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## Structure Reports

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# Di-4-pyridyl disulfide–isophthalic acid (1/1)

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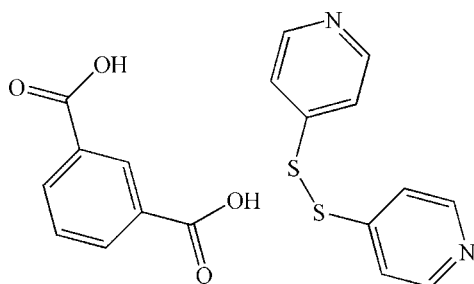
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 Key indicators: single-crystal X-ray study;  $T = 295$  K; mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å;  $R$  factor = 0.043;  $wR$  factor = 0.135; data-to-parameter ratio = 17.2.

In the title 1:1 cocrystal,  $\text{C}_{10}\text{H}_8\text{N}_2\text{S}_2 \cdot \text{C}_8\text{H}_6\text{O}_4$ , the asymmetric unit contains an isophthalic acid molecule and a 4,4'-dipyridyl disulfide molecule. The two carboxyl groups of isophthalic acid interact with neighbouring 4,4'-dipyridyl disulfide molecules through  $\text{O}-\text{H} \cdots \text{N}$  hydrogen bonds, forming a one-dimensional zigzag chain. Neighbouring chains are linked to each other *via*  $\pi-\pi$  stacking interactions between the pyridyl rings of adjacent 4,4'-dipyridyl disulfide molecules [centroid-centroid distance = 3.7346 (6) Å], resulting in a layered motif. The dihedral angle between pyridine rings of 84.13 (7)° and the  $\text{C}-\text{S}-\text{S}-\text{C}$  torsion angle of 91.95 (1)° confirm the *gauche* conformation of 4,4'-dipyridyl disulfide.

## Related literature

For ligands with two 4-pyridyl donors, see: Biradha *et al.* (2006); Sun *et al.* (2006); He *et al.* (2008); Suen *et al.* (2005). For related structures, see: Ranjbar *et al.* (2007).



## Experimental

### Crystal data

$\text{C}_{10}\text{H}_8\text{N}_2\text{S}_2 \cdot \text{C}_8\text{H}_6\text{O}_4$   
 $M_r = 386.43$   
 Monoclinic,  $P2_1/c$   
 $a = 5.9616$  (12) Å

$b = 10.024$  (2) Å  
 $c = 29.797$  (6) Å  
 $\beta = 93.71$  (3)°  
 $V = 1776.9$  (6) Å<sup>3</sup>

$Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.33$  mm<sup>-1</sup>

$T = 295$  K  
 $0.29 \times 0.20 \times 0.11$  mm

### Data collection

Rigaku R-Axis RAPID diffractometer  
 Absorption correction: multi-scan (ABSCOR; Higashi, 1995)  
 $T_{\min} = 0.920$ ,  $T_{\max} = 0.964$

16923 measured reflections  
 4039 independent reflections  
 2330 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.048$

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.043$   
 $wR(F^2) = 0.135$   
 $S = 1.08$   
 4039 reflections

235 parameters  
 H-atom parameters constrained  
 $\Delta\rho_{\max} = 0.30$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.41$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$\text{O2}-\text{H2C} \cdots \text{N1}^i$	0.99	1.64	2.629 (3)	175
$\text{O4}-\text{H4C} \cdots \text{N2}^{ii}$	0.81	1.85	2.651 (3)	176

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

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2004); 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: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2150).

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## supporting information

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## Di-4-pyridyl disulfide–isophthalic acid (1/1)

Dan-Jun Wang, Jun-Ming Zhao, Jie Zhang and Jian-Li Lin

### S1. Comment

The ligands having two 4-pyridyl donors, *e.g.*, 4,4'-bipyridine (Biradha *et al.*, 2006), 1,2-bis(4-pyridyl)ethane (Sun *et al.*, 2006) and 1,3-bis(4-pyridyl)propane (He *et al.*, 2008) have been intensively employed for the construction of coordination polymers. Compared with the above examples, di-4-pyridyl disulfide is seldom used for research. It shows a twisted structure, with a C—S—S—C torsion angle of approximately 90°. More importantly, the ligand has axial chirality, which generates *M*- and *P*- enantiomers as shown in Fig. 3. It means that the use of this ligand possibly can produce the complex with a non-centrosymmetric space group (Suen *et al.*, 2005). As we know, some special properties, *e.g.*, triboluminescence, second harmonic generation and ferroelectricity are only found in these materials. For this consideration, we mixed this ligand and carboxylate ligand hoping to gain coordination polymer with special properties. However, a crystal suitable for X-ray diffraction was obtained during the synthesis unexpectedly. In this paper we report the crystal structure of the title cocrystal.

