

Received 15 October 2014
Accepted 22 October 2014

Edited by A. J. Lough, University of Toronto,
Canada

Keywords: crystal structure; nicotine; proton-transfer salts; 3,5-dinitrosalicylic acid; 5-sulfosalicylic acid; hydrogen-bonding; π - π interactions

CCDC references: 1030394; 1030395

Supporting information: this article has supporting information at journals.iucr.org/e

Crystal structures and hydrogen bonding in the proton-transfer salts of nicotine with 3,5-dinitrosalicylic acid and 5-sulfosalicylic acid

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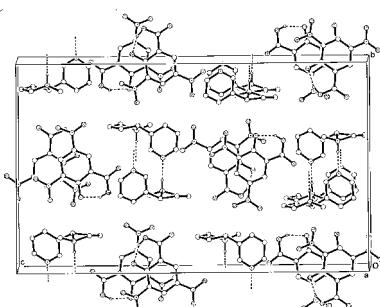
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The structures of the 1:1 anhydrous salts of nicotine (NIC) with 3,5-dinitrosalicylic acid (DNSA) and 5-sulfosalicylic acid (5-SSA), namely (*1R,2S*)-1-methyl-2-(pyridin-3-yl)-1*H*-pyrrolidin-1-i um 2-carboxy-4,6-dinitrophenolate, $C_{10}H_{15}N_2^+ \cdot C_7H_3N_2O_7^-$, (I), and (*1R,2S*)-1-methyl-2-(pyridin-3-yl)-1*H*-pyrrolidin-1-i um 3-carboxy-4-hydroxybenzenesulfonate, $C_{10}H_{15}N_2^+ \cdot C_7H_5O_6S^-$, (II), are reported. The asymmetric units of both (I) and (II) comprise two independent nicotinium cations (*C* and *D*) and either two DNSA or two 5-SSA anions (*A* and *B*), respectively. One of the DNSA anions shows a 25% rotational disorder in the benzene ring system. In the crystal of (I), inter-unit pyrrolidinium N—H \cdots N_{pyridine} hydrogen bonds generate zigzag NIC cation chains which extend along *a*, while the DNSA anions are not involved in any formal inter-species hydrogen bonding but instead form π - π -associated stacks which are parallel to the NIC cation chains along *a* [ring-centroid separation = 3.857 (2) Å]. Weak C—H \cdots O interactions between chain substructures give an overall three-dimensional structure. In the crystal of (II), *A* and *B* anions form independent zigzag chains with *C* and *D* cations, respectively, through carboxylic acid O—H \cdots N_{pyridine} hydrogen bonds. These chains, which extend along *b*, are pseudocentrosymmetrically related and give π - π interactions between the benzene rings of anions *A* and *B* and the pyridine rings of the NIC cations *C* and *D*, respectively [ring centroid separations = 3.6422 (19) and 3.7117 (19) Å]. Also present are weak C—H \cdots O hydrogen-bonding interactions between the chains, giving an overall three-dimensional structure.

1. Chemical context

Nicotine [3-(2S-1-methylpyrrolidin-2-yl)pyridine] is well known as a toxic liquid alkaloid which is found in the leaves of the tobacco plants *Nicotiana tabacum* and *N. rustica* (Rodgman & Parfetti, 2009). Because of these properties, nicotine and its compounds have been of commercial interest and have been used in the past as insecticides and as veterinary ectoparasiticides (usually as the sulfate) (Ujváry, 1999), as well as in limited medical applications as the bitartrate (Eudermol) for the treatment of smoking-withdrawal syndrome (Enzell *et al.*, 1977). However, its veterinary use is restricted due to its toxicity with even topical applications, resulting in the total ban on its use in the USA early in 2014.

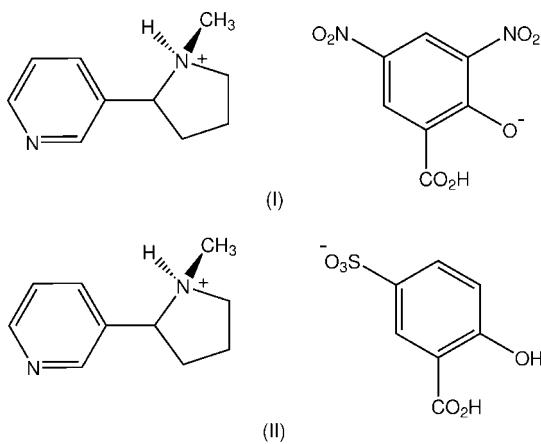
As a Lewis base, nicotine is potentially capable of forming both monocationic and dicationic species ($pK_{a1} = 3.10$ and $pK_{a2} = 8.01$) and the sulfate, dihydrochloride, bitartrate and bipicrate salts have been reported (O'Neil, 2001). However, the only example of a simple dicationic salt in the crystallographic literature is the dihydroiodide (Koo & Kim, 1965). Some metal complexes with the dication as a counter-ion are known, *e.g.* tetrachloridocopper(II) nicotinate (Choi *et al.*,



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2002). More commonly, monocationic salt structures are reported, e.g. the iodide (Barlow *et al.*, 1986), the picrate (Arnaud *et al.*, 2007) and the salicylate (Kim & Jeffrey, 1971).

3,5-Dinitrosalicylic acid (DNSA) ($pK_a = 2.18$) and 3-carboxy-4-hydroxybenzenesulfonic acid (5-sulfosalicylic acid: 5-SSA) ($pK_a < 1$) are capable of forming salts with most Lewis bases and have been used for the formation of crystalline salts suitable for X-ray analysis, e.g. with 5-SSA (Baskar Raj *et al.*, 2003; Smith *et al.*, 2006) and with DNSA, where the majority of the salts formed are phenolates rather than carboxylates (Smith *et al.*, 2007). The title salts $C_{10}H_{15}N_2^+ \cdot C_7H_3N_2O_7^-$, (I) and $C_{10}H_{15}N_2^+ \cdot C_7H_5O_6S^-$, (II) were prepared from the reaction of nicotine (NIC) with DNSA and with 5-SSA, respectively, and the structures are reported herein.



2. Structural commentary

In both the nicotinium salts of DNSA (I) and 5-SSA (II), proton-transfer to the pyrrolidine N-atom of nicotine has occurred as expected, generating an N11(R) chiral centre relative to the known C21(S) centre. Also, in both (I) and (II) (Figs. 1 and 2), the asymmetric units comprise two indepen-

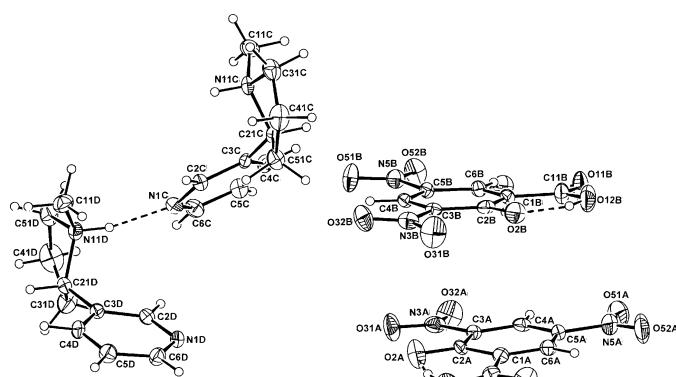


Figure 1

The molecular conformation and atom labelling for the two NIC cations (C and D) and the two DNSA anions (A and B) in the asymmetric unit of (I), with displacement ellipsoids drawn at the 40% probability level. Inter-species hydrogen bonds are shown as dashed lines (see Table 1).

