

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

2-Methyl-4-oxo-6,7,8,9-tetrahydrothieno[2',3':4,5]pyrimidino[1,2-a]pyridine-3-carboxylic acid

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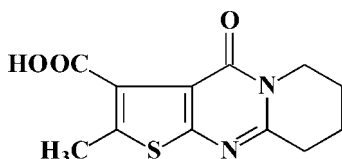
Received 17 January 2011; accepted 2 March 2011

Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.007$ Å; disorder in main residue; R factor = 0.063; wR factor = 0.169; data-to-parameter ratio = 9.9.

There are two independent molecules in the asymmetric unit of the title compound, $\text{C}_{12}\text{H}_{12}\text{N}_2\text{O}_3\text{S}$. With the exception of the methylene groups, a mean plane fitted through all non-H atoms of each molecule has an r.m.s. deviation of 0.035 Å for one molecule and 0.120 Å for the second. In one of the independent molecules, the methylene groups was refined using a disorder model with an occupancy ratio of 0.53:0.47 (14). Each molecule features an intramolecular O—H...O hydrogen bond, which generates an $S(7)$ ring.

Related literature

For the synthesis of thieno[2,3-*d*]pyrimidin-4-ones and their derivatives, see: Litvinov (2004); Elmuradov *et al.* (2010); Csukonyi *et al.* (1986). For the physiological activity of thieno[2,3-*d*]pyrimidin-4-ones and their derivatives, see: Lilienkampf *et al.* (2007). For a related structure, see: Bozorov *et al.* (2010). For hydrogen-bond motifs, see: Bernstein *et al.* (1995).



Experimental

Crystal data

$\text{C}_{12}\text{H}_{12}\text{N}_2\text{O}_3\text{S}$

$M_r = 264.30$

Monoclinic, $P2_1/n$
 $a = 7.2550$ (15) Å
 $b = 20.506$ (4) Å
 $c = 15.824$ (3) Å
 $\beta = 96.93$ (3)°
 $V = 2337.0$ (8) Å³

$Z = 8$
Cu $K\alpha$ radiation
 $\mu = 2.50$ mm⁻¹
 $T = 296$ K
 $0.60 \times 0.40 \times 0.15$ mm

Data collection

Stoe STADI4 diffractometer
Absorption correction: ψ scan
(Blessing, 1987)
 $T_{\min} = 0.346$, $T_{\max} = 0.687$
4297 measured reflections
3438 independent reflections

2420 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.047$
 $\theta_{\text{max}} = 60.0^\circ$
3 standard reflections every 60 min
intensity decay: 4.7%

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.063$
 $wR(F^2) = 0.169$
 $S = 1.10$
3438 reflections
348 parameters
23 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.20$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.28$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O2}-\text{H2O}\cdots\text{O1}$	0.87 (2)	1.63 (2)	2.501 (5)	177 (7)
$\text{O52}-\text{H52O}\cdots\text{O51}$	0.87 (2)	1.71 (3)	2.518 (6)	154 (4)

Data collection: *STADI4* (Stoe & Cie, 1997); cell refinement: *STADI4*; data reduction: *X-RED* (Stoe & Cie, 1997); 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: *publCIF* (Westrip, 2010).

We thank the Academy of Sciences of the Republic of Uzbekistan for supporting this study (grants FA-F3-T045 and FA-F3-T047)

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NK2082).

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supporting information

Acta Cryst. (2011). E67, o824 [doi:10.1107/S1600536811007902]

2-Methyl-4-oxo-6,7,8,9-tetrahydrothieno[2',3':4,5]pyrimidino[1,2-*a*]pyridine-3-carboxylic acid

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S1. Comment

Among heterocyclic compounds, the thieno[2,3-*d*]pyrimidin-4-one family (Litvinov, 2004; Elmuradov *et al.*, 2010; Csukonyi *et al.*, 1986) has been shown to possess various physiological activity (Lilienkamp *et al.*, 2007). The reaction of 2,3-dimethylthieno[2',3':4,5]pyrimidino[1,2-*a*]pyridin-4-one with nitric acid in ratio of reagents - substrate:HNO₃ - 1:4 in concentrated sulfuric acid leads to the formation of 3-hydroxycarbonyl-2-methylthieno[2',3':4,5]pyrimidino[1,2-*a*]pyridin-4-one (Figure 1). We report here the synthesis and crystal structure.

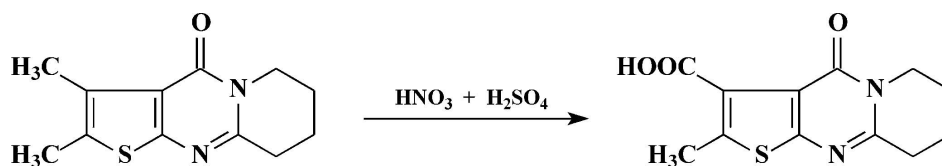
The asymmetric unit contains two crystallographically unique molecules (Figure 2). With the exception of the methylene group, a mean plane fitted through all non-H atoms of each molecule has an rms deviation of 0.035 for one molecule, and 0.120 for the second. In one of the unique molecules the methylene group was refined using a disorder model with an occupancy ratio of 0.53:0.47 (14). An *S*(7) intramolecular O—H...O hydrogen bond is observed in each unique molecule (Bernstein *et al.*, 1995).