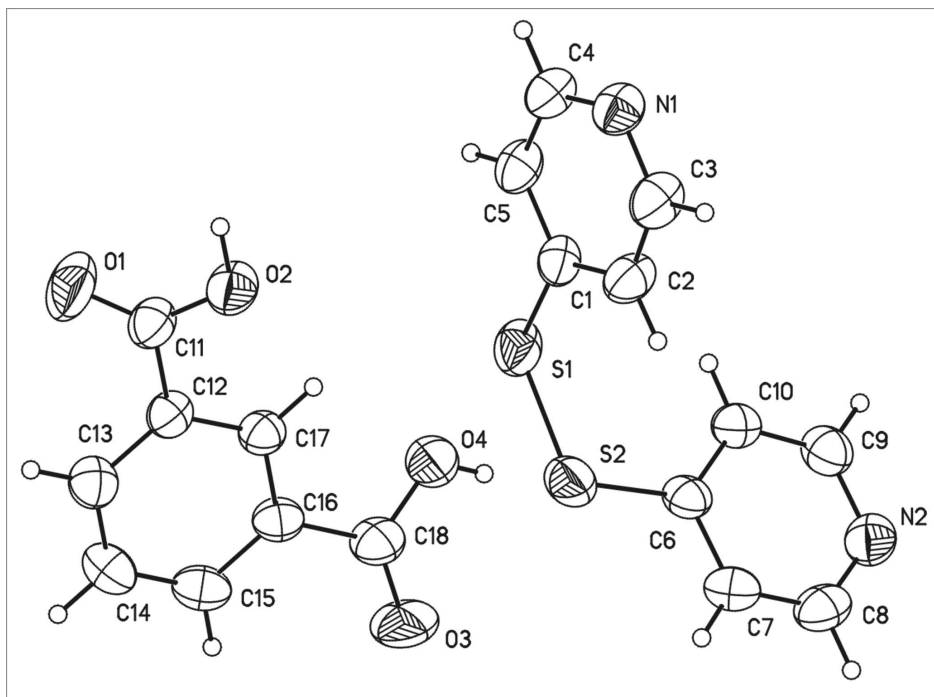
The asymmetric unit of the title cocrystal consists of one isophthalic acid molecule and one *P*- form di-4-pyridyl disulfide molecule (Fig. 1). The two carboxylic groups of the isophthalic acid are hydrogen bonded with the corresponding di-4-pyridyl disulfide molecules (O2—H2C $\cdots$ N1<sup>i</sup> and O4—H4C $\cdots$ N2<sup>ii</sup> (Table 1)) generating a one-dimensional zigzag chain along the *c* axis. The neighbouring chains are further linked to each other *via*  $\pi$ — $\pi$  packing interactions between the pyridyl rings of adjacent di-4-pyridyl disulfide molecules resulting in a two-dimensional layered structure (Fig. 2). The centroid-centroid distance is 3.7346 (6) Å, the C—S—S—C torsion angle is 91.95 (1)°, and the pyridyl ring planes form a dihedral angle of 84.13 (7)°. The crystal structures of closely related cocrystals have been reported (Ranjbar *et al.*, 2007).

