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Bis(4-fluoroanilinium) sulfate

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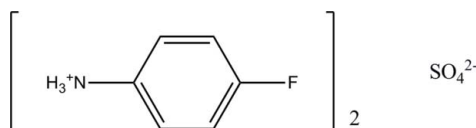
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.038; wR factor = 0.103; data-to-parameter ratio = 22.5.

In the crystal of the title molecular salt, $2\text{C}_6\text{H}_7\text{FN}^+\cdot\text{SO}_4^{2-}$, the cations and anions are linked by $\text{N}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds into sheets parallel to the ab plane. The crystal studied was found to be a racemic twin with a 0.50 (10):0.50 (10) domain ratio.

Related literature

For related literature on phase transition dielectric materials, see: Fu *et al.* (2007, 2008, 2009); Fu & Xiong (2008). For hydrogen bonding studies, see: Zimmerman & Corbin (2000); Brunsveld *et al.* (2001); Desiraju (2002); Steiner (2002); Desiraju & Steiner (1999); Boutobba *et al.* (2010). For reference bond-length data, see: Allen *et al.* (1987). For a related crystal structure, see: Boutobba *et al.* (2010)



Experimental

Crystal data

 $2\text{C}_6\text{H}_7\text{FN}^+\cdot\text{SO}_4^{2-}$ $M_r = 320.31$ Orthorhombic, $P2_12_12_1$ $a = 6.2907$ (5) Å $b = 7.4155$ (6) Å $c = 30.168$ (3) Å $V = 1407.3$ (2) Å³ $Z = 4$ Mo $K\alpha$ radiation $\mu = 0.27$ mm⁻¹ $T = 296$ K $0.58 \times 0.12 \times 0.07$ mm

Data collection

Bruker SMART APEXII DUO

CCD area-detector diffractometer

Absorption correction: multi-scan

(SADABS; Bruker, 2009)

 $T_{\min} = 0.859$, $T_{\max} = 0.982$

31062 measured reflections

4292 independent reflections

3503 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.044$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.038$ $wR(F^2) = 0.103$ $S = 0.98$

4292 reflections

191 parameters

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.18$ e Å⁻³ $\Delta\rho_{\text{min}} = -0.39$ e Å⁻³

Absolute structure: Flack (1983),

1794 Friedel pairs

Flack parameter: 0.50 (10)

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H3N1}\cdots\text{O3}$	0.97	2.09	2.887 (3)	138
$\text{N2}-\text{H2N2}\cdots\text{O3}$	0.85	1.90	2.741 (3)	168
$\text{N1}-\text{H1N1}\cdots\text{O1}^{\text{i}}$	0.86	1.92	2.761 (2)	162
$\text{N1}-\text{H2N1}\cdots\text{O3}^{\text{ii}}$	0.82	2.25	2.967 (2)	146
$\text{N1}-\text{H2N1}\cdots\text{O4}^{\text{ii}}$	0.82	2.55	3.151 (3)	131
$\text{N1}-\text{H3N1}\cdots\text{O2}^{\text{iii}}$	0.97	2.24	2.925 (2)	126
$\text{N2}-\text{H1N2}\cdots\text{O4}^{\text{iv}}$	0.94	1.78	2.7121 (18)	170
$\text{N2}-\text{H3N2}\cdots\text{O2}^{\text{v}}$	0.99	1.72	2.702 (2)	170
$\text{C11}-\text{H11A}\cdots\text{O4}^{\text{iii}}$	0.93	2.53	3.374 (3)	151

Symmetry codes: (i) $x, y + 1, z$; (ii) $x - \frac{1}{2}, -y + \frac{3}{2}, -z$; (iii) $x + \frac{1}{2}, -y + \frac{3}{2}, -z$; (iv) $x + \frac{1}{2}, -y + \frac{1}{2}, -z$; (v) $x + 1, y, z$.