S2. Experimental

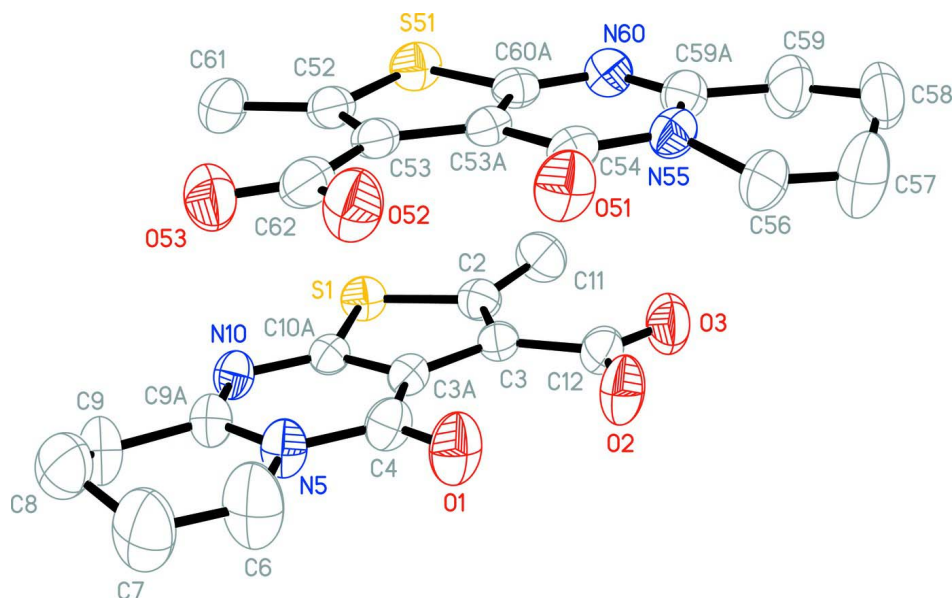
Into a flask supplied with a mixer was poured 2 ml H₂SO₄ and cooled by an ice bath (0.25 h). Then 1 g (4.26 mmole) 2,3-dimethylthieno[2',3':4,5]pyrimidino[1,2-*a*]pyridin-4-one was added in portions, mixed before complete dissolution, nitrating acid consisting of 1.89 g (1.4 ml, *d*=1.35 g/ml) (17.08 mmole) HNO₃ and 1.4 ml (*d*=1.835 g/ml) H₂SO₄ were added (0.5 h) drop wise. A reactionary mixture were mixed 1 h at room temperature and left 2 days at room temperature and a mixture is decomposed in cold. The formed yellow crystals were filtered off and washed with water and dried. Yield 1.0 g (89%), m.p. 478–479 K (ethanol). The yellow crystals suitable for X-ray analysis were obtained from absolute methanol at room temperature.

S3. Refinement

C-bound H atoms were placed geometrically (with C—H distances of 0.97 Å for CH₂; 0.96 Å for CH₃; and 0.93 Å for C_{ar}) and included in the refinement in a riding motion approximation with $U_{\text{iso}}=1.2U_{\text{eq}}(\text{C})$ [$U_{\text{iso}}=1.5U_{\text{eq}}(\text{C})$ for methyl H atoms]. O-bound H atoms involved in the intermolecular hydrogen bonding were found by difference Fourier synthesis and refined isotropically with a distance restrains of 0.85 (2) Å. [O2—H2O = 0.870 (20) Å, O52—H52O = 0.871 (19) Å]. Atoms C56, C57, C58 and C59 were refined using a disorder model with an occupancy ratio of 0.53:0.47 (14)


Figure 1

Reaction scheme.


Figure 2

The asymmetric unit of the title compound, with H atoms and the minor disorder component omitted. Displacement ellipsoids are at the 30% probability level.