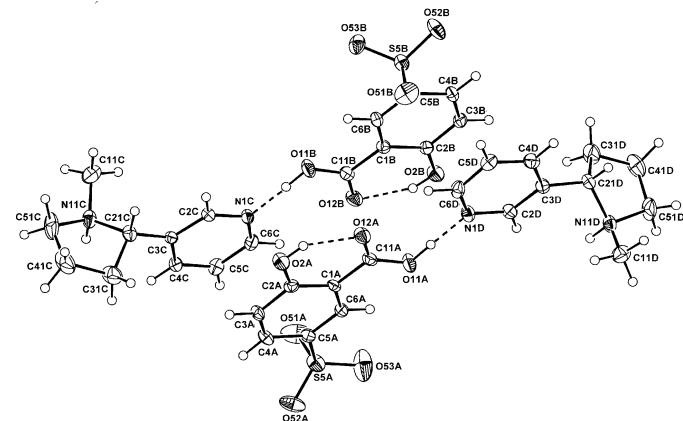


Figure 2

The molecular conformation and atom labelling for the two NIC cations (C and D) and the two 5-SSA anions (A and B) in the asymmetric unit of (II), with displacement ellipsoids drawn at the 40% probability level. Inter-species hydrogen bonds are shown as dashed lines (see Table 2).

dent NIC⁺ cations (C and D) and either, for (I), two DNSA phenolate monoanions or two 5-SSA carboxylate monoanions (A and B) (Figs. 1, 2). With (II), the two independent anion and cation pairs are pseudo-centrosymmetrically related but the presence of the inversion centre is obviated by the fact that both of the NIC cations have the same N11(R), C21(S) absolute configuration.

In (I), the nicotinium C and D cations are conformationally similar but in (II), they are different. However, in both, the pyrrolidinium plane is significantly rotated with respect to that of the benzene ring [the torsion angles C2C/D—C3C/D—C21C/D—N11C/D are $-71.9(4)$ (C) and $-68.8(4)$ ° (D) in (I) and $-45.7(4)$ (C) and $125.7(3)$ ° (D) in (II)]. This conformation with the two rings *antiplanar* is usual for cationic nicotine structures, e.g. Arnaud *et al.* (2007). The substituent carboxyl and nitro groups of the DNSA anions in (I) are essentially coplanar with the benzene ring, with the maximum deviation among the three defining torsion angles for each anion (C2A/B—C1A/B—C11A/B—O2A/B, C2A/B—C3A/B—N3A/B—O32A/B and C4A/B—C5A/B—N5A/B—O52A/B) being for the C3B nitro group [$173.7(3)$ °]. In the B anion, there is 25% rotational disorder about the C1···C4 ring vector, which generates a second phenolic O-component (O21B). This phenomenon has precedence in DNSA salt structures, e.g. with the nicotinamide salt (Koman *et al.*, 2003; 24% disorder). The C3 nitro group is most often associated with deviation from planarity in the DNSA phenolate salts (Smith *et al.*, 2007) and is the more interactive and sterically crowded group. In the case of (I), the uncommon planarity is probably associated with the presence of anion π-bonding associations.

With the 5-SSA anions, the carboxylic acid group is essentially coplanar with the benzene ring, which is expected in this salicylic acid species, invariably having the short intramolecular carboxylic acid O—H···O_{phenol} hydrogen bond (Table 2) (Smith *et al.*, 2006). This interaction is also present in the phenolate anion in (I) in which the carboxylic acid H-atom is *anti*-related (Table 1).

Table 1
Hydrogen-bond geometry (\AA , $^\circ$) for (I).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O12A—H12A \cdots O2A	0.84	1.71	2.475 (4)	150
O12B—H12B \cdots O2B	0.84	1.63	2.411 (4)	152
N11C—H11C \cdots N1D ⁱ	0.93	1.89	2.809 (4)	169
N11D—H11D \cdots N1C	0.93	1.90	2.817 (5)	168
C2C—H2C \cdots O11A ⁱⁱ	0.95	2.42	3.228 (5)	143
C4C—H4C \cdots O31A ⁱ	0.95	2.59	3.452 (5)	151
C6C—H6C \cdots O32A ⁱⁱⁱ	0.95	2.27	3.054 (5)	139
C11C—H13C \cdots O32B ⁱ	0.98	2.48	3.151 (6)	126
C11D—H14D \cdots O51A ^{iv}	0.98	2.55	3.373 (6)	141
C21C—H21C \cdots O2A ⁱ	1.00	2.27	3.163 (5)	148
C21D—H21D \cdots O11B ^v	1.00	2.44	3.307 (5)	144
C51C—H52C \cdots O11A ⁱⁱ	0.99	2.54	3.534 (6)	177

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

Table 2
Hydrogen-bond geometry (\AA , $^\circ$) for (II).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O2A—H2A \cdots O12A	0.84	1.80	2.549 (4)	147
O2B—H2B \cdots O12B	0.84	1.82	2.561 (4)	146
O11A—H11A \cdots N1D	0.95	1.60	2.555 (4)	179
O11B—H11B \cdots N1C	0.95	1.61	2.558 (4)	179
N11C—H11C \cdots O51B ⁱ	0.93	2.32	3.022 (5)	132
N11C—H11C \cdots O53B ⁱ	0.93	2.15	3.029 (5)	157
N11D—H11D \cdots O52A ⁱⁱ	0.93	1.85	2.735 (4)	158
C11D—H12D \cdots O2B ⁱⁱⁱ	0.98	2.51	3.491 (5)	174
C2C—H2C \cdots O53B ⁱ	0.95	2.29	3.201 (5)	160
C2D—H2D \cdots O53A ^{iv}	0.95	2.45	3.359 (4)	160
C11C—H12C \cdots O2A ^v	0.98	2.52	3.481 (5)	165
C11C—H13C \cdots O52B ^{vi}	0.98	2.46	3.290 (5)	142
C11D—H13D \cdots O51A ^{iv}	0.98	2.37	3.251 (5)	150
C21C—H21C \cdots O52B ^{vi}	1.00	2.42	3.331 (5)	151

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

3. Supramolecular features

In the supramolecular structure of (I), the two independent NIC cations *C* and *D* interact through N1C—H \cdots N1Dⁱ and N1D—H \cdots N1C hydrogen bonds (Table 1), giving zigzag chains extending along *a* (Fig. 3). With the DNAA anions, there are no formal hydrogen-bonding interactions either between *A* and *B* anions or with the NIC chain structures. Instead, these anions form π -bonded stacks which are parallel to the NIC⁺ chains down *a* [ring-centroid separation = 3.857 (2) \AA]. The presence of π - π stacking is unusual in DNAA cation structures. In the crystal, there are a number of intermolecular CC/D—H \cdots OA/B hydrogen-bonding interactions, which give an overall three-dimensional structure.

In the crystal of (II), the independent *A* and *B* 5-SSA anions form carboxylic acid O—H \cdots N_{pyridine} hydrogen bonds with the *D* and *C* NIC cations, respectively (Table 2) (see Fig. 2). These cation–anion subunits are then extended into independent chain structures through pyrrolidinium N—H \cdots O_{sulfonate} hydrogen bonds, which with anion *C* is three-centre (O51Bⁱ and O53Bⁱ) and with anion *D*, linear (O52Aⁱⁱ). These give independent zigzag chain substructures which extend along *b*. Although there are no formal hydrogen-

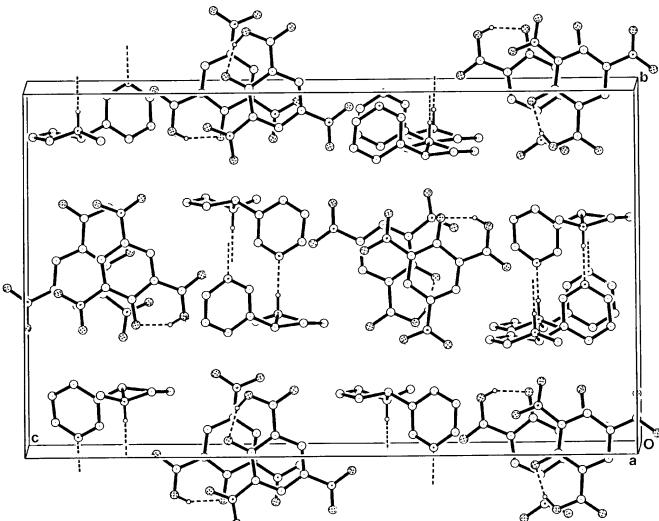


Figure 3

The alternating hydrogen-bonded C-D cationic columns and π -bonded A-B anion stacks in the structure of (I), viewed along the stacks in the unit cell.

bonding links between the two chains, there are π - π interactions between 5-SSA anion *A* and *B* benzene rings and *C* and *D* NIC cation pyridine rings, respectively [ring-centroid separations = 3.6422 (19) and 3.7117 (19) \AA] (Fig. 4). The presence of a number of intermolecular C—H \cdots O hydrogen-bonding interactions to carboxyl, nitro and phenolic O-atom acceptors gives rise to an overall three-dimensional structure.