Data collection: APEX2 (Bruker, 2009); cell refinement: SAINT (Bruker, 2009); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL and PLATON (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: WN2446).

References

- Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). *J. Chem. Soc. Perkin Trans. 2*, pp. S1–19.
- Boutobba, Z., Direm, A. & Benali-Cherif, N. (2010). *Acta Cryst. E* **66**, o595–o596.
- Bruker (2009). *SADABS, APEX2* and *SAINTE*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Brunsveld, L., Folmer, B. J. B., Meijer, E. W. & Sijbesma, R. P. (2001). *Chem. Rev.* **101**, 4071–4097.
- Desiraju, G. R. (2002). *Acc. Chem. Res.* **35**, 565–573.
- Desiraju, G. R. & Steiner, T. (1999). *The Weak Hydrogen Bond in Structural Chemistry and Biology*, p. 507. New York: Oxford University Press.
- Flack, H. D. (1983). *Acta Cryst. A* **39**, 876–881.
- Fu, D.-W., Ge, J.-Z., Dai, J., Ye, H.-Y. & Qu, Z.-R. (2009). *Inorg. Chem. Commun.* **12**, 994–997.
- Fu, D.-W., Song, Y.-M., Wang, G.-X., Ye, Q., Xiong, R.-G., Akutagawa, T., Nakamura, T., Chan, P. W. H. & Huang, S. P. D. (2007). *J. Am. Chem. Soc.* **129**, 5346–5347.
- Fu, D.-W. & Xiong, R.-G. (2008). *Dalton Trans.* pp. 3946–3948.
- Fu, D.-W., Zhang, W. & Xiong, R.-G. (2008). *Cryst. Growth Des.* **8**, 3461–3464.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Spek, A. L. (2009). *Acta Cryst. D* **65**, 148–155.
- Steiner, T. (2002). *Angew. Chem. Int. Ed.* **41**, 48–76.
- Zimmerman, S. C. & Corbin, P. S. (2000). *Struct. Bond.* **96**, 63–94.

* Thomson Reuters ResearcherID: A-3561-2009.

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Bis(4-fluoroanilinium) sulfate

Hoong-Kun Fun, Suhana Arshad, Dinesha, Sandeep Laxmeshwar and G. K. Nagaraja

S1. Comment

Amine salts have attracted much attention as phase transition dielectric materials for their application in memory storage (Fu *et al.* 2007; Fu & Xiong 2008; Fu *et al.* 2008; Fu *et al.* 2009). Hydrogen bonding is one of the most versatile non-covalent forces in supramolecular chemistry and crystal engineering (Zimmerman & Corbin, 2000; Brunsveld *et al.*, 2001; Desiraju, 2002). Therefore, in the past decades assessment of discrete hydrogen bonding patterns has received great attention (Steiner, 2002; Desiraju & Steiner, 1999; Boutobba *et al.*, 2010) because of their widespread occurrence in biological systems.

The asymmetric unit of the title compound (Fig 1), contains two crystallographically independent 4-fluoroanilinium cations and a sulfate anion. The bond lengths (Allen *et al.*, 1987) and angles are within normal ranges and comparable to those in a closely related crystal structure (Boutobba *et al.*, 2010).

The cations and anions are linked *via* intermolecular N1—H3N1···O3 and N2—H2N2···O3 hydrogen bonds (Table 1). In the crystal packing (Fig. 2), the intermolecular N1—H1N1···O1, N1—H2N1···O3, N1—H2N1···O4, N1—H3N1···O2, N2—H1N2···O4, N2—H3N2···O2 and C11—H11A···O4 hydrogen bonds (Table 1) link the molecules into sheets parallel to the *ab* plane.

S2. Experimental

To a solution of 4-fluoroaniline (10 mmol) in absolute ethanol was added sulfuric acid (5 drops) and the mixture refluxed for 4 h. After cooling the mixture to room temperature, a white solid appeared. This crude product was recrystallized from dimethylformamide to afford the desired product. *M.p.*: 151–153°C.