2-Methyl-4-oxo-6,7,8,9-tetrahydrothieno[2',3':4,5]pyrimidino[1,2- a]pyridine-3-carboxylic acid

Crystal data

 $C_{12}H_{12}N_2O_3S$
 $M_r = 264.30$

 Monoclinic, $P2_1/n$

 Hall symbol: $-P\ 2_1/n$
 $a = 7.2550\ (15)\ \text{\AA}$
 $b = 20.506\ (4)\ \text{\AA}$
 $c = 15.824\ (3)\ \text{\AA}$
 $\beta = 96.93\ (3)^\circ$
 $V = 2337.0\ (8)\ \text{\AA}^3$
 $Z = 8$
 $F(000) = 1104$
 $D_x = 1.502\ \text{Mg m}^{-3}$

 Cu $K\alpha$ radiation, $\lambda = 1.54184\ \text{\AA}$

Cell parameters from 12 reflections

 $\theta = 10\text{--}20^\circ$
 $\mu = 2.50\ \text{mm}^{-1}$
 $T = 296\ \text{K}$

Prism, yellow

 $0.60 \times 0.40 \times 0.15\ \text{mm}$

Data collection

Stoe STADI4

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

 Scan width (ω) = $1.56\text{--}1.80$, scan ratio $2\theta:\omega =$
 $1.00\ I(\text{Net})$ and $\sigma(I)$ calculated according to

Blessing (1987)

 Absorption correction: ψ scan

(Blessing, 1987)

 $T_{\min} = 0.346$, $T_{\max} = 0.687$

4297 measured reflections

3438 independent reflections

 2420 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.047$

$\theta_{\max} = 60.0^\circ$, $\theta_{\min} = 3.5^\circ$
 $h = -8 \rightarrow 8$
 $k = 0 \rightarrow 23$

$l = 0 \rightarrow 17$
 3 standard reflections every 60 min
 intensity decay: 4.7%

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.063$
 $wR(F^2) = 0.169$
 $S = 1.10$
 3438 reflections
 348 parameters
 23 restraints
 Primary atom site location: structure-invariant
 direct methods
 Secondary atom site location: difference Fourier
 map

Hydrogen site location: inferred from
 neighbouring sites
 H atoms treated by a mixture of independent
 and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0601P)^2 + 2.3488P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.20 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.28 \text{ e } \text{\AA}^{-3}$
 Extinction correction: *SHELXL*,
 $F_c^* = kF_c[1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
 Extinction coefficient: 0.0018 (3)

Special details

Experimental. ψ Scan Reflections used $\mu * R = 0.00$

H K L, θ , χ , I_{\min}/I_{\max} : -1 -1 0 13.0 84.8 0.563

¹H NMR (400 MHz, CDCl₃): 15.46 (1H, s, OH), 4.06 (2H, t, J=6.1 Hz, H-6), 3.01 (2H, t, J=6.6 Hz, H-9), 2.76 (3H, s, CH₃-2), 1.93–2.07 (4H, m, H-7, 8).

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 $F^2 > 2\sigma(F^2)$ 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>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
S1	0.04758 (17)	0.79444 (6)	0.42793 (7)	0.0671 (4)	
O1	0.2734 (6)	0.78157 (17)	0.7344 (2)	0.0980 (12)	
O2	0.3136 (6)	0.66799 (17)	0.6820 (3)	0.0995 (13)	
H2O	0.304 (9)	0.7074 (15)	0.701 (4)	0.119*	
O3	0.2595 (6)	0.61070 (18)	0.5666 (3)	0.1034 (13)	
N5	0.1590 (6)	0.87759 (18)	0.6804 (2)	0.0722 (11)	
N10	0.0491 (5)	0.89415 (17)	0.5367 (2)	0.0619 (10)	
C2	0.1269 (6)	0.7205 (2)	0.4683 (3)	0.0651 (12)	
C3	0.1810 (6)	0.7225 (2)	0.5538 (3)	0.0580 (11)	
C3A	0.1573 (5)	0.7869 (2)	0.5876 (2)	0.0538 (10)	
C4	0.2008 (7)	0.8123 (2)	0.6711 (3)	0.0680 (13)	
C6	0.2045 (12)	0.9053 (3)	0.7664 (3)	0.132 (3)	
H6A	0.1358	0.8810	0.8048	0.159*	
H6B	0.3354	0.8977	0.7841	0.159*	
C7	0.1711 (14)	0.9690 (3)	0.7773 (4)	0.146 (3)	
H7A	0.2672	0.9851	0.8202	0.175*	