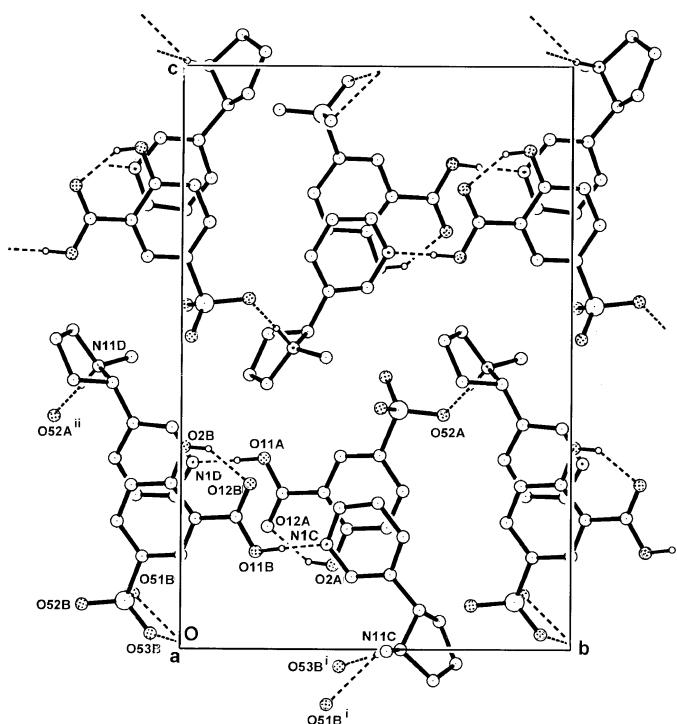


Figure 4

The hydrogen-bonded A-C and B-D chain structures in (II), extending along *b*. Non-associative H atoms have been omitted. For symmetry codes, see Table 2.

Table 3

Experimental details.

	(I)	(II)
Crystal data		
Chemical formula	$C_{10}H_{15}N_2^+ \cdot C_7H_3N_2O_7^-$	$C_{10}H_{15}N_2^+ \cdot C_7H_5O_6S^-$
M_r	390.35	380.41
Crystal system, space group	Orthorhombic, $P2_12_12_1$	Monoclinic, $P2_1$
Temperature (K)	200	200
a, b, c (Å)	6.8096 (5), 17.6403 (15), 29.3604 (19)	7.1568 (3), 12.6416 (5), 19.1519 (8)
α, β, γ (°)	90, 90, 90	90, 93.729 (4), 90
V (Å ³)	3526.9 (4)	1729.07 (12)
Z	8	4
Radiation type	Mo $K\alpha$	Mo $K\alpha$
μ (mm ⁻¹)	0.12	0.23
Crystal size (mm)	0.40 × 0.10 × 0.08	0.35 × 0.30 × 0.12
Data collection		
Diffractometer	Oxford Diffraction Gemini-S CCD detector	Oxford Diffraction Gemini-S CCD detector
Absorption correction	Multi-scan (<i>CrysAlis PRO</i> ; Agilent, 2013)	Multi-scan (<i>CrysAlis PRO</i> ; Agilent, 2013)
T_{min}, T_{max}	0.807, 0.980	0.909, 0.981
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	8840, 6476, 4303	7764, 5104, 4424
R_{int}	0.028	0.031
(sin θ/λ) _{max} (Å ⁻¹)	0.617	0.680
Refinement		
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.072, 0.122, 1.07	0.046, 0.108, 1.01
No. of reflections	6476	5104
No. of parameters	508	469
No. of restraints	2	1
H-atom treatment	H-atom parameters constrained	H-atom parameters constrained
$\Delta\rho_{\text{max}}, \Delta\rho_{\text{min}}$ (e Å ⁻³)	0.44, -0.19	0.49, -0.36
Absolute structure	Flack (1983), 2983 Friedel pairs	Flack (1983), 4361 Friedel pairs
Absolute structure parameter	-0.2 (16)	-0.02 (9)

Computer programs: *CrysAlis PRO* (Agilent, 2013), *SHELXS97* and *SHELXL97* (Sheldrick, 2008) within *WinGX* (Farrugia, 2012), *SIR92* (Altomare *et al.*, 1993) and *PLATON* (Spek, 2009).

4. Synthesis and crystallization

The title salts (I) and (II) were prepared by refluxing equimolar quantities of nicotine (160 mg) and the respective acids, 3,5-dinitrosalicylic acid (230 mg) for (I) or 3-carboxy-4-hydroxybenzenesulfonic acid (220 mg) for (II) in 30 ml of ethanol for 10 min, after which room temperature evaporation of the solutions gave, for (I), thin yellow needles and for (II) colourless prisms, from which specimens were cleaved for the X-ray analyses.

5. Refinement details

Crystal data, data collection and structure refinement details are summarized in Table 3. H atoms on all potentially interactive O—H and N—H groups in all molecular species, were located by difference-Fourier methods but these and the carbon-bound H-atoms were subsequently set as riding on the parent atoms in the refinement in calculated positions [O—H = 0.88, N—H = 0.94, C—H = 0.95–1.00 Å] and with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O}$ or methyl-C) or $1.2U_{\text{eq}}(\text{C}, \text{N})$.

The site occupancy factors for the rotationally disordered phenolate components (O2B) and its other component (O21B) in anion B of (I) were determined as 0.752 (4): 0.248 (4) and were subsequently set at 0.75:0.25 in the refinement.

In both structures, the known C21(S) absolute configuration was invoked. The Flack parameter for (I) [0.2 (16)] has no physical meaning. The absolute structure of compound (II) was confirmed by resonant scattering [Flack parameter = -0.02 (9)].

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supporting information

Acta Cryst. (2014). E70, 430-434 [doi:10.1107/S1600536814023253]

Crystal structures and hydrogen bonding in the proton-transfer salts of nicotine with 3,5-dinitrosalicylic acid and 5-sulfosalicylic acid

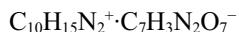
Graham Smith and Urs D. Wermuth

Computing details

For both compounds, data collection: *CrysAlis PRO* (Agilent, 2013); cell refinement: *CrysAlis PRO* (Agilent, 2013); data reduction: *CrysAlis PRO* (Agilent, 2013). Program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008) for (I); *SIR92* (Altomare *et al.*, 1993) for (II). For both compounds, program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008) within *WinGX* (Farrugia, 2012); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *PLATON* (Spek, 2009).

(I) (1*R*,2*S*)-1-Methyl-2-(pyridin-3-yl)pyrrolidin-1-ium 2-carboxy-4,6-dinitrophenolate

Crystal data



$M_r = 390.35$

Orthorhombic, $P2_12_12_1$

Hall symbol: P 2ac 2ab

$a = 6.8096 (5)$ Å

$b = 17.6403 (15)$ Å

$c = 29.3604 (19)$ Å

$V = 3526.9 (4)$ Å³

$Z = 8$

$F(000) = 1632$

$D_x = 1.470 \text{ Mg m}^{-3}$

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

Cell parameters from 1206 reflections

$\theta = 3.6\text{--}22.4^\circ$

$\mu = 0.12 \text{ mm}^{-1}$

$T = 200$ K

Needle, yellow

$0.40 \times 0.10 \times 0.08$ mm

Data collection

Oxford Diffraction Gemini-S CCD-detector
diffractometer

Radiation source: Enhance (Mo) X-ray source

Graphite monochromator

Detector resolution: 16.077 pixels mm⁻¹

ω scans

Absorption correction: multi-scan
(*CrysAlis PRO*; Agilent, 2013)

$T_{\min} = 0.807$, $T_{\max} = 0.980$

8840 measured reflections

6476 independent reflections

4303 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.028$

$\theta_{\max} = 26.0^\circ$, $\theta_{\min} = 3.2^\circ$

$h = -7 \rightarrow 8$

$k = -21 \rightarrow 9$

$l = -20 \rightarrow 36$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.072$

