *et al.*, 2010, 2011; Zhao *et al.*, 2010). Here we present the synthesis and the crystal structure of the title compound (I), namely, 1,4-bis[5-(pyridin-4-yl)-1,3,4-oxadiazol-2-ylsulfanyl]butane.

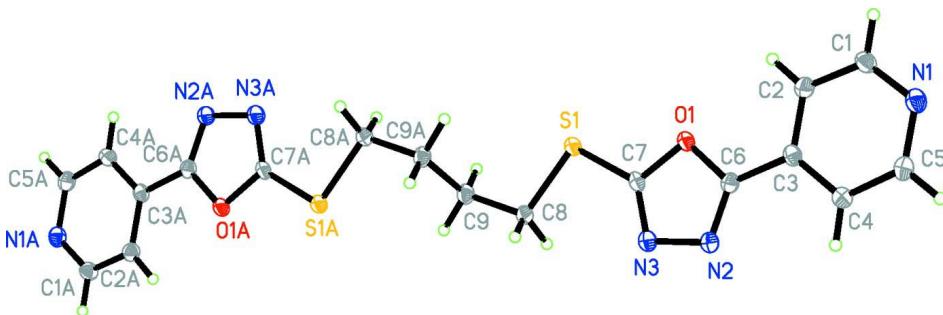
The molecular structure of title compound (I) (Fig. 1)reveals a twofold rotational axis through the mid of the C-C bond of butane group. Therefore, an asymmetric unit comprises a half of the molecule. 1,3,4-Oxadiazole moiety is planar with an r.m.s. deviation of 0.0033 (2) Å and maximum deviation of 0.0052 (2) Å for the atom C7. The dihedral angle between the oxadiazole and its attached pyridinyl ring [r.m.s. deviation = 0.0062 (2) Å] of 5.3 (3)° indicates that they are almost coplanar. As a result of π - π conjugation, the C_{sp^2} -S bond [S1—C7 = 1.722 (13) Å] is significantly shorter than the C_{sp^3} -S bond [S1—C8 = 1.817 (12) Å].

S2. Experimental

A suspension of 5-(pyridin-4-yl)-1,3,4-oxadiazole-2-thiol (2.0 mmol) and 1,1-dibromobutane (1.0 mmol) in ethanol (10 mL) was stirred at room temperature. The reaction progress was monitored *via* TLC. The resulting precipitate was filtered off, washed with cold ethanol, dried and purified to give the target product as light-yellow solid in 87% yield. Crystals of (I) suitable for single-crystal X-ray analysis were grown by slow evaporation of a solution in chloroform-ethanol (1:1).

S3. Refinement

All H atoms were positioned geometrically and refined as riding ($C-H = 0.95-0.99$ Å) and allowed to ride on their parent atoms, with $U_{iso}(H) = 1.2U_{eq}(\text{parent})$.

**Figure 1**

View of the molecule of (I) showing the atom-labelling scheme [symmetry code: (A)- x , - $y + 1$, - $z + 1$]. Displacement ellipsoids are drawn at the 50% probability level.

4-{5-[{4-[(5-(pyridin-4-yl)-1,3,4-oxadiazol-2-yl)sulfanyl]butyl}sulfanyl]-1,3,4-oxadiazol-2-yl}pyridine

Crystal data



$M_r = 412.49$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 4.9780(6)$ Å

$b = 5.7933(7)$ Å

$c = 31.003(4)$ Å

$\beta = 92.588(5)^\circ$

$V = 893.20(18)$ Å³

$Z = 2$

$F(000) = 428$

$D_x = 1.534$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 2798 reflections

$\theta = 2.6\text{--}27.9^\circ$

$\mu = 0.33$ mm⁻¹

$T = 113$ K

Prism, light-yellow

0.20 × 0.18 × 0.10 mm

Data collection

Rigaku Saturn CCD area-detector
diffractometer

Radiation source: rotating anode

Multilayer monochromator

Detector resolution: 14.63 pixels mm⁻¹

φ and ω scans

Absorption correction: multi-scan
(*CrystalClear*; Rigaku/MSC, 2005)