H7B	0.0546	0.9722	0.8013	0.175*	
C8	0.1608 (11)	1.0132 (3)	0.7067 (4)	0.114 (2)	
H8A	0.2849	1.0221	0.6927	0.136*	
H8B	0.1070	1.0542	0.7223	0.136*	
C9	0.0451 (9)	0.9853 (2)	0.6305 (4)	0.0967 (18)	
H9A	-0.0848	0.9889	0.6388	0.116*	
H9B	0.0642	1.0110	0.5809	0.116*	
C9A	0.0868 (7)	0.9159 (2)	0.6132 (3)	0.0679 (12)	
C10A	0.0859 (6)	0.8302 (2)	0.5262 (2)	0.0533 (10)	
C11	0.1308 (8)	0.6661 (3)	0.4061 (3)	0.0911 (17)	
H11A	0.2570	0.6528	0.4036	0.137*	
H11B	0.0770	0.6804	0.3509	0.137*	
H11C	0.0610	0.6299	0.4239	0.137*	
C12	0.2532 (7)	0.6630 (3)	0.6007 (4)	0.0745 (14)	
S51	0.54855 (18)	0.78491 (7)	0.40855 (7)	0.0753 (4)	
O51	0.7745 (6)	0.74580 (18)	0.71060 (19)	0.0938 (12)	
O52	0.7039 (7)	0.8660 (2)	0.7032 (3)	0.1067 (14)	
H52O	0.756 (9)	0.8281 (18)	0.715 (4)	0.129*	
O53	0.5648 (7)	0.9368 (2)	0.6164 (3)	0.1136 (15)	
N55	0.7534 (5)	0.65997 (19)	0.6203 (2)	0.0672 (10)	
N60	0.6566 (5)	0.66871 (19)	0.4739 (2)	0.0684 (10)	
C52	0.5490 (6)	0.8492 (2)	0.4774 (3)	0.0683 (13)	
C53A	0.6617 (5)	0.7637 (2)	0.5653 (2)	0.0535 (10)	
C53	0.6134 (6)	0.8322 (2)	0.5592 (3)	0.0615 (11)	
C54	0.7323 (6)	0.7256 (2)	0.6369 (3)	0.0642 (12)	
C56	0.792 (9)	0.6213 (13)	0.7004 (13)	0.089 (7)	0.468 (14)
H56A	0.9111	0.6347	0.7298	0.106*	0.468 (14)
H56B	0.6979	0.6312	0.7370	0.106*	0.468 (14)
C57	0.795 (7)	0.5546 (14)	0.6872 (16)	0.138 (8)	0.468 (14)
H57A	0.6742	0.5381	0.6969	0.166*	0.468 (14)
H57B	0.8840	0.5366	0.7319	0.166*	0.468 (14)
C58	0.838 (3)	0.5262 (9)	0.6076 (14)	0.104 (4)	0.468 (14)
H58A	0.8013	0.4807	0.6053	0.125*	0.468 (14)
H58B	0.9703	0.5285	0.6050	0.125*	0.468 (14)
C59	0.737 (8)	0.5619 (7)	0.5322 (15)	0.107 (3)	0.468 (14)
H59A	0.8012	0.5534	0.4832	0.129*	0.468 (14)
H59B	0.6134	0.5432	0.5203	0.129*	0.468 (14)
C56'	0.832 (7)	0.6170 (11)	0.6917 (12)	0.089 (7)	0.532 (14)
H56C	0.9576	0.6318	0.7108	0.106*	0.532 (14)
H56D	0.7597	0.6230	0.7387	0.106*	0.532 (14)
C57'	0.840 (6)	0.5505 (13)	0.6745 (15)	0.138 (8)	0.532 (14)
H57C	0.8124	0.5275	0.7251	0.166*	0.532 (14)
H57D	0.9668	0.5401	0.6665	0.166*	0.532 (14)
C58'	0.719 (3)	0.5234 (7)	0.6019 (12)	0.104 (4)	0.532 (14)
H58C	0.5906	0.5249	0.6130	0.125*	0.532 (14)
H58D	0.7519	0.4782	0.5932	0.125*	0.532 (14)
C59'	0.744 (7)	0.5631 (6)	0.5233 (13)	0.107 (3)	0.532 (14)
H59C	0.8676	0.5563	0.5077	0.129*	0.532 (14)

H59D	0.6549	0.5490	0.4762	0.129*	0.532 (14)
C59A	0.7158 (7)	0.6339 (2)	0.5402 (3)	0.0706 (13)	
C60A	0.6315 (6)	0.7326 (2)	0.4885 (3)	0.0587 (11)	
C61	0.4852 (9)	0.9139 (3)	0.4415 (4)	0.0991 (19)	
H61A	0.3715	0.9259	0.4628	0.149*	
H61B	0.4649	0.9112	0.3805	0.149*	
H61C	0.5783	0.9462	0.4582	0.149*	
C62	0.6225 (8)	0.8821 (3)	0.6276 (4)	0.0836 (15)	