$wR(F^2) = 0.122$

$S = 1.07$

6476 reflections

508 parameters

2 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

$$w = 1/[\sigma^2(F_o^2) + (0.0329P)^2]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

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

$$\Delta\rho_{\max} = 0.44 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.19 \text{ e } \text{\AA}^{-3}$$

Absolute structure: Flack (1983), 2983 Friedel pairs

Absolute structure parameter: -0.2 (16)

Special details

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
O2B	0.7440 (5)	-0.0370 (2)	0.16873 (13)	0.0543 (14)	0.750
O11B	0.7783 (6)	-0.17264 (18)	0.06034 (13)	0.0812 (15)	
O12B	0.7503 (5)	-0.16123 (18)	0.13451 (12)	0.0722 (15)	
O31B	0.7213 (6)	0.1013 (2)	0.19951 (11)	0.0867 (16)	
O32B	0.8016 (5)	0.19443 (18)	0.15674 (11)	0.0645 (14)	
O51B	0.9215 (5)	0.16351 (18)	0.00036 (11)	0.0653 (14)	
O52B	0.9011 (5)	0.05377 (19)	-0.03182 (10)	0.0614 (11)	
N3B	0.7691 (5)	0.1272 (2)	0.16280 (13)	0.0440 (14)	
N5B	0.8927 (5)	0.0950 (2)	0.00167 (12)	0.0433 (12)	
C1B	0.7883 (5)	-0.0493 (2)	0.09179 (14)	0.0327 (14)	
C2B	0.7715 (5)	-0.0035 (2)	0.13069 (13)	0.0337 (14)	
C3B	0.7884 (5)	0.0757 (2)	0.12439 (13)	0.0283 (12)	
C4B	0.8272 (5)	0.1069 (2)	0.08283 (13)	0.0300 (11)	
C5B	0.8439 (5)	0.0611 (2)	0.04550 (13)	0.0287 (12)	
C6B	0.8230 (5)	-0.0181 (2)	0.04906 (14)	0.0333 (12)	
C11B	0.7727 (7)	-0.1329 (3)	0.09462 (19)	0.0543 (19)	
O21B	0.8109 (16)	-0.0648 (7)	0.0169 (4)	0.0543 (14)	0.250
O2A	0.2427 (6)	0.14566 (18)	0.18109 (11)	0.0750 (14)	
O11A	0.1135 (5)	0.0131 (2)	0.28861 (10)	0.0655 (14)	
O12A	0.1240 (5)	0.1263 (2)	0.25992 (10)	0.0650 (12)	
O31A	0.3353 (5)	0.1552 (2)	0.09522 (11)	0.0723 (16)	
O32A	0.3389 (5)	0.0535 (2)	0.05553 (11)	0.0784 (15)	
O51A	0.2682 (6)	-0.18960 (19)	0.11877 (14)	0.0817 (16)	
O52A	0.2223 (6)	-0.20342 (18)	0.19127 (14)	0.0889 (15)	
N3A	0.3211 (5)	0.0872 (2)	0.09182 (13)	0.0470 (14)	
N5A	0.2457 (6)	-0.1639 (2)	0.15757 (18)	0.0603 (16)	
C1A	0.2002 (5)	0.0247 (2)	0.21164 (13)	0.0333 (12)	
C2A	0.2429 (5)	0.0748 (2)	0.17458 (14)	0.0333 (14)	
C3A	0.2821 (5)	0.0401 (2)	0.13212 (13)	0.0287 (12)	
C4A	0.2834 (5)	-0.0373 (2)	0.12566 (14)	0.0350 (16)	
C5A	0.2445 (5)	-0.0816 (2)	0.16347 (16)	0.0383 (14)	
C6A	0.2026 (5)	-0.0513 (2)	0.20552 (14)	0.0370 (16)	

C11A	0.1438 (6)	0.0551 (3)	0.25629 (15)	0.0397 (16)	
N1C	0.9322 (4)	0.46384 (19)	0.08412 (11)	0.0342 (11)	
N11C	1.3932 (4)	0.34882 (17)	0.16230 (10)	0.0325 (10)	
C2C	1.0225 (5)	0.4293 (2)	0.11887 (13)	0.0297 (12)	
C3C	1.0826 (5)	0.3546 (2)	0.11738 (12)	0.0267 (12)	
C4C	1.0488 (5)	0.3140 (2)	0.07781 (13)	0.0367 (12)	
C5C	0.9526 (6)	0.3489 (3)	0.04199 (14)	0.0400 (16)	
C6C	0.8990 (6)	0.4232 (3)	0.04661 (14)	0.0413 (16)	
C11C	1.5274 (6)	0.3279 (3)	0.12449 (13)	0.0437 (17)	
C21C	1.1861 (5)	0.3191 (2)	0.15719 (12)	0.0297 (12)	
C31C	1.4563 (6)	0.3212 (3)	0.20791 (14)	0.0483 (17)	
C41C	1.2739 (7)	0.3266 (3)	0.23712 (14)	0.0563 (18)	
C51C	1.0999 (6)	0.3309 (3)	0.20426 (12)	0.0447 (16)	
N1D	0.4269 (4)	0.50744 (17)	0.16513 (11)	0.0331 (10)	
N11D	0.8855 (4)	0.62248 (19)	0.08623 (11)	0.0349 (11)	
C2D	0.5212 (5)	0.5403 (2)	0.13071 (14)	0.0330 (12)	
C3D	0.5708 (5)	0.6165 (2)	0.12973 (13)	0.0300 (12)	
C4D	0.5191 (5)	0.6592 (2)	0.16646 (14)	0.0373 (14)	
C5D	0.4196 (6)	0.6276 (2)	0.20240 (14)	0.0423 (17)	
C6D	0.3794 (6)	0.5509 (2)	0.20076 (14)	0.0393 (14)	
C11D	1.0141 (6)	0.6465 (3)	0.12373 (14)	0.0507 (16)	
C21D	0.6765 (5)	0.6508 (2)	0.08995 (12)	0.0330 (12)	
C31D	0.5957 (6)	0.6372 (3)	0.04297 (14)	0.0560 (17)	
C41D	0.7691 (8)	0.6443 (3)	0.01105 (15)	0.075 (2)	
C51D	0.9507 (7)	0.6480 (3)	0.03995 (14)	0.0557 (17)	
H6B	0.83210	-0.04960	0.02280	0.0400*	0.750
H12B	0.73670	-0.12610	0.15360	0.1080*	0.750
H4B	0.84250	0.16020	0.07990	0.0360*	
H2B	0.74970	-0.02470	0.16000	0.0410*	0.250
H11B	0.79170	-0.14500	0.03730	0.1220*	0.250
H4A	0.30970	-0.05920	0.09670	0.0420*	
H6A	0.17520	-0.08390	0.23050	0.0450*	
H12A	0.17450	0.14760	0.23720	0.0970*	
H2C	1.04650	0.45760	0.14580	0.0360*	
H4C	1.09110	0.26290	0.07530	0.0440*	
H5C	0.92430	0.32190	0.01480	0.0480*	
H6C	0.83470	0.44720	0.02180	0.0500*	
H11C	1.38680	0.40140	0.16350	0.0390*	
H12C	1.47550	0.34780	0.09570	0.0660*	
H13C	1.53770	0.27260	0.12260	0.0660*	
H14C	1.65760	0.34960	0.13010	0.0660*	
H21C	1.19410	0.26330	0.15160	0.0350*	
H31C	1.56270	0.35340	0.22040	0.0580*	
H32C	1.50340	0.26820	0.20610	0.0580*	
H41C	1.27900	0.37250	0.25650	0.0680*	
H42C	1.26220	0.28150	0.25700	0.0680*	
H51C	1.00240	0.29100	0.21130	0.0530*	
H52C	1.03470	0.38100	0.20630	0.0530*	