$T_{\min} = 0.937$, $T_{\max} = 0.962$

8437 measured reflections

2128 independent reflections

1811 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.035$

$\theta_{\max} = 27.9^\circ$, $\theta_{\min} = 2.6^\circ$

$h = -4 \rightarrow 6$

$k = -7 \rightarrow 7$

$l = -38 \rightarrow 40$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.032$

$wR(F^2) = 0.089$

$S = 1.10$

2128 reflections

127 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

$w = 1/[\sigma^2(F_o^2) + (0.0531P)^2 + 0.054P]$
where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.40$ e Å⁻³

$\Delta\rho_{\min} = -0.19$ e Å⁻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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.25227 (6)	0.35176 (5)	0.422550 (9)	0.01776 (12)
O1	0.61211 (17)	0.17771 (15)	0.37253 (3)	0.0163 (2)
N1	1.3455 (2)	-0.0526 (2)	0.27364 (3)	0.0197 (2)
N2	0.7346 (2)	-0.15583 (19)	0.40229 (3)	0.0186 (2)
N3	0.5348 (2)	-0.05632 (19)	0.42735 (3)	0.0189 (2)
C1	1.1858 (3)	0.1307 (2)	0.27715 (4)	0.0190 (3)
H1	1.2063	0.2551	0.2576	0.023*
C2	0.9910 (3)	0.1507 (2)	0.30774 (4)	0.0179 (3)
H2	0.8797	0.2836	0.3086	0.021*
C3	0.9634 (2)	-0.0287 (2)	0.33688 (4)	0.0155 (3)
C4	1.1268 (3)	-0.2220 (2)	0.33375 (4)	0.0185 (3)
H4	1.1125	-0.3479	0.3531	0.022*
C5	1.3118 (3)	-0.2262 (2)	0.30146 (4)	0.0199 (3)
H5	1.4206	-0.3598	0.2990	0.024*
C6	0.7718 (2)	-0.0138 (2)	0.37112 (4)	0.0153 (3)
C7	0.4729 (2)	0.1379 (2)	0.40882 (4)	0.0152 (3)
C8	0.1260 (3)	0.2236 (2)	0.47114 (4)	0.0177 (3)
H8A	0.0331	0.0764	0.4640	0.021*
H8B	0.2770	0.1908	0.4921	0.021*
C9	-0.0695 (3)	0.3927 (2)	0.49063 (4)	0.0178 (3)
H9A	-0.2052	0.4403	0.4680	0.021*
H9B	-0.1652	0.3137	0.5137	0.021*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0216 (2)	0.01715 (19)	0.01492 (18)	0.00366 (11)	0.00549 (12)	0.00101 (11)
O1	0.0174 (4)	0.0179 (5)	0.0140 (4)	0.0017 (3)	0.0045 (3)	0.0003 (3)
N1	0.0180 (5)	0.0223 (6)	0.0191 (5)	-0.0024 (4)	0.0030 (4)	-0.0030 (4)
N2	0.0208 (6)	0.0179 (6)	0.0176 (5)	0.0020 (4)	0.0053 (4)	-0.0002 (4)
N3	0.0201 (5)	0.0193 (6)	0.0177 (5)	0.0022 (4)	0.0053 (4)	-0.0003 (4)
C1	0.0194 (7)	0.0202 (7)	0.0176 (6)	-0.0024 (5)	0.0037 (5)	0.0011 (5)
C2	0.0178 (6)	0.0173 (6)	0.0186 (6)	0.0012 (5)	0.0023 (5)	-0.0005 (5)
C3	0.0146 (6)	0.0185 (6)	0.0132 (5)	-0.0014 (5)	-0.0001 (4)	-0.0016 (4)
C4	0.0205 (6)	0.0185 (6)	0.0166 (6)	0.0009 (5)	0.0022 (5)	0.0009 (5)
C5	0.0193 (6)	0.0200 (6)	0.0207 (6)	0.0025 (5)	0.0029 (5)	-0.0026 (5)