S3. Refinement

N-bound H atoms were located from a difference Fourier map, fixed at their found location and refined using a riding model with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{N})$ [N—H = 0.8198 to 0.9875 Å]. The remaining H atoms were positioned geometrically [C—H = 0.93 Å] and refined using a riding model with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. The studied crystal is an inversion twin, the refined ratio of twin components being 0.50 (10): 0.50 (10).

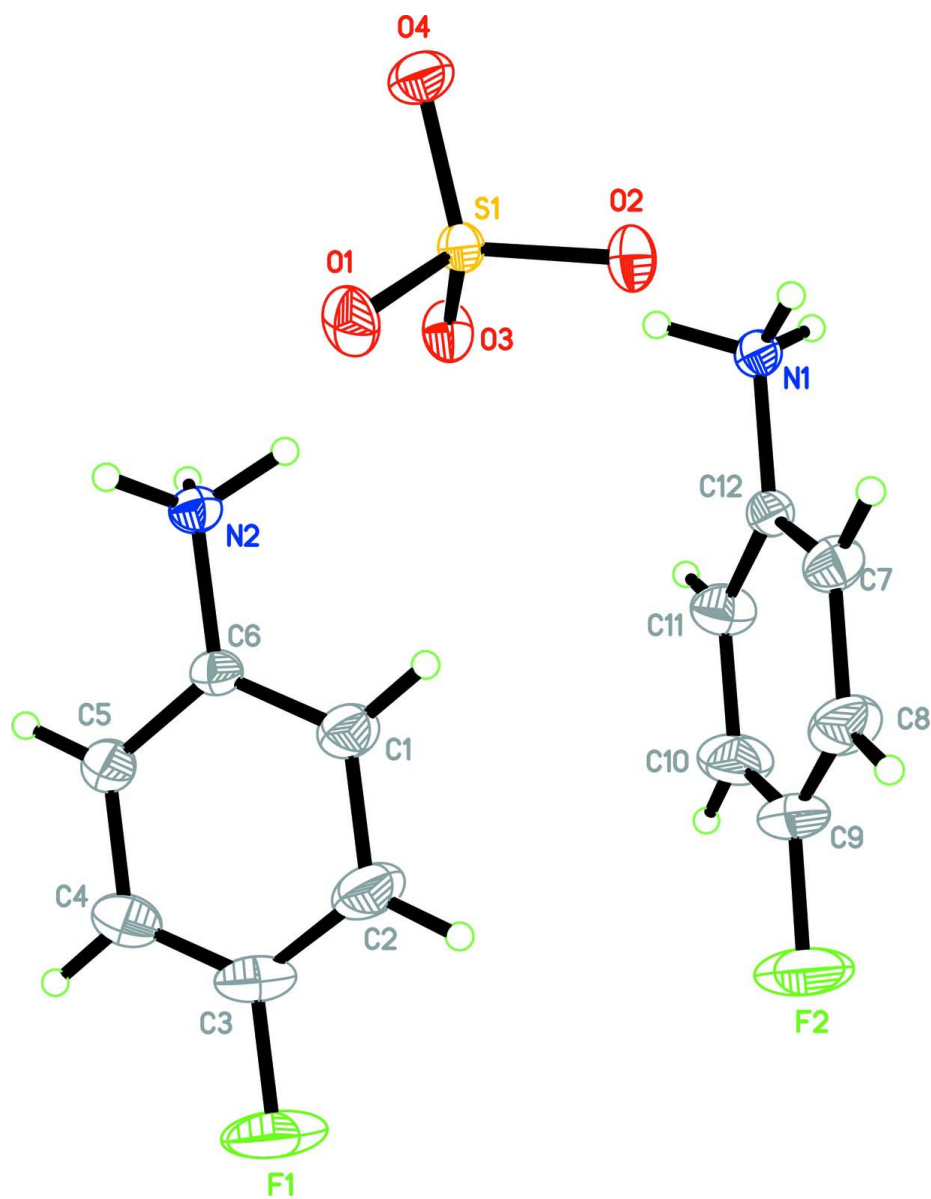
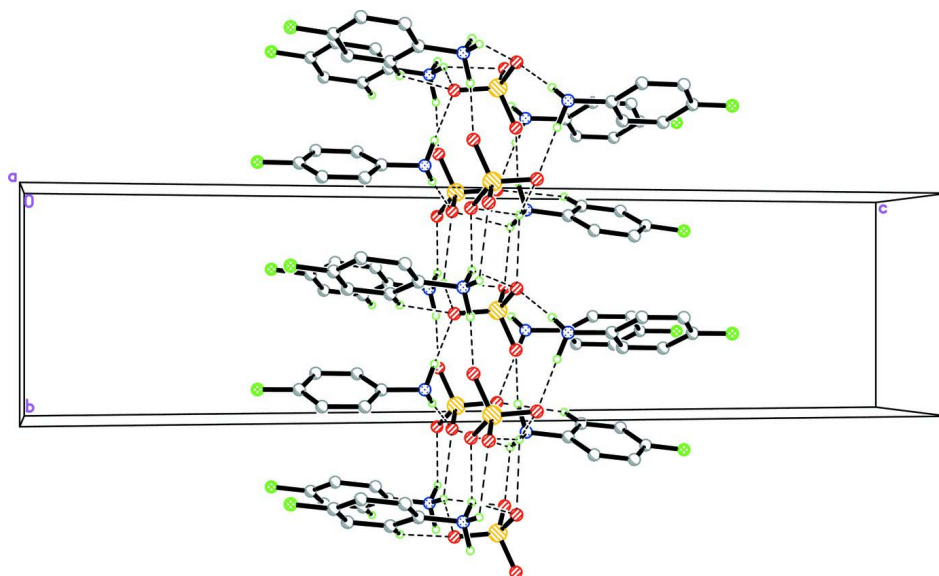