H2D	0.55660	0.50980	0.10530	0.0400*
H4D	0.55240	0.71140	0.16720	0.0450*
H5D	0.37950	0.65750	0.22760	0.0510*
H6D	0.31540	0.52800	0.22600	0.0470*
H11D	0.88190	0.56980	0.08620	0.0420*
H12D	1.14600	0.62570	0.11900	0.0760*
H13D	0.96120	0.62770	0.15270	0.0760*
H14D	1.02080	0.70200	0.12450	0.0760*
H21D	0.68180	0.70690	0.09490	0.0400*
H31D	0.49390	0.67530	0.03550	0.0670*
H32D	0.53690	0.58600	0.04080	0.0670*
H41D	0.77510	0.60010	-0.00960	0.0900*
H42D	0.75700	0.69090	-0.00760	0.0900*
H51D	1.05380	0.61390	0.02780	0.0670*
H52D	1.00290	0.70030	0.04110	0.0670*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O2B	0.059 (2)	0.060 (3)	0.044 (2)	0.001 (2)	0.001 (2)	0.008 (2)
O11B	0.126 (3)	0.0287 (19)	0.089 (3)	-0.005 (2)	0.007 (3)	-0.007 (2)
O12B	0.093 (3)	0.0386 (19)	0.085 (3)	-0.010 (2)	-0.006 (2)	0.021 (2)
O31B	0.134 (3)	0.082 (3)	0.044 (2)	-0.006 (3)	0.031 (2)	-0.015 (2)
O32B	0.092 (3)	0.0375 (19)	0.064 (2)	0.0084 (19)	-0.008 (2)	-0.0172 (19)
O51B	0.101 (3)	0.039 (2)	0.056 (2)	-0.018 (2)	0.0012 (19)	0.0146 (18)
O52B	0.095 (2)	0.060 (2)	0.0291 (17)	0.011 (2)	0.0128 (17)	0.0013 (17)
N3B	0.040 (2)	0.053 (3)	0.039 (2)	0.003 (2)	-0.0037 (18)	-0.009 (2)
N5B	0.046 (2)	0.047 (2)	0.037 (2)	0.001 (2)	0.0006 (18)	0.008 (2)
C1B	0.030 (2)	0.028 (2)	0.040 (3)	0.0029 (19)	-0.0067 (19)	0.005 (2)
C2B	0.024 (2)	0.048 (3)	0.029 (2)	0.002 (2)	0.0015 (18)	0.013 (2)
C3B	0.023 (2)	0.033 (2)	0.029 (2)	0.0040 (18)	0.0008 (18)	-0.005 (2)
C4B	0.0269 (19)	0.026 (2)	0.037 (2)	0.0032 (18)	-0.0050 (19)	0.000 (2)
C5B	0.032 (2)	0.029 (2)	0.025 (2)	0.0009 (18)	0.0003 (17)	0.0052 (19)
C6B	0.035 (2)	0.027 (2)	0.038 (2)	0.0014 (19)	0.001 (2)	-0.008 (2)
C11B	0.056 (3)	0.031 (3)	0.076 (4)	-0.004 (3)	-0.008 (3)	0.013 (3)
O21B	0.059 (2)	0.060 (3)	0.044 (2)	0.001 (2)	0.001 (2)	0.008 (2)
O2A	0.110 (3)	0.0401 (19)	0.075 (2)	0.004 (2)	-0.011 (2)	0.006 (2)
O11A	0.081 (2)	0.078 (3)	0.0375 (19)	-0.013 (2)	0.0043 (18)	0.002 (2)
O12A	0.085 (2)	0.059 (2)	0.051 (2)	0.004 (2)	0.0026 (17)	-0.012 (2)
O31A	0.104 (3)	0.040 (2)	0.073 (3)	0.011 (2)	0.027 (2)	0.026 (2)
O32A	0.110 (3)	0.095 (3)	0.0302 (18)	-0.012 (3)	0.008 (2)	0.006 (2)
O51A	0.097 (3)	0.044 (2)	0.104 (3)	-0.001 (2)	0.008 (3)	-0.031 (2)
O52A	0.117 (3)	0.0316 (19)	0.118 (3)	0.004 (2)	0.024 (3)	0.019 (2)
N3A	0.035 (2)	0.062 (3)	0.044 (2)	0.006 (2)	0.0001 (18)	0.014 (2)
N5A	0.057 (3)	0.034 (2)	0.090 (3)	-0.002 (2)	0.006 (3)	-0.003 (3)
C1A	0.032 (2)	0.035 (2)	0.033 (2)	0.000 (2)	-0.0052 (19)	0.004 (2)
C2A	0.028 (2)	0.023 (2)	0.049 (3)	0.005 (2)	-0.010 (2)	0.005 (2)
C3A	0.028 (2)	0.033 (2)	0.025 (2)	0.0035 (19)	0.0006 (18)	0.011 (2)

C4A	0.023 (2)	0.043 (3)	0.039 (3)	0.005 (2)	-0.0039 (19)	0.002 (2)
C5A	0.031 (2)	0.025 (2)	0.059 (3)	0.001 (2)	-0.002 (2)	0.003 (2)
C6A	0.032 (2)	0.039 (3)	0.040 (3)	0.003 (2)	0.003 (2)	0.015 (2)
C11A	0.037 (2)	0.042 (3)	0.040 (3)	-0.004 (2)	-0.004 (2)	-0.003 (2)
N1C	0.0364 (18)	0.036 (2)	0.0303 (19)	0.0046 (17)	-0.0010 (16)	-0.0028 (18)
N11C	0.0431 (18)	0.0196 (16)	0.0348 (18)	-0.0007 (16)	-0.0043 (17)	0.0030 (16)
C2C	0.036 (2)	0.030 (2)	0.023 (2)	0.0014 (19)	0.0002 (17)	0.001 (2)
C3C	0.028 (2)	0.020 (2)	0.032 (2)	-0.0048 (18)	0.0048 (17)	0.0000 (19)
C4C	0.042 (2)	0.029 (2)	0.039 (2)	-0.003 (2)	0.007 (2)	-0.006 (2)
C5C	0.050 (3)	0.041 (3)	0.029 (2)	0.000 (2)	-0.007 (2)	-0.003 (2)
C6C	0.045 (3)	0.045 (3)	0.034 (2)	0.008 (2)	-0.004 (2)	0.005 (2)
C11C	0.041 (3)	0.046 (3)	0.044 (3)	0.009 (2)	0.006 (2)	0.003 (2)
C21C	0.036 (2)	0.019 (2)	0.034 (2)	-0.0047 (17)	0.0022 (19)	0.0007 (19)
C31C	0.060 (3)	0.044 (3)	0.041 (3)	0.005 (2)	-0.017 (2)	0.010 (2)
C41C	0.087 (4)	0.053 (3)	0.029 (2)	-0.006 (3)	-0.004 (3)	0.003 (2)
C51C	0.061 (3)	0.042 (3)	0.031 (2)	0.007 (2)	0.008 (2)	0.009 (2)
N1D	0.0353 (17)	0.0241 (17)	0.040 (2)	-0.0021 (16)	-0.0002 (17)	0.0004 (18)
N11D	0.0427 (19)	0.0271 (18)	0.0349 (19)	-0.0011 (17)	0.0062 (16)	0.0028 (17)
C2D	0.033 (2)	0.029 (2)	0.037 (2)	0.003 (2)	0.0023 (19)	-0.003 (2)
C3D	0.029 (2)	0.026 (2)	0.035 (2)	0.0002 (19)	-0.0037 (19)	0.002 (2)
C4D	0.041 (2)	0.022 (2)	0.049 (3)	-0.0043 (19)	0.001 (2)	-0.005 (2)
C5D	0.049 (3)	0.038 (3)	0.040 (3)	0.000 (2)	0.009 (2)	-0.009 (2)
C6D	0.042 (2)	0.035 (2)	0.041 (3)	-0.004 (2)	0.008 (2)	-0.001 (2)
C11D	0.039 (2)	0.056 (3)	0.057 (3)	-0.008 (2)	-0.008 (2)	0.002 (3)
C21D	0.035 (2)	0.025 (2)	0.039 (2)	0.0035 (19)	0.0007 (19)	0.001 (2)
C31D	0.063 (3)	0.060 (3)	0.045 (3)	-0.012 (3)	-0.016 (3)	0.019 (3)
C41D	0.113 (4)	0.074 (4)	0.038 (3)	0.010 (4)	0.001 (3)	0.000 (3)
C51D	0.071 (3)	0.045 (3)	0.051 (3)	0.008 (3)	0.031 (3)	0.013 (3)