C6	0.0142 (6)	0.0160 (6)	0.0158 (6)	0.0011 (4)	0.0003 (4)	-0.0017 (4)
C7	0.0155 (6)	0.0188 (6)	0.0114 (5)	-0.0013 (4)	0.0028 (4)	-0.0014 (4)
C8	0.0215 (6)	0.0162 (6)	0.0158 (6)	-0.0010 (5)	0.0055 (5)	-0.0007 (5)
C9	0.0173 (6)	0.0182 (6)	0.0183 (6)	-0.0013 (5)	0.0042 (5)	-0.0030 (5)

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

S1—C7	1.7218 (13)	C2—H2	0.9500
S1—C8	1.8171 (12)	C3—C4	1.3902 (17)
O1—C6	1.3669 (14)	C3—C6	1.4615 (16)
O1—C7	1.3675 (14)	C4—C5	1.3906 (17)
N1—C1	1.3337 (17)	C4—H4	0.9500
N1—C5	1.3404 (17)	C5—H5	0.9500
N2—C6	1.2888 (16)	C8—C9	1.5251 (17)
N2—N3	1.4121 (15)	C8—H8A	0.9900
N3—C7	1.2941 (16)	C8—H8B	0.9900
C1—C2	1.3914 (18)	C9—C9 ⁱ	1.525 (2)
C1—H1	0.9500	C9—H9A	0.9900
C2—C3	1.3878 (17)	C9—H9B	0.9900
C7—S1—C8	99.16 (6)	C4—C5—H5	118.1
C6—O1—C7	101.91 (9)	N2—C6—O1	113.00 (10)
C1—N1—C5	116.91 (11)	N2—C6—C3	128.89 (11)
C6—N2—N3	106.29 (10)	O1—C6—C3	118.07 (10)
C7—N3—N2	105.69 (10)	N3—C7—O1	113.10 (11)
N1—C1—C2	123.91 (12)	N3—C7—S1	131.13 (10)
N1—C1—H1	118.0	O1—C7—S1	115.76 (9)
C2—C1—H1	118.0	C9—C8—S1	108.47 (9)
C3—C2—C1	118.29 (12)	C9—C8—H8A	110.0
C3—C2—H2	120.9	S1—C8—H8A	110.0
C1—C2—H2	120.9	C9—C8—H8B	110.0
C2—C3—C4	118.84 (11)	S1—C8—H8B	110.0
C2—C3—C6	121.06 (11)	H8A—C8—H8B	108.4
C4—C3—C6	120.08 (11)	C9 ⁱ —C9—C8	112.82 (13)
C3—C4—C5	118.21 (12)	C9 ⁱ —C9—H9A	109.0
C3—C4—H4	120.9	C8—C9—H9A	109.0
C5—C4—H4	120.9	C9 ⁱ —C9—H9B	109.0
N1—C5—C4	123.80 (12)	C8—C9—H9B	109.0
N1—C5—H5	118.1	H9A—C9—H9B	107.8
C6—N2—N3—C7	-0.50 (14)	C2—C3—C6—N2	-174.70 (13)
C5—N1—C1—C2	0.29 (19)	C4—C3—C6—N2	3.7 (2)
N1—C1—C2—C3	1.2 (2)	C2—C3—C6—O1	2.71 (17)
C1—C2—C3—C4	-1.37 (19)	C4—C3—C6—O1	-178.93 (11)
C1—C2—C3—C6	177.01 (11)	N2—N3—C7—O1	0.93 (14)
C2—C3—C4—C5	0.23 (18)	N2—N3—C7—S1	-177.95 (10)
C6—C3—C4—C5	-178.17 (11)	C6—O1—C7—N3	-0.96 (13)
C1—N1—C5—C4	-1.56 (19)	C6—O1—C7—S1	178.10 (8)

C3—C4—C5—N1	1.3 (2)	C8—S1—C7—N3	-0.56 (14)
N3—N2—C6—O1	-0.09 (14)	C8—S1—C7—O1	-179.42 (9)
N3—N2—C6—C3	177.42 (12)	C7—S1—C8—C9	177.46 (9)
C7—O1—C6—N2	0.61 (13)	S1—C8—C9—C9 ⁱ	-69.35 (15)
C7—O1—C6—C3	-177.20 (11)		

Symmetry code: (i) $-x, -y+1, -z+1$.