Figure 1

The molecular structure of the title compound with atom labels with 30% probability displacement ellipsoids. Hydrogen atoms are shown as spheres of arbitrary radius.

**Figure 2**

The crystal packing of the title compound. Dashed lines represent hydrogen bonds. H atoms not involved in the hydrogen bond interactions have been omitted for clarity.

Bis(4-fluoroanilinium) sulfate

Crystal data



$M_r = 320.31$

Orthorhombic, $P2_12_12_1$

Hall symbol: P 2ac 2ab

$a = 6.2907 (5) \text{ \AA}$

$b = 7.4155 (6) \text{ \AA}$

$c = 30.168 (3) \text{ \AA}$

$V = 1407.3 (2) \text{ \AA}^3$

$Z = 4$

$F(000) = 664$

$D_x = 1.512 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 5694 reflections

$\theta = 2.7\text{--}27.5^\circ$

$\mu = 0.27 \text{ mm}^{-1}$

$T = 296 \text{ K}$

Needle, colourless

$0.58 \times 0.12 \times 0.07 \text{ mm}$

Data collection

Bruker SMART APEXII DUO CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2009)

$T_{\min} = 0.859$, $T_{\max} = 0.982$

31062 measured reflections

4292 independent reflections

3503 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.044$

$\theta_{\max} = 30.6^\circ$, $\theta_{\min} = 2.7^\circ$

$h = -8 \rightarrow 8$

$k = -10 \rightarrow 10$

$l = -43 \rightarrow 43$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.038$

$wR(F^2) = 0.103$

$S = 0.98$

4292 reflections

191 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
 $w = 1/[\sigma^2(F_o^2) + (0.0492P)^2 + 0.377P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 0.18 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.39 \text{ e } \text{\AA}^{-3}$
 Absolute structure: Flack (1983), 1794 Friedel pairs
 Absolute structure parameter: 0.50 (10)