Geometric parameters (Å, °)

O2B—C2B	1.277 (5)	C4A—C5A	1.383 (6)
O11B—C11B	1.227 (7)	C5A—C6A	1.375 (6)
O12B—C11B	1.283 (7)	C4A—H4A	0.9500
O21B—C6B	1.256 (13)	C6A—H6A	0.9500
O31B—N3B	1.215 (5)	C2C—C3C	1.381 (5)
O32B—N3B	1.220 (5)	C3C—C21C	1.502 (5)
O51B—N5B	1.225 (5)	C3C—C4C	1.384 (5)
O52B—N5B	1.224 (5)	C4C—C5C	1.384 (6)
O11B—H11B	0.8400	C5C—C6C	1.367 (7)
O12B—H12B	0.8400	C21C—C51C	1.516 (5)
O2A—C2A	1.265 (5)	C31C—C41C	1.512 (6)
O11A—C11A	1.222 (6)	C41C—C51C	1.530 (6)
O12A—C11A	1.268 (6)	C2C—H2C	0.9500
O31A—N3A	1.208 (5)	C4C—H4C	0.9500
O32A—N3A	1.226 (5)	C5C—H5C	0.9500
O51A—N5A	1.236 (6)	C6C—H6C	0.9500
O52A—N5A	1.221 (6)	C11C—H14C	0.9800

O12A—H12A	0.8400	C11C—H12C	0.9800
N3B—C3B	1.454 (5)	C11C—H13C	0.9800
N5B—C5B	1.457 (5)	C21C—H21C	1.0000
N3A—C3A	1.470 (5)	C31C—H32C	0.9900
N5A—C5A	1.462 (5)	C31C—H31C	0.9900
N1C—C6C	1.333 (6)	C41C—H41C	0.9900
N1C—C2C	1.338 (5)	C41C—H42C	0.9900
N11C—C11C	1.485 (5)	C51C—H52C	0.9900
N11C—C21C	1.512 (4)	C51C—H51C	0.9900
N11C—C31C	1.488 (5)	C2D—C3D	1.386 (5)
N11C—H11C	0.9300	C3D—C21D	1.499 (5)
N1D—C2D	1.330 (5)	C3D—C4D	1.362 (5)
N1D—C6D	1.337 (5)	C4D—C5D	1.372 (6)
N11D—C51D	1.499 (5)	C5D—C6D	1.381 (5)
N11D—C11D	1.469 (5)	C21D—C31D	1.504 (5)
N11D—C21D	1.512 (4)	C31D—C41D	1.513 (7)
N11D—H11D	0.9300	C41D—C51D	1.501 (7)
C1B—C6B	1.390 (6)	C2D—H2D	0.9500
C1B—C11B	1.481 (6)	C4D—H4D	0.9500
C1B—C2B	1.404 (5)	C5D—H5D	0.9500
C2B—C3B	1.414 (5)	C6D—H6D	0.9500
C3B—C4B	1.364 (5)	C11D—H12D	0.9800
C4B—C5B	1.366 (5)	C11D—H13D	0.9800
C5B—C6B	1.408 (5)	C11D—H14D	0.9800
C2B—H2B	0.9500	C21D—H21D	1.0000
C4B—H4B	0.9500	C31D—H31D	0.9900
C6B—H6B	0.9500	C31D—H32D	0.9900
C1A—C6A	1.353 (5)	C41D—H41D	0.9900
C1A—C2A	1.432 (5)	C41D—H42D	0.9900
C1A—C11A	1.468 (6)	C51D—H51D	0.9900
C2A—C3A	1.414 (5)	C51D—H52D	0.9900
C3A—C4A	1.379 (5)		
C11B—O11B—H11B	109.00	C3C—C21C—C51C	118.1 (3)
C11B—O12B—H12B	110.00	N11C—C21C—C51C	102.9 (3)
C11A—O12A—H12A	109.00	N11C—C31C—C41C	104.6 (3)
O31B—N3B—O32B	122.9 (4)	C31C—C41C—C51C	106.4 (3)
O31B—N3B—C3B	118.5 (3)	C21C—C51C—C41C	105.6 (3)
O32B—N3B—C3B	118.6 (3)	N1C—C2C—H2C	118.00
O52B—N5B—C5B	118.4 (3)	C3C—C2C—H2C	119.00
O51B—N5B—O52B	123.6 (4)	C5C—C4C—H4C	120.00
O51B—N5B—C5B	118.0 (3)	C3C—C4C—H4C	120.00
O32A—N3A—C3A	116.3 (3)	C4C—C5C—H5C	121.00
O31A—N3A—C3A	120.6 (4)	C6C—C5C—H5C	121.00
O31A—N3A—O32A	123.1 (4)	C5C—C6C—H6C	118.00
O51A—N5A—O52A	123.6 (4)	N1C—C6C—H6C	118.00
O52A—N5A—C5A	118.1 (4)	N11C—C11C—H13C	110.00
O51A—N5A—C5A	118.3 (4)	H12C—C11C—H13C	109.00

C2C—N1C—C6C	117.6 (4)	N11C—C11C—H12C	109.00
C11C—N11C—C21C	114.4 (3)	H13C—C11C—H14C	110.00
C11C—N11C—C31C	114.4 (3)	N11C—C11C—H14C	110.00
C21C—N11C—C31C	104.2 (3)	H12C—C11C—H14C	109.00
C21C—N11C—H11C	108.00	C51C—C21C—H21C	108.00
C11C—N11C—H11C	108.00	C3C—C21C—H21C	108.00
C31C—N11C—H11C	108.00	N11C—C21C—H21C	108.00
C2D—N1D—C6D	117.5 (3)	C41C—C31C—H32C	111.00
C21D—N11D—C51D	104.2 (3)	N11C—C31C—H31C	111.00
C11D—N11D—C51D	114.6 (3)	C41C—C31C—H31C	111.00
C11D—N11D—C21D	114.3 (3)	H31C—C31C—H32C	109.00
C51D—N11D—H11D	108.00	N11C—C31C—H32C	111.00
C21D—N11D—H11D	108.00	C51C—C41C—H41C	110.00
C11D—N11D—H11D	108.00	H41C—C41C—H42C	109.00
C2B—C1B—C6B	121.4 (3)	C31C—C41C—H42C	110.00
C2B—C1B—C11B	121.5 (4)	C31C—C41C—H41C	110.00
C6B—C1B—C11B	117.2 (4)	C51C—C41C—H42C	110.00
C1B—C2B—C3B	117.1 (3)	H51C—C51C—H52C	109.00
O2B—C2B—C3B	125.7 (4)	C21C—C51C—H51C	111.00
O2B—C2B—C1B	117.2 (3)	C41C—C51C—H52C	111.00
N3B—C3B—C2B	120.6 (3)	C41C—C51C—H51C	111.00
N3B—C3B—C4B	117.3 (3)	C21C—C51C—H52C	111.00
C2B—C3B—C4B	122.1 (3)	N1D—C2D—C3D	123.8 (4)
C3B—C4B—C5B	119.7 (3)	C2D—C3D—C21D	121.6 (3)
N5B—C5B—C6B	119.7 (3)	C2D—C3D—C4D	117.2 (3)
C4B—C5B—C6B	121.2 (4)	C4D—C3D—C21D	121.2 (3)
N5B—C5B—C4B	119.0 (3)	C3D—C4D—C5D	120.8 (3)
O21B—C6B—C5B	127.0 (7)	C4D—C5D—C6D	118.0 (4)
C1B—C6B—C5B	118.5 (3)	N1D—C6D—C5D	122.8 (4)
O21B—C6B—C1B	114.1 (6)	C3D—C21D—C31D	118.3 (3)
O12B—C11B—C1B	116.6 (4)	N11D—C21D—C3D	112.0 (3)
O11B—C11B—O12B	122.0 (5)	N11D—C21D—C31D	103.0 (3)
O11B—C11B—C1B	121.4 (5)	C21D—C31D—C41D	105.6 (3)
C3B—C2B—H2B	121.00	C31D—C41D—C51D	107.2 (4)
C1B—C2B—H2B	122.00	N11D—C51D—C41D	104.8 (4)
C5B—C4B—H4B	120.00	N1D—C2D—H2D	118.00
C3B—C4B—H4B	120.00	C3D—C2D—H2D	118.00
C5B—C6B—H6B	121.00	C3D—C4D—H4D	120.00
C1B—C6B—H6B	121.00	C5D—C4D—H4D	120.00
C2A—C1A—C11A	120.4 (3)	C4D—C5D—H5D	121.00
C2A—C1A—C6A	120.6 (4)	C6D—C5D—H5D	121.00
C6A—C1A—C11A	119.0 (4)	N1D—C6D—H6D	119.00
C1A—C2A—C3A	116.2 (3)	C5D—C6D—H6D	119.00
O2A—C2A—C3A	124.2 (4)	N11D—C11D—H12D	109.00
O2A—C2A—C1A	119.7 (4)	N11D—C11D—H13D	109.00
N3A—C3A—C2A	119.9 (3)	N11D—C11D—H14D	109.00
N3A—C3A—C4A	116.6 (3)	H12D—C11D—H13D	109.00
C2A—C3A—C4A	123.5 (3)	H12D—C11D—H14D	110.00