### S2. Experimental

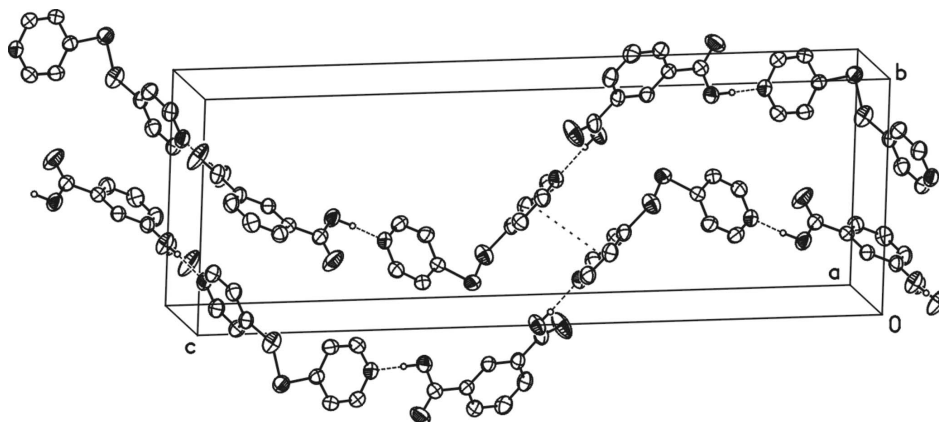
Dropwise addition of Na<sub>2</sub>CO<sub>3</sub> (0.5 ml 1.0 *M*) to an aqueous solution of Zn(NO<sub>3</sub>)<sub>2</sub>·6H<sub>2</sub>O (0.0808 g, 0.25 mmol) in 4 ml H<sub>2</sub>O produced white precipitate, which was then centrifuged and washed with distilled water six times. The collected precipitate was subsequently moved to a stirred suspension of isophthalic acid (0.0817 g, 0.5 mmol) in a mixed solvent composed of EtOH (10 ml) and H<sub>2</sub>O (20 ml), and further stirred at 353 K for 1 h, followed by the addition of an ethanolic solution of 0.1120 g (0.5 mmol) di-4-pyridyl disulfide in 5 ml EtOH. The resulting mixture was further stirred at 343 K for 30 min and filtered off. Slow evaporation of the colorless filtrate at room temperature for one week gave colorless block crystals (yield: 0.05 g).

### S3. Refinement

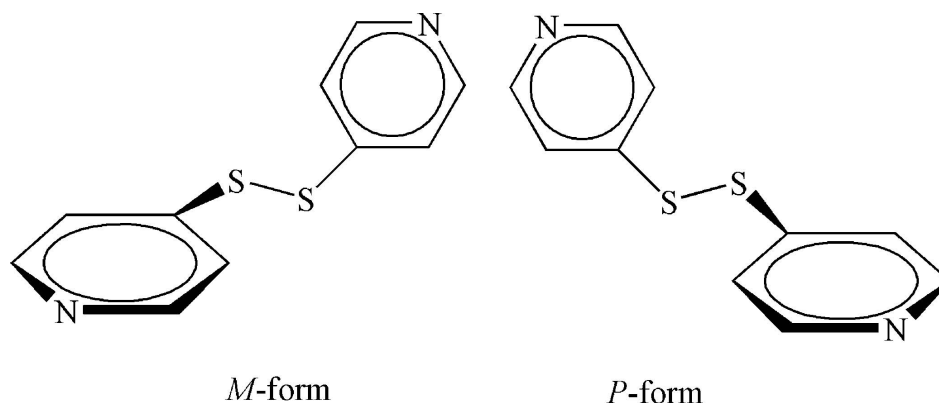
H atoms bonded to C atoms were placed in geometrically calculated position and were refined using a riding model, with  $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$ . H atoms attached to O atoms were found in a difference Fourier synthesis and were refined using a riding model, with the O—H distances fixed as initially found and with  $U_{\text{iso}}(\text{H})$  values set at  $1.2 U_{\text{eq}}(\text{O})$ .

**Figure 1**

A view of the molecular structure of the title cocrystal, displacement ellipsoids are drawn at the 45% probability level.

**Figure 2**

The crystal packing diagram, showing the  $\pi$ — $\pi$  stacking and hydrogen bonds as dash lines.

**Figure 3**

The *M*- and *P*-enantiomers of di-4-pyridyl disulfide.

### Di-4-pyridyl disulfide–isophthalic acid (1/1)

#### Crystal data

$C_{10}H_8N_2S_2 \cdot C_8H_6O_4$

$M_r = 386.43$

Monoclinic,  $P2_1/c$

Hall symbol:  $-P\ 2ybc$

$a = 5.9616$  (12) Å

$b = 10.024$  (2) Å

$c = 29.797$  (6) Å

$\beta = 93.71$  (3)°

$V = 1776.9$  (6) Å<sup>3</sup>

$Z = 4$

$F(000) = 800$

$D_x = 1.445$  Mg m<sup>-3</sup>

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

Cell parameters from 16923 reflections

$\theta = 3.4$ – $27.4$ °

$\mu = 0.33$  mm<sup>-1</sup>

$T = 295$  K

Platelet, colorless

$0.29 \times 0.20 \times 0.11$  mm

#### Data collection

Rigaku R-AXIS RAPID

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 0 pixels mm<sup>-1</sup>

$\omega$  scans

Absorption correction: multi-scan

(*ABSCOR*; Higashi, 1995)