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0690 (8)	0.0763 (8)	0.0541 (6)	0.0001 (6)	0.0000 (5)	0.0014 (6)
O1	0.138 (3)	0.083 (2)	0.064 (2)	0.010 (2)	-0.025 (2)	0.0097 (18)
O2	0.132 (3)	0.069 (2)	0.091 (3)	0.012 (2)	-0.014 (2)	0.020 (2)
O3	0.135 (4)	0.062 (2)	0.114 (3)	0.012 (2)	0.018 (2)	0.004 (2)
N5	0.094 (3)	0.060 (2)	0.058 (2)	0.001 (2)	-0.009 (2)	-0.0030 (18)
N10	0.065 (2)	0.056 (2)	0.062 (2)	0.0013 (18)	-0.0050 (18)	0.0081 (18)
C2	0.062 (3)	0.065 (3)	0.068 (3)	-0.006 (2)	0.011 (2)	-0.007 (2)
C3	0.051 (3)	0.056 (3)	0.069 (3)	-0.001 (2)	0.012 (2)	0.004 (2)
C3A	0.050 (2)	0.056 (3)	0.054 (2)	-0.004 (2)	0.0020 (18)	0.006 (2)
C4	0.076 (3)	0.066 (3)	0.058 (3)	-0.004 (2)	-0.004 (2)	0.006 (2)
C6	0.230 (9)	0.094 (5)	0.064 (4)	0.010 (5)	-0.022 (4)	-0.022 (3)
C7	0.252 (11)	0.097 (5)	0.089 (5)	-0.004 (6)	0.016 (5)	-0.026 (4)
C8	0.164 (7)	0.083 (4)	0.097 (4)	-0.021 (4)	0.030 (4)	-0.023 (4)
C9	0.115 (5)	0.058 (3)	0.112 (4)	0.006 (3)	-0.008 (4)	-0.001 (3)
C9A	0.072 (3)	0.055 (3)	0.074 (3)	0.002 (2)	-0.003 (2)	0.002 (2)
C10A	0.048 (2)	0.056 (3)	0.055 (2)	-0.003 (2)	0.0022 (18)	0.003 (2)
C11	0.100 (4)	0.091 (4)	0.083 (4)	0.001 (3)	0.010 (3)	-0.018 (3)
C12	0.071 (3)	0.061 (3)	0.093 (4)	-0.004 (3)	0.014 (3)	0.011 (3)
S51	0.0716 (8)	0.0943 (10)	0.0577 (7)	0.0027 (7)	-0.0013 (6)	0.0137 (6)
O51	0.132 (3)	0.094 (3)	0.0509 (19)	-0.001 (2)	-0.0072 (19)	0.0014 (18)
O52	0.162 (4)	0.082 (3)	0.076 (3)	-0.003 (3)	0.016 (3)	-0.009 (2)
O53	0.153 (4)	0.072 (3)	0.121 (3)	0.013 (3)	0.041 (3)	-0.002 (2)
N55	0.072 (3)	0.067 (3)	0.062 (2)	-0.003 (2)	0.0049 (19)	0.0116 (19)
N60	0.075 (3)	0.068 (3)	0.062 (2)	-0.006 (2)	0.0056 (19)	-0.008 (2)
C52	0.058 (3)	0.078 (3)	0.070 (3)	0.002 (2)	0.014 (2)	0.014 (2)
C53A	0.047 (2)	0.062 (3)	0.052 (2)	-0.006 (2)	0.0081 (19)	0.005 (2)
C53	0.055 (3)	0.066 (3)	0.066 (3)	-0.007 (2)	0.019 (2)	0.000 (2)
C54	0.063 (3)	0.073 (3)	0.056 (3)	-0.007 (2)	0.008 (2)	0.004 (2)
C56	0.105 (19)	0.086 (5)	0.074 (5)	0.014 (6)	0.009 (7)	0.031 (4)
C57	0.202 (19)	0.090 (6)	0.116 (9)	0.012 (8)	-0.011 (10)	0.030 (6)
C58	0.107 (10)	0.064 (4)	0.138 (7)	0.019 (9)	0.005 (12)	0.012 (5)
C59	0.147 (7)	0.059 (4)	0.120 (6)	0.002 (4)	0.030 (6)	-0.002 (4)
C56'	0.105 (19)	0.086 (5)	0.074 (5)	0.014 (6)	0.009 (7)	0.031 (4)
C57'	0.202 (19)	0.090 (6)	0.116 (9)	0.012 (8)	-0.011 (10)	0.030 (6)
C58'	0.107 (10)	0.064 (4)	0.138 (7)	0.019 (9)	0.005 (12)	0.012 (5)
C59'	0.147 (7)	0.059 (4)	0.120 (6)	0.002 (4)	0.030 (6)	-0.002 (4)

C59A	0.073 (3)	0.064 (3)	0.075 (3)	-0.004 (2)	0.009 (3)	-0.001 (3)
C60A	0.047 (2)	0.075 (3)	0.054 (2)	-0.003 (2)	0.0043 (19)	0.004 (2)
C61	0.107 (5)	0.085 (4)	0.108 (4)	0.012 (3)	0.024 (3)	0.039 (3)
C62	0.090 (4)	0.080 (4)	0.085 (4)	-0.013 (3)	0.027 (3)	0.005 (3)

Geometric parameters (Å, °)