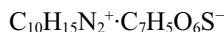
C3A—C4A—C5A	116.6 (4)	H13D—C11D—H14D	110.00
C4A—C5A—C6A	122.7 (3)	N11D—C21D—H21D	108.00
N5A—C5A—C4A	117.7 (4)	C3D—C21D—H21D	108.00
N5A—C5A—C6A	119.6 (4)	C31D—C21D—H21D	108.00
C1A—C6A—C5A	120.5 (4)	C21D—C31D—H31D	111.00
O11A—C11A—O12A	121.2 (4)	C21D—C31D—H32D	111.00
O11A—C11A—C1A	121.1 (4)	C41D—C31D—H31D	111.00
O12A—C11A—C1A	117.7 (4)	C41D—C31D—H32D	111.00
C3A—C4A—H4A	122.00	H31D—C31D—H32D	109.00
C5A—C4A—H4A	122.00	C31D—C41D—H41D	110.00
C5A—C6A—H6A	120.00	C31D—C41D—H42D	110.00
C1A—C6A—H6A	120.00	C51D—C41D—H41D	110.00
N1C—C2C—C3C	123.1 (3)	C51D—C41D—H42D	110.00
C2C—C3C—C4C	118.1 (3)	H41D—C41D—H42D	109.00
C4C—C3C—C21C	121.0 (3)	N11D—C51D—H51D	111.00
C2C—C3C—C21C	120.8 (3)	N11D—C51D—H52D	111.00
C3C—C4C—C5C	119.1 (4)	C41D—C51D—H51D	111.00
C4C—C5C—C6C	118.5 (4)	C41D—C51D—H52D	111.00
N1C—C6C—C5C	123.5 (4)	H51D—C51D—H52D	109.00
N11C—C21C—C3C	111.7 (3)		
O32B—N3B—C3B—C2B	173.7 (3)	N5B—C5B—C6B—C1B	176.1 (3)
O31B—N3B—C3B—C4B	175.0 (4)	C4B—C5B—C6B—C1B	-1.2 (5)
O32B—N3B—C3B—C4B	-5.0 (5)	C2A—C1A—C11A—O11A	-177.5 (4)
O31B—N3B—C3B—C2B	-6.3 (5)	C2A—C1A—C6A—C5A	-0.6 (5)
O52B—N5B—C5B—C6B	5.0 (5)	C11A—C1A—C2A—O2A	4.0 (5)
O52B—N5B—C5B—C4B	-177.7 (3)	C11A—C1A—C6A—C5A	176.5 (3)
O51B—N5B—C5B—C6B	-175.7 (3)	C11A—C1A—C2A—C3A	-175.6 (3)
O51B—N5B—C5B—C4B	1.7 (5)	C6A—C1A—C11A—O11A	5.5 (6)
O31A—N3A—C3A—C2A	5.8 (5)	C2A—C1A—C11A—O12A	3.8 (5)
O31A—N3A—C3A—C4A	-175.1 (4)	C6A—C1A—C2A—O2A	-179.0 (4)
O32A—N3A—C3A—C2A	-174.6 (3)	C6A—C1A—C11A—O12A	-173.2 (4)
O32A—N3A—C3A—C4A	4.5 (5)	C6A—C1A—C2A—C3A	1.4 (5)
O51A—N5A—C5A—C6A	174.3 (4)	C1A—C2A—C3A—N3A	178.0 (3)
O52A—N5A—C5A—C6A	-4.9 (6)	O2A—C2A—C3A—N3A	-1.6 (6)
O51A—N5A—C5A—C4A	-4.7 (6)	C1A—C2A—C3A—C4A	-1.0 (5)
O52A—N5A—C5A—C4A	176.0 (4)	O2A—C2A—C3A—C4A	179.4 (4)
C2C—N1C—C6C—C5C	-0.2 (6)	N3A—C3A—C4A—C5A	-179.3 (3)
C6C—N1C—C2C—C3C	0.3 (5)	C2A—C3A—C4A—C5A	-0.3 (5)
C31C—N11C—C21C—C3C	168.2 (3)	C3A—C4A—C5A—C6A	1.2 (5)
C31C—N11C—C21C—C51C	40.5 (4)	C3A—C4A—C5A—N5A	-179.8 (3)
C11C—N11C—C21C—C3C	-66.1 (4)	C4A—C5A—C6A—C1A	-0.8 (5)
C11C—N11C—C31C—C41C	-162.7 (4)	N5A—C5A—C6A—C1A	-179.8 (3)
C11C—N11C—C21C—C51C	166.3 (3)	N1C—C2C—C3C—C4C	0.5 (5)
C21C—N11C—C31C—C41C	-36.9 (4)	N1C—C2C—C3C—C21C	179.1 (3)
C6D—N1D—C2D—C3D	-0.5 (5)	C4C—C3C—C21C—N11C	106.6 (4)
C2D—N1D—C6D—C5D	1.8 (6)	C2C—C3C—C21C—C51C	47.1 (5)
C21D—N11D—C51D—C41D	-33.4 (4)	C2C—C3C—C21C—N11C	-71.9 (4)

C51D—N11D—C21D—C31D	39.4 (4)	C4C—C3C—C21C—C51C	-134.4 (4)
C11D—N11D—C21D—C3D	-66.6 (4)	C2C—C3C—C4C—C5C	-1.6 (5)
C11D—N11D—C21D—C31D	165.2 (4)	C21C—C3C—C4C—C5C	179.9 (3)
C51D—N11D—C21D—C3D	167.6 (3)	C3C—C4C—C5C—C6C	1.8 (6)
C11D—N11D—C51D—C41D	-159.0 (4)	C4C—C5C—C6C—N1C	-0.9 (6)
C6B—C1B—C11B—O11B	-4.3 (6)	C3C—C21C—C51C—C41C	-151.8 (4)
C11B—C1B—C2B—C3B	-180.0 (3)	N11C—C21C—C51C—C41C	-28.2 (4)
C11B—C1B—C2B—O2B	0.6 (5)	N11C—C31C—C41C—C51C	18.9 (5)
C2B—C1B—C11B—O12B	-2.5 (6)	C31C—C41C—C51C—C21C	6.1 (5)
C6B—C1B—C2B—C3B	0.9 (5)	N1D—C2D—C3D—C21D	-179.7 (3)
C6B—C1B—C11B—O12B	176.7 (4)	N1D—C2D—C3D—C4D	0.1 (5)
C6B—C1B—C2B—O2B	-178.5 (3)	C2D—C3D—C4D—C5D	-0.9 (5)
C2B—C1B—C11B—O11B	176.6 (4)	C21D—C3D—C4D—C5D	179.0 (3)
C11B—C1B—C6B—C5B	-178.3 (3)	C4D—C3D—C21D—C31D	-129.0 (4)
C2B—C1B—C6B—C5B	0.9 (5)	C4D—C3D—C21D—N11D	111.4 (4)
C1B—C2B—C3B—C4B	-2.6 (5)	C2D—C3D—C21D—C31D	50.8 (5)
C1B—C2B—C3B—N3B	178.9 (3)	C2D—C3D—C21D—N11D	-68.8 (4)
O2B—C2B—C3B—N3B	-1.8 (6)	C3D—C4D—C5D—C6D	2.0 (6)
O2B—C2B—C3B—C4B	176.8 (4)	C4D—C5D—C6D—N1D	-2.5 (6)
C2B—C3B—C4B—C5B	2.3 (5)	N11D—C21D—C31D—C41D	-30.0 (4)
N3B—C3B—C4B—C5B	-179.1 (3)	C3D—C21D—C31D—C41D	-154.1 (4)
C3B—C4B—C5B—N5B	-177.7 (3)	C21D—C31D—C41D—C51D	9.8 (5)
C3B—C4B—C5B—C6B	-0.4 (5)	C31D—C41D—C51D—N11D	14.6 (5)