$T_{\min} = 0.920$ ,  $T_{\max} = 0.964$

16923 measured reflections

4039 independent reflections

2330 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.048$

$\theta_{\max} = 27.4$ °,  $\theta_{\min} = 3.4$ °

$h = -7 \rightarrow 7$

$k = -12 \rightarrow 12$

$l = -38 \rightarrow 38$

#### Refinement

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.043$

$wR(F^2) = 0.135$

$S = 1.08$

4039 reflections

235 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

$w = 1/[\sigma^2(F_o^2) + (0.0469P)^2 + 0.8478P]$

where  $P = (F_o^2 + 2F_c^2)/3$

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

$\Delta\rho_{\max} = 0.30$  e Å<sup>-3</sup>

$\Delta\rho_{\min} = -0.41$  e Å<sup>-3</sup>

*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 ( $\text{\AA}^2$ )*

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.1566 (4)	0.1783 (3)	0.32831 (8)	0.0472 (6)
C2	0.1371 (5)	0.0516 (3)	0.31039 (10)	0.0570 (7)
H2A	0.2419	-0.0142	0.3187	0.068*
C3	-0.0411 (5)	0.0247 (3)	0.27995 (10)	0.0616 (8)
H3A	-0.0545	-0.0611	0.2682	0.074*
N1	-0.1955 (4)	0.1147 (2)	0.26642 (7)	0.0541 (6)
C4	-0.1742 (5)	0.2369 (3)	0.28378 (9)	0.0542 (7)
H4A	-0.2806	0.3011	0.2748	0.065*
C5	-0.0019 (5)	0.2723 (3)	0.31427 (9)	0.0526 (7)
H5A	0.0079	0.3589	0.3254	0.063*
S1	0.36292 (14)	0.23223 (8)	0.36981 (3)	0.0629 (2)
S2	0.59614 (12)	0.08517 (9)	0.37425 (3)	0.0664 (3)
C6	0.5151 (4)	-0.0234 (3)	0.41697 (8)	0.0467 (6)
C7	0.6649 (4)	-0.1247 (3)	0.42909 (9)	0.0543 (7)
H7A	0.7977	-0.1335	0.4146	0.065*
C8	0.6162 (5)	-0.2124 (3)	0.46265 (10)	0.0588 (8)
H8A	0.7195	-0.2791	0.4708	0.071*
N2	0.4262 (4)	-0.2055 (3)	0.48407 (7)	0.0571 (6)
C9	0.2814 (5)	-0.1082 (3)	0.47191 (9)	0.0558 (7)
H9A	0.1483	-0.1031	0.4865	0.067*
C10	0.3170 (4)	-0.0152 (3)	0.43917 (9)	0.0520 (7)
H10A	0.2119	0.0513	0.4321	0.062*
O1	0.6473 (5)	0.7527 (3)	0.28620 (9)	0.0996 (9)
O2	0.5074 (3)	0.5519 (2)	0.29642 (6)	0.0634 (6)
H2C	0.3967	0.5742	0.2715	0.095*
C11	0.6474 (5)	0.6489 (3)	0.30626 (9)	0.0528 (7)
C12	0.8127 (4)	0.6179 (3)	0.34480 (8)	0.0452 (6)
C13	1.0212 (5)	0.6802 (3)	0.34736 (9)	0.0541 (7)
H13A	1.0553	0.7426	0.3257	0.065*
C14	1.1780 (5)	0.6495 (3)	0.38191 (10)	0.0603 (8)
H14A	1.3179	0.6909	0.3833	0.072*
C15	1.1292 (4)	0.5580 (3)	0.41451 (9)	0.0574 (8)
H15A	1.2368	0.5368	0.4374	0.069*
C16	0.9196 (4)	0.4977 (3)	0.41308 (8)	0.0467 (6)
C17	0.7617 (4)	0.5277 (3)	0.37792 (8)	0.0451 (6)