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	-0.00432 (8)	0.47390 (4)	0.013801 (11)	0.02691 (9)
F1	0.5089 (5)	0.3730 (3)	0.25139 (4)	0.1033 (6)
F2	-0.0181 (4)	0.8372 (3)	0.22704 (4)	0.1096 (7)
O1	-0.0040 (3)	0.29560 (14)	0.03416 (4)	0.0452 (3)
O2	-0.2002 (2)	0.5713 (2)	0.02606 (5)	0.0372 (3)
O3	0.1801 (2)	0.5795 (2)	0.02941 (5)	0.0381 (4)
O4	0.0060 (3)	0.45980 (17)	-0.03476 (4)	0.0470 (3)
N1	-0.0142 (3)	0.92624 (16)	0.04544 (4)	0.0309 (3)
H1N1	0.0125	1.0369	0.0384	0.046*
H2N1	-0.1252	0.8982	0.0330	0.046*
H3N1	0.0935	0.8514	0.0315	0.046*
N2	0.4960 (3)	0.37944 (17)	0.06828 (4)	0.0332 (3)
H1N2	0.5163	0.2639	0.0563	0.050*
H2N2	0.3856	0.4288	0.0573	0.050*
H3N2	0.6038	0.4605	0.0553	0.050*
C1	0.3283 (3)	0.4459 (4)	0.13955 (7)	0.0492 (5)
H1A	0.2115	0.4926	0.1245	0.059*
C2	0.3318 (4)	0.4435 (4)	0.18574 (7)	0.0639 (7)
H2A	0.2179	0.4881	0.2020	0.077*
C3	0.5047 (5)	0.3749 (3)	0.20625 (6)	0.0634 (6)
C4	0.6765 (5)	0.3079 (4)	0.18406 (8)	0.0646 (7)
H4A	0.7923	0.2611	0.1994	0.077*
C5	0.6742 (4)	0.3113 (3)	0.13811 (7)	0.0487 (5)
H5A	0.7901	0.2686	0.1221	0.058*
C6	0.4990 (4)	0.37857 (19)	0.11639 (5)	0.0326 (3)
C7	-0.1933 (3)	0.8307 (3)	0.11348 (6)	0.0434 (5)
H7A	-0.3113	0.7982	0.0967	0.052*
C8	-0.1934 (4)	0.8076 (4)	0.15908 (7)	0.0583 (6)
H8A	-0.3101	0.7575	0.1734	0.070*