S1—C10A	1.712 (4)	N55—C54	1.382 (6)
S1—C2	1.717 (5)	N55—C56	1.492 (13)
O1—C4	1.245 (5)	N55—C56'	1.492 (12)
O2—C12	1.313 (6)	N60—C59A	1.299 (6)
O2—H2O	0.87 (2)	N60—C60A	1.346 (6)
O3—C12	1.204 (6)	C52—C53	1.367 (6)
N5—C9A	1.374 (5)	C52—C61	1.495 (6)
N5—C4	1.384 (6)	C53A—C60A	1.368 (5)
N5—C6	1.474 (6)	C53A—C54	1.420 (6)
N10—C9A	1.288 (5)	C53A—C53	1.447 (6)
N10—C10A	1.352 (5)	C53—C62	1.485 (7)
C2—C3	1.363 (6)	C54—O51	1.241 (5)
C2—C11	1.491 (6)	C56—C57	1.385 (13)
C3—C3A	1.444 (6)	C56—H56A	0.9700
C3—C12	1.489 (6)	C56—H56B	0.9700
C3A—C10A	1.370 (5)	C57—C58	1.45 (2)
C3A—C4	1.421 (6)	C57—H57A	0.9700
C4—O1	1.245 (5)	C57—H57B	0.9700
C6—C7	1.342 (7)	C58—C59	1.512 (18)
C6—H6A	0.9700	C58—H58A	0.9700
C6—H6B	0.9700	C58—H58B	0.9700
C7—C8	1.435 (8)	C59—C59A	1.490 (13)
C7—H7A	0.9700	C59—H59A	0.9700
C7—H7B	0.9700	C59—H59B	0.9700
C8—C9	1.498 (7)	C56'—C57'	1.392 (12)
C8—H8A	0.9700	C56'—H56C	0.9700
C8—H8B	0.9700	C56'—H56D	0.9700
C9—C9A	1.487 (6)	C57'—C58'	1.47 (2)
C9—H9A	0.9700	C57'—H57C	0.9700
C9—H9B	0.9700	C57'—H57D	0.9700
C11—H11A	0.9600	C58'—C59'	1.516 (18)
C11—H11B	0.9600	C58'—H58C	0.9700
C11—H11C	0.9600	C58'—H58D	0.9700
S51—C52	1.709 (5)	C59'—C59A	1.494 (12)
S51—C60A	1.711 (4)	C59'—H59C	0.9700
O51—C54	1.241 (5)	C59'—H59D	0.9700
O52—C62	1.312 (7)	C61—H61A	0.9600
O52—H52O	0.871 (19)	C61—H61B	0.9600
O53—C62	1.202 (6)	C61—H61C	0.9600
N55—C59A	1.372 (6)		

C10A—S1—C2	91.7 (2)	C52—C53—C53A	111.3 (4)
C12—O2—H2O	113 (4)	C52—C53—C62	119.6 (5)
C9A—N5—C4	122.6 (4)	C53A—C53—C62	129.1 (4)
C9A—N5—C6	121.0 (4)	O51—C54—N55	118.9 (4)
C4—N5—C6	116.4 (4)	O51—C54—N55	118.9 (4)
C9A—N10—C10A	115.4 (4)	O51—C54—C53A	126.1 (5)
C3—C2—C11	130.6 (5)	O51—C54—C53A	126.1 (5)
C3—C2—S1	112.8 (3)	N55—C54—C53A	115.0 (4)
C11—C2—S1	116.6 (4)	C57—C56—N55	114 (2)
C2—C3—C3A	111.1 (4)	C57—C56—H56A	108.8
C2—C3—C12	120.8 (4)	N55—C56—H56A	108.8
C3A—C3—C12	128.0 (4)	C57—C56—H56B	108.8
C10A—C3A—C4	116.4 (4)	N55—C56—H56B	108.8
C10A—C3A—C3	112.5 (4)	H56A—C56—H56B	107.7
C4—C3A—C3	131.0 (4)	C56—C57—C58	122 (3)
O1—C4—N5	118.8 (4)	C56—C57—H57A	106.8
O1—C4—N5	118.8 (4)	C58—C57—H57A	106.8
O1—C4—C3A	126.0 (4)	C56—C57—H57B	106.8
O1—C4—C3A	126.0 (4)	C58—C57—H57B	106.8
N5—C4—C3A	115.2 (4)	H57A—C57—H57B	106.6
C7—C6—N5	118.0 (6)	C57—C58—C59	110.9 (19)
C7—C6—H6A	107.8	C57—C58—H58A	109.5
N5—C6—H6A	107.8	C59—C58—H58A	109.5
C7—C6—H6B	107.8	C57—C58—H58B	109.5
N5—C6—H6B	107.8	C59—C58—H58B	109.5
H6A—C6—H6B	107.1	H58A—C58—H58B	108.0
C6—C7—C8	120.6 (6)	C59A—C59—C58	117.1 (17)
C6—C7—H7A	107.2	C59A—C59—H59A	108.0
C8—C7—H7A	107.2	C58—C59—H59A	108.0
C6—C7—H7B	107.2	C59A—C59—H59B	108.0
C8—C7—H7B	107.2	C58—C59—H59B	108.0
H7A—C7—H7B	106.8	H59A—C59—H59B	107.3
C7—C8—C9	110.9 (5)	C57'—C56'—N55	116.8 (18)
C7—C8—H8A	109.5	C57'—C56'—H56C	108.1
C9—C8—H8A	109.5	N55—C56'—H56C	108.1
C7—C8—H8B	109.5	C57'—C56'—H56D	108.1
C9—C8—H8B	109.5	N55—C56'—H56D	108.1
H8A—C8—H8B	108.1	H56C—C56'—H56D	107.3
C9A—C9—C8	114.1 (5)	C56'—C57'—C58'	119 (2)
C9A—C9—H9A	108.7	C56'—C57'—H57C	107.4
C8—C9—H9A	108.7	C58'—C57'—H57C	107.4
C9A—C9—H9B	108.7	C56'—C57'—H57D	107.4
C8—C9—H9B	108.7	C58'—C57'—H57D	107.5
H9A—C9—H9B	107.6	H57C—C57'—H57D	107.0
N10—C9A—N5	123.2 (4)	C57'—C58'—C59'	108.7 (19)
N10—C9A—C9	118.6 (4)	C57'—C58'—H58C	110.0
N5—C9A—C9	118.2 (4)	C59'—C58'—H58C	110.0
N10—C10A—C3A	127.2 (4)	C57'—C58'—H58D	110.0