H17A	0.6214	0.4868	0.3767	0.054*
C18	0.8684 (5)	0.3991 (3)	0.44862 (9)	0.0566 (7)
O3	1.0115 (4)	0.3410 (3)	0.47123 (9)	0.0996 (9)
O4	0.6531 (3)	0.3815 (2)	0.45246 (7)	0.0695 (6)
H4C	0.6275	0.3313	0.4726	0.104*

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C1	0.0550 (16)	0.0416 (15)	0.0445 (14)	-0.0047 (12)	0.0007 (12)	0.0048 (12)
C2	0.0646 (18)	0.0438 (17)	0.0606 (17)	0.0055 (14)	-0.0107 (14)	-0.0019 (13)
C3	0.078 (2)	0.0459 (17)	0.0589 (17)	-0.0006 (16)	-0.0134 (16)	-0.0092 (14)
N1	0.0640 (15)	0.0536 (15)	0.0433 (12)	-0.0002 (12)	-0.0078 (11)	0.0003 (11)
C4	0.0624 (17)	0.0485 (17)	0.0509 (15)	0.0066 (14)	-0.0027 (14)	0.0038 (13)
C5	0.0681 (18)	0.0391 (15)	0.0502 (15)	0.0009 (14)	-0.0005 (14)	-0.0019 (12)
S1	0.0717 (5)	0.0503 (5)	0.0638 (5)	-0.0141 (4)	-0.0174 (4)	0.0058 (4)
S2	0.0482 (4)	0.0838 (6)	0.0672 (5)	-0.0068 (4)	0.0036 (3)	0.0218 (4)
C6	0.0386 (13)	0.0557 (17)	0.0447 (13)	-0.0022 (12)	-0.0055 (11)	-0.0001 (12)
C7	0.0433 (15)	0.0660 (19)	0.0527 (16)	0.0059 (14)	-0.0033 (12)	-0.0024 (14)
C8	0.0587 (17)	0.0601 (19)	0.0552 (16)	0.0083 (15)	-0.0144 (14)	0.0001 (15)
N2	0.0612 (15)	0.0606 (16)	0.0477 (13)	-0.0051 (13)	-0.0106 (11)	0.0067 (11)
C9	0.0489 (15)	0.067 (2)	0.0510 (15)	-0.0048 (15)	0.0008 (13)	0.0040 (14)
C10	0.0434 (14)	0.0571 (18)	0.0551 (15)	0.0022 (13)	-0.0007 (12)	0.0064 (14)
O1	0.118 (2)	0.0680 (17)	0.1053 (19)	-0.0274 (15)	-0.0499 (16)	0.0421 (15)
O2	0.0698 (13)	0.0577 (13)	0.0589 (12)	-0.0097 (11)	-0.0244 (10)	0.0090 (10)
C11	0.0612 (17)	0.0465 (17)	0.0495 (15)	-0.0013 (14)	-0.0059 (13)	0.0043 (13)
C12	0.0509 (15)	0.0405 (15)	0.0435 (13)	0.0004 (12)	-0.0026 (12)	-0.0027 (11)
C13	0.0570 (17)	0.0544 (18)	0.0507 (15)	-0.0046 (14)	0.0011 (13)	0.0000 (13)
C14	0.0458 (15)	0.071 (2)	0.0634 (18)	-0.0101 (15)	-0.0001 (14)	-0.0047 (16)
C15	0.0455 (15)	0.072 (2)	0.0526 (16)	0.0031 (15)	-0.0103 (13)	-0.0050 (15)
C16	0.0452 (14)	0.0530 (17)	0.0408 (13)	0.0033 (12)	-0.0054 (11)	-0.0026 (12)
C17	0.0446 (14)	0.0447 (15)	0.0449 (13)	0.0002 (12)	-0.0044 (11)	-0.0011 (12)
C18	0.0541 (16)	0.069 (2)	0.0454 (15)	0.0018 (15)	-0.0095 (13)	0.0063 (14)
O3	0.0644 (14)	0.135 (2)	0.0961 (18)	0.0153 (16)	-0.0159 (13)	0.0616 (18)
O4	0.0580 (12)	0.0864 (16)	0.0624 (12)	-0.0072 (11)	-0.0107 (10)	0.0307 (11)