C9	-0.0178 (5)	0.8602 (4)	0.18240 (6)	0.0636 (7)
C10	0.1576 (4)	0.9328 (4)	0.16300 (8)	0.0671 (7)
H10A	0.2739	0.9677	0.1800	0.081*
C11	0.1593 (4)	0.9536 (3)	0.11737 (7)	0.0496 (5)
H11A	0.2778	1.0015	0.1032	0.059*
C12	-0.0157 (3)	0.9027 (2)	0.09331 (5)	0.0317 (3)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.02748 (16)	0.02319 (15)	0.03006 (16)	-0.0005 (2)	0.0004 (2)	0.00064 (11)
F1	0.1150 (14)	0.1674 (18)	0.0274 (6)	-0.006 (2)	-0.0039 (9)	0.0034 (8)
F2	0.1171 (15)	0.1814 (19)	0.0301 (6)	0.013 (2)	0.0030 (10)	0.0157 (9)
O1	0.0549 (8)	0.0257 (5)	0.0550 (7)	0.0033 (10)	0.0022 (10)	0.0096 (5)
O2	0.0269 (6)	0.0333 (9)	0.0515 (8)	0.0032 (6)	0.0029 (6)	-0.0008 (7)
O3	0.0295 (6)	0.0324 (9)	0.0524 (8)	-0.0017 (6)	-0.0039 (6)	-0.0042 (7)
O4	0.0639 (8)	0.0467 (7)	0.0303 (6)	0.0010 (14)	0.0026 (8)	-0.0026 (5)
N1	0.0369 (7)	0.0261 (5)	0.0298 (6)	0.0007 (9)	0.0015 (7)	0.0008 (4)
N2	0.0321 (6)	0.0393 (6)	0.0283 (6)	-0.0015 (10)	0.0003 (9)	0.0001 (4)
C1	0.0423 (10)	0.0681 (15)	0.0373 (10)	0.0054 (11)	0.0034 (8)	-0.0018 (10)
C2	0.0595 (14)	0.095 (2)	0.0372 (11)	0.0064 (16)	0.0121 (10)	-0.0068 (13)
C3	0.0767 (16)	0.0857 (16)	0.0279 (8)	-0.008 (2)	-0.0039 (15)	0.0008 (8)
C4	0.0707 (17)	0.0807 (19)	0.0424 (12)	0.0137 (15)	-0.0198 (11)	0.0016 (12)
C5	0.0482 (12)	0.0584 (14)	0.0394 (10)	0.0137 (11)	-0.0061 (9)	-0.0058 (10)
C6	0.0364 (8)	0.0336 (7)	0.0279 (6)	-0.0029 (12)	-0.0022 (10)	-0.0012 (5)
C7	0.0436 (11)	0.0487 (12)	0.0380 (10)	-0.0036 (10)	0.0068 (8)	-0.0001 (9)
C8	0.0661 (16)	0.0679 (16)	0.0409 (11)	-0.0040 (14)	0.0180 (11)	0.0081 (11)
C9	0.0770 (18)	0.0866 (16)	0.0272 (8)	0.010 (2)	0.0032 (13)	0.0061 (9)
C10	0.0642 (15)	0.097 (2)	0.0400 (11)	-0.0012 (16)	-0.0178 (11)	-0.0017 (14)
C11	0.0458 (11)	0.0629 (14)	0.0399 (10)	-0.0086 (11)	-0.0051 (9)	0.0039 (10)
C12	0.0372 (9)	0.0281 (6)	0.0298 (7)	0.0017 (10)	0.0002 (9)	0.0012 (5)

Geometric parameters (Å, °)

S1—O1	1.4579 (11)	C2—C3	1.350 (4)
S1—O4	1.4700 (11)	C2—H2A	0.9300
S1—O2	1.4755 (14)	C3—C4	1.365 (4)
S1—O3	1.4767 (14)	C4—C5	1.387 (3)
F1—C3	1.362 (2)	C4—H4A	0.9300
F2—C9	1.357 (2)	C5—C6	1.376 (3)
N1—C12	1.4548 (19)	C5—H5A	0.9300
N1—H1N1	0.8644	C7—C12	1.379 (3)
N1—H2N1	0.8198	C7—C8	1.386 (3)
N1—H3N1	0.9724	C7—H7A	0.9300
N2—C6	1.4513 (17)	C8—C9	1.366 (4)
N2—H1N2	0.9393	C8—H8A	0.9300
N2—H2N2	0.8524	C9—C10	1.360 (4)
N2—H3N2	0.9875	C10—C11	1.385 (3)