N10—C10A—S1	121.0 (3)	C59'—C58'—H58D	110.0
C3A—C10A—S1	111.8 (3)	H58C—C58'—H58D	108.3
C2—C11—H11A	109.5	C59A—C59'—C58'	110.1 (15)
C2—C11—H11B	109.5	C59A—C59'—H59C	109.6
H11A—C11—H11B	109.5	C58'—C59'—H59C	109.6
C2—C11—H11C	109.5	C59A—C59'—H59D	109.6
H11A—C11—H11C	109.5	C58'—C59'—H59D	109.6
H11B—C11—H11C	109.5	H59C—C59'—H59D	108.2
O3—C12—O2	118.8 (5)	N60—C59A—N55	122.8 (4)
O3—C12—C3	122.5 (5)	N60—C59A—C59	120.2 (10)
O2—C12—C3	118.7 (5)	N55—C59A—C59	117.0 (10)
C52—S51—C60A	92.2 (2)	N60—C59A—C59'	115.4 (9)
C62—O52—H52O	124 (3)	N55—C59A—C59'	121.9 (9)
C59A—N55—C54	122.7 (4)	N60—C60A—C53A	126.6 (4)
C59A—N55—C56	124.9 (12)	N60—C60A—S51	121.8 (3)
C54—N55—C56	111.7 (12)	C53A—C60A—S51	111.6 (3)
C59A—N55—C56'	119.0 (10)	C52—C61—H61A	109.5
C54—N55—C56'	118.2 (10)	C52—C61—H61B	109.5
C59A—N60—C60A	115.8 (4)	H61A—C61—H61B	109.5
C53—C52—C61	129.8 (5)	C52—C61—H61C	109.5
C53—C52—S51	112.4 (4)	H61A—C61—H61C	109.5
C61—C52—S51	117.8 (4)	H61B—C61—H61C	109.5
C60A—C53A—C54	117.1 (4)	O53—C62—O52	118.6 (6)
C60A—C53A—C53	112.4 (4)	O53—C62—C53	123.5 (6)
C54—C53A—C53	130.5 (4)	O52—C62—C53	117.9 (5)
C10A—S1—C2—C3	0.1 (4)	C59A—N55—C54—O51	-177.8 (4)
C10A—S1—C2—C11	-178.4 (4)	C56—N55—C54—O51	11 (3)
C11—C2—C3—C3A	178.0 (5)	C56'—N55—C54—O51	-2 (2)
S1—C2—C3—C3A	-0.2 (5)	C59A—N55—C54—O51	-177.8 (4)
C11—C2—C3—C12	-2.1 (8)	C56—N55—C54—O51	11 (3)
S1—C2—C3—C12	179.7 (3)	C56'—N55—C54—O51	-2 (2)
C2—C3—C3A—C10A	0.3 (5)	C59A—N55—C54—C53A	2.4 (6)
C12—C3—C3A—C10A	-179.6 (4)	C56—N55—C54—C53A	-169 (3)
C2—C3—C3A—C4	-177.3 (4)	C56'—N55—C54—C53A	178 (2)
C12—C3—C3A—C4	2.8 (8)	C60A—C53A—C54—O51	177.1 (5)
C9A—N5—C4—O1	177.0 (5)	C53—C53A—C54—O51	-3.2 (8)
C6—N5—C4—O1	-0.2 (8)	C60A—C53A—C54—O51	177.1 (5)
C9A—N5—C4—O1	177.0 (5)	C53—C53A—C54—O51	-3.2 (8)
C6—N5—C4—O1	-0.2 (8)	C60A—C53A—C54—N55	-3.1 (6)
C9A—N5—C4—C3A	-2.2 (7)	C53—C53A—C54—N55	176.6 (4)
C6—N5—C4—C3A	-179.5 (5)	C59A—N55—C56—C57	3 (6)
C10A—C3A—C4—O1	-176.9 (5)	C54—N55—C56—C57	174 (4)
C3—C3A—C4—O1	0.6 (8)	C56'—N55—C56—C57	-65 (10)
C10A—C3A—C4—O1	-176.9 (5)	N55—C56—C57—C58	25 (7)
C3—C3A—C4—O1	0.6 (8)	C56—C57—C58—C59	-44 (6)
C10A—C3A—C4—N5	2.2 (6)	C57—C58—C59—C59A	37 (5)
C3—C3A—C4—N5	179.7 (4)	C59A—N55—C56'—C57'	-9 (5)