*Geometric parameters (Å, °)*

C1—C5	1.380 (4)	C9—H9A	0.9300
C1—C2	1.380 (4)	C10—H10A	0.9300
C1—S1	1.771 (3)	O1—C11	1.200 (3)
C2—C3	1.378 (4)	O2—C11	1.302 (3)
C2—H2A	0.9300	O2—H2C	0.9857
C3—N1	1.333 (4)	C11—C12	1.497 (4)
C3—H3A	0.9300	C12—C17	1.387 (4)
N1—C4	1.333 (4)	C12—C13	1.388 (4)
C4—C5	1.373 (4)	C13—C14	1.379 (4)
C4—H4A	0.9300	C13—H13A	0.9300

C5—H5A	0.9300	C14—C15	1.381 (4)
S1—S2	2.0248 (13)	C14—H14A	0.9300
S2—C6	1.766 (3)	C15—C16	1.386 (4)
C6—C7	1.385 (4)	C15—H15A	0.9300
C6—C10	1.393 (3)	C16—C17	1.395 (3)
C7—C8	1.376 (4)	C16—C18	1.495 (4)
C7—H7A	0.9300	C17—H17A	0.9300
C8—N2	1.338 (4)	C18—O3	1.203 (3)
C8—H8A	0.9300	C18—O4	1.308 (3)
N2—C9	1.337 (4)	O4—H4C	0.8043
C9—C10	1.376 (4)		
C5—C1—C2	118.2 (3)	C10—C9—H9A	118.0
C5—C1—S1	115.7 (2)	C9—C10—C6	118.1 (3)
C2—C1—S1	126.0 (2)	C9—C10—H10A	120.9
C3—C2—C1	118.5 (3)	C6—C10—H10A	120.9
C3—C2—H2A	120.8	C11—O2—H2C	112.9
C1—C2—H2A	120.8	O1—C11—O2	123.7 (3)
N1—C3—C2	123.7 (3)	O1—C11—C12	122.7 (3)
N1—C3—H3A	118.2	O2—C11—C12	113.5 (2)
C2—C3—H3A	118.2	C17—C12—C13	119.4 (2)
C3—N1—C4	117.3 (2)	C17—C12—C11	121.2 (2)
N1—C4—C5	122.8 (3)	C13—C12—C11	119.5 (2)
N1—C4—H4A	118.6	C14—C13—C12	120.1 (3)
C5—C4—H4A	118.6	C14—C13—H13A	119.9
C4—C5—C1	119.5 (3)	C12—C13—H13A	119.9
C4—C5—H5A	120.2	C13—C14—C15	120.6 (3)
C1—C5—H5A	120.2	C13—C14—H14A	119.7
C1—S1—S2	105.44 (10)	C15—C14—H14A	119.7
C6—S2—S1	106.08 (10)	C14—C15—C16	119.9 (3)
C7—C6—C10	118.1 (3)	C14—C15—H15A	120.0
C7—C6—S2	116.0 (2)	C16—C15—H15A	120.0
C10—C6—S2	125.9 (2)	C15—C16—C17	119.5 (3)
C8—C7—C6	119.7 (3)	C15—C16—C18	119.4 (2)
C8—C7—H7A	120.2	C17—C16—C18	121.0 (2)
C6—C7—H7A	120.2	C12—C17—C16	120.4 (2)
N2—C8—C7	122.6 (3)	C12—C17—H17A	119.8
N2—C8—H8A	118.7	C16—C17—H17A	119.8
C7—C8—H8A	118.7	O3—C18—O4	123.4 (3)
C9—N2—C8	117.4 (2)	O3—C18—C16	123.2 (3)
N2—C9—C10	124.0 (3)	O4—C18—C16	113.4 (2)
N2—C9—H9A	118.0	C18—O4—H4C	112.6

*Hydrogen-bond geometry (Å, °)*

<i>D</i> —H $\cdots$ <i>A</i>	<i>D</i> —H	H $\cdots$ <i>A</i>	<i>D</i> $\cdots$ <i>A</i>	<i>D</i> —H $\cdots$ <i>A</i>
O2—H2C $\cdots$ N1 <sup>i</sup>	0.99	1.64	2.629 (3)	175

O4—H4C···N2 <sup>ii</sup>	0.81	1.85	2.651 (3)	176
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Symmetry codes: (i)  $-x, y+1/2, -z+1/2$ ; (ii)  $-x+1, -y, -z+1$ .