C1—C6	1.376 (3)	C10—H10A	0.9300
C1—C2	1.394 (3)	C11—C12	1.371 (3)
C1—H1A	0.9300	C11—H11A	0.9300
O1—S1—O4	110.81 (7)	C3—C4—C5	118.4 (2)
O1—S1—O2	109.84 (10)	C3—C4—H4A	120.8
O4—S1—O2	108.76 (10)	C5—C4—H4A	120.8
O1—S1—O3	110.21 (10)	C6—C5—C4	119.4 (2)
O4—S1—O3	108.71 (10)	C6—C5—H5A	120.3
O2—S1—O3	108.45 (7)	C4—C5—H5A	120.3
C12—N1—H1N1	111.1	C1—C6—C5	121.03 (16)
C12—N1—H2N1	114.8	C1—C6—N2	119.73 (19)
H1N1—N1—H2N1	107.0	C5—C6—N2	119.24 (19)
C12—N1—H3N1	111.5	C12—C7—C8	119.1 (2)
H1N1—N1—H3N1	107.5	C12—C7—H7A	120.5
H2N1—N1—H3N1	104.5	C8—C7—H7A	120.5
C6—N2—H1N2	112.3	C9—C8—C7	118.4 (2)
C6—N2—H2N2	113.7	C9—C8—H8A	120.8
H1N2—N2—H2N2	110.6	C7—C8—H8A	120.8
C6—N2—H3N2	113.0	F2—C9—C10	118.5 (3)
H1N2—N2—H3N2	108.0	F2—C9—C8	118.3 (3)
H2N2—N2—H3N2	98.3	C10—C9—C8	123.19 (19)
C6—C1—C2	119.4 (2)	C9—C10—C11	118.5 (2)
C6—C1—H1A	120.3	C9—C10—H10A	120.7
C2—C1—H1A	120.3	C11—C10—H10A	120.7
C3—C2—C1	118.4 (2)	C12—C11—C10	119.3 (2)
C3—C2—H2A	120.8	C12—C11—H11A	120.4
C1—C2—H2A	120.8	C10—C11—H11A	120.4
C2—C3—F1	118.5 (3)	C11—C12—C7	121.54 (17)
C2—C3—C4	123.4 (2)	C11—C12—N1	119.15 (19)
F1—C3—C4	118.1 (3)	C7—C12—N1	119.31 (18)
C6—C1—C2—C3	0.2 (4)	C12—C7—C8—C9	1.1 (4)
C1—C2—C3—F1	179.8 (2)	C7—C8—C9—F2	-179.9 (2)
C1—C2—C3—C4	0.1 (5)	C7—C8—C9—C10	-0.5 (4)
C2—C3—C4—C5	0.4 (5)	F2—C9—C10—C11	179.0 (3)
F1—C3—C4—C5	-179.3 (2)	C8—C9—C10—C11	-0.4 (5)
C3—C4—C5—C6	-1.1 (4)	C9—C10—C11—C12	0.8 (4)
C2—C1—C6—C5	-0.9 (4)	C10—C11—C12—C7	-0.2 (4)
C2—C1—C6—N2	179.6 (2)	C10—C11—C12—N1	179.4 (2)
C4—C5—C6—C1	1.4 (4)	C8—C7—C12—C11	-0.8 (3)
C4—C5—C6—N2	-179.1 (2)	C8—C7—C12—N1	179.62 (19)

Hydrogen-bond geometry (\AA , $^\circ$)

<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>
N1—H3N1 \cdots O3	0.97	2.09	2.887 (3)	138
N2—H2N2 \cdots O3	0.85	1.90	2.741 (3)	168

N1—H1M1…O1 ⁱ	0.86	1.92	2.761 (2)	162
N1—H2M1…O3 ⁱⁱ	0.82	2.25	2.967 (2)	146
N1—H2M1…O4 ⁱⁱ	0.82	2.55	3.151 (3)	131
N1—H3M1…O2 ⁱⁱⁱ	0.97	2.24	2.925 (2)	126
N2—H1N2…O4 ^{iv}	0.94	1.78	2.7121 (18)	170
N2—H3N2…O2 ^v	0.99	1.72	2.702 (2)	170
C11—H11A…O4 ⁱⁱⁱ	0.93	2.53	3.374 (3)	151

Symmetry codes: (i) $x, y+1, z$; (ii) $x-1/2, -y+3/2, -z$; (iii) $x+1/2, -y+3/2, -z$; (iv) $x+1/2, -y+1/2, -z$; (v) $x+1, y, z$.