C9A—N5—C6—C7	0.7 (11)	C54—N55—C56'—C57'	176 (3)
C4—N5—C6—C7	178.0 (8)	C56—N55—C56'—C57'	111 (15)
N5—C6—C7—C8	-23.3 (14)	N55—C56'—C57'—C58'	-20 (6)
C6—C7—C8—C9	45.4 (12)	C56'—C57'—C58'—C59'	52 (4)
C7—C8—C9—C9A	-45.3 (8)	C57'—C58'—C59'—C59A	-53 (4)
C10A—N10—C9A—N5	-0.1 (7)	C60A—N60—C59A—N55	-0.8 (7)
C10A—N10—C9A—C9	-178.2 (5)	C60A—N60—C59A—C59	176 (3)
C4—N5—C9A—N10	1.2 (8)	C60A—N60—C59A—C59'	-180 (2)
C6—N5—C9A—N10	178.3 (6)	C54—N55—C59A—N60	-0.4 (7)
C4—N5—C9A—C9	179.4 (5)	C56—N55—C59A—N60	170 (3)
C6—N5—C9A—C9	-3.6 (8)	C56'—N55—C59A—N60	-176 (2)
C8—C9—C9A—N10	-155.3 (5)	C54—N55—C59A—C59	-177 (3)
C8—C9—C9A—N5	26.4 (8)	C56—N55—C59A—C59	-7 (4)
C9A—N10—C10A—C3A	0.2 (7)	C56'—N55—C59A—C59	7 (4)
C9A—N10—C10A—S1	-178.8 (3)	C54—N55—C59A—C59'	179 (2)
C4—C3A—C10A—N10	-1.3 (7)	C56—N55—C59A—C59'	-11 (4)
C3—C3A—C10A—N10	-179.3 (4)	C56'—N55—C59A—C59'	3 (3)
C4—C3A—C10A—S1	177.7 (3)	C58—C59—C59A—N60	169 (2)
C3—C3A—C10A—S1	-0.2 (5)	C58—C59—C59A—N55	-14 (5)
C2—S1—C10A—N10	179.2 (4)	C58—C59—C59A—C59'	131 (38)
C2—S1—C10A—C3A	0.1 (3)	C58'—C59'—C59A—N60	-152.3 (19)
C2—C3—C12—O3	-3.2 (8)	C58'—C59'—C59A—N55	29 (4)
C3A—C3—C12—O3	176.6 (5)	C58'—C59'—C59A—C59	-9 (33)
C2—C3—C12—O2	176.1 (5)	C59A—N60—C60A—C53A	-0.1 (7)
C3A—C3—C12—O2	-4.0 (7)	C59A—N60—C60A—S51	-178.3 (4)
C60A—S51—C52—C53	1.0 (4)	C54—C53A—C60A—N60	2.2 (7)
C60A—S51—C52—C61	179.9 (4)	C53—C53A—C60A—N60	-177.5 (4)
C61—C52—C53—C53A	-179.5 (5)	C54—C53A—C60A—S51	-179.5 (3)
S51—C52—C53—C53A	-0.7 (5)	C53—C53A—C60A—S51	0.8 (5)
C61—C52—C53—C62	1.5 (8)	C52—S51—C60A—N60	177.4 (4)
S51—C52—C53—C62	-179.7 (4)	C52—S51—C60A—C53A	-1.0 (3)
C60A—C53A—C53—C52	0.0 (5)	C52—C53—C62—O53	4.9 (8)
C54—C53A—C53—C52	-179.7 (4)	C53A—C53—C62—O53	-173.9 (5)
C60A—C53A—C53—C62	178.8 (5)	C52—C53—C62—O52	-172.6 (5)
C54—C53A—C53—C62	-0.8 (8)	C53A—C53—C62—O52	8.6 (8)

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

<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>
O2—H2O...O1	0.87 (2)	1.63 (2)	2.501 (5)	177 (7)
O52—H52O...O51	0.87 (2)	1.71 (3)	2.518 (6)	154 (4)