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

$D—H\cdots A$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
O12A—H12A…O2A	0.84	1.71	2.475 (4)	150
O12B—H12B…O2B	0.84	1.63	2.411 (4)	152
N11C—H11C…N1D ⁱ	0.93	1.89	2.809 (4)	169
N11D—H11D…N1C	0.93	1.90	2.817 (5)	168
C2C—H2C…O11A ⁱⁱ	0.95	2.42	3.228 (5)	143
C4C—H4C…O31A ⁱ	0.95	2.59	3.452 (5)	151
C5D—H5D…O31B ⁱⁱ	0.95	2.46	3.071 (5)	122
C6C—H6C…O32A ⁱⁱⁱ	0.95	2.27	3.054 (5)	139
C6D—H6D…O31B ⁱⁱ	0.95	2.55	3.136 (5)	120
C11C—H13C…O32B ⁱ	0.98	2.48	3.151 (6)	126
C11D—H14D…O51A ^{iv}	0.98	2.55	3.373 (6)	141
C21C—H21C…O2A ⁱ	1.00	2.27	3.163 (5)	148
C21D—H21D…O11B ^v	1.00	2.44	3.307 (5)	144
C51C—H52C…O11A ⁱⁱ	0.99	2.54	3.534 (6)	177

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

(II) (1*R*,2*S*)-1-Methyl-2-(pyridin-3-yl)pyrrolidin-1-i um 3-carboxy-4-hydroxybenzenesulfonate*Crystal data*

$M_r = 380.41$

Monoclinic, $P2_1$

Hall symbol: P 2yb

$a = 7.1568 (3)$ Å

$b = 12.6416 (5)$ Å

$c = 19.1519 (8)$ Å

$\beta = 93.729 (4)^\circ$

$V = 1729.07 (12)$ Å³

$Z = 4$

$F(000) = 800$

$D_x = 1.461 \text{ Mg m}^{-3}$

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

Cell parameters from 2239 reflections

$\theta = 3.4\text{--}27.5^\circ$

$\mu = 0.23 \text{ mm}^{-1}$

$T = 200$ K

Prism, colourless

0.35 × 0.30 × 0.12 mm

Data collection

Oxford Diffraction Gemini-S CCD-detector
diffractometer

Radiation source: Enhance (Mo) X-ray source

Graphite monochromator

Detector resolution: 16.077 pixels mm⁻¹

ω scans

Absorption correction: multi-scan
(*CrysAlis PRO*; Agilent, 2013)

$T_{\min} = 0.909$, $T_{\max} = 0.981$

7764 measured reflections

5104 independent reflections

4424 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.031$

$\theta_{\max} = 28.9^\circ$, $\theta_{\min} = 3.2^\circ$

$h = -8 \rightarrow 9$

$k = -17 \rightarrow 8$

$l = -24 \rightarrow 14$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.046$

$wR(F^2) = 0.108$

$S = 1.01$

5104 reflections

469 parameters

1 restraint

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

$w = 1/[\sigma^2(F_o^2) + (0.0448P)^2 + 0.6152P]$
where $P = (F_o^2 + 2F_c^2)/3$

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

$\Delta\rho_{\max} = 0.49 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.36 \text{ e } \text{\AA}^{-3}$

Absolute structure: Flack (1983), 4361 Friedel
pairs

Absolute structure parameter: -0.02 (9)

Special details

Geometry. Bond distances, angles etc. have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

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 (Å²)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
S5A	0.68704 (12)	0.55842 (7)	0.41012 (4)	0.0263 (3)
O2A	0.3397 (4)	0.3912 (2)	0.14516 (12)	0.0340 (8)
O11A	0.3204 (4)	0.2139 (2)	0.32641 (13)	0.0348 (8)

O12A	0.2584 (4)	0.2236 (2)	0.21069 (13)	0.0337 (8)
O51A	0.8728 (4)	0.5131 (3)	0.41339 (15)	0.0508 (10)
O52A	0.6922 (4)	0.6735 (2)	0.40180 (14)	0.0422 (9)
O53A	0.5792 (4)	0.5253 (2)	0.46723 (12)	0.0393 (9)
C1A	0.4129 (4)	0.3676 (3)	0.26886 (17)	0.0221 (9)
C2A	0.4124 (5)	0.4284 (3)	0.20674 (17)	0.0222 (10)
C3A	0.4845 (5)	0.5304 (3)	0.20917 (17)	0.0240 (10)
C4A	0.5643 (4)	0.5705 (3)	0.27120 (17)	0.0239 (9)
C5A	0.5696 (4)	0.5108 (3)	0.33217 (17)	0.0211 (9)
C6A	0.4930 (5)	0.4106 (3)	0.33044 (18)	0.0234 (10)
C11A	0.3243 (5)	0.2614 (3)	0.26686 (19)	0.0259 (11)
S5B	0.30149 (14)	-0.14181 (8)	0.08257 (5)	0.0345 (3)
O2B	0.6834 (4)	0.0076 (2)	0.34578 (12)	0.0344 (8)
O11B	0.7029 (4)	0.1928 (2)	0.16767 (13)	0.0347 (8)
O12B	0.7680 (4)	0.1780 (2)	0.28273 (13)	0.0335 (8)
O51B	0.1062 (4)	-0.1205 (3)	0.09470 (17)	0.0625 (11)
O52B	0.3370 (4)	-0.2529 (2)	0.07618 (16)	0.0480 (10)
O53B	0.3571 (6)	-0.0759 (3)	0.02651 (16)	0.0801 (15)
C1B	0.6053 (4)	0.0375 (3)	0.22345 (17)	0.0206 (9)
C2B	0.6058 (5)	-0.0256 (3)	0.28333 (17)	0.0233 (10)
C3B	0.5299 (5)	-0.1274 (3)	0.28007 (18)	0.0255 (10)
C4B	0.4433 (5)	-0.1631 (3)	0.21848 (19)	0.0261 (10)
C5B	0.4331 (5)	-0.0982 (3)	0.15869 (17)	0.0231 (10)
C6B	0.5145 (4)	0.0004 (3)	0.16130 (17)	0.0222 (10)
C11B	0.6976 (5)	0.1427 (3)	0.22658 (19)	0.0260 (11)
N1C	0.8536 (4)	0.3758 (3)	0.18074 (15)	0.0270 (9)
N11C	0.9801 (4)	0.5668 (3)	-0.00002 (14)	0.0295 (9)
C2C	0.8558 (5)	0.4408 (3)	0.12534 (19)	0.0253 (10)
C3C	0.9371 (4)	0.5393 (3)	0.12903 (18)	0.0257 (10)
C4C	1.0161 (5)	0.5724 (3)	0.19371 (18)	0.0294 (10)
C5C	1.0096 (5)	0.5074 (3)	0.25102 (18)	0.0296 (11)
C6C	0.9288 (5)	0.4092 (3)	0.24258 (19)	0.0297 (11)
C11C	1.1769 (5)	0.5314 (4)	-0.0030 (2)	0.0392 (14)
C21C	0.9336 (5)	0.6148 (3)	0.06816 (18)	0.0290 (11)
C31C	0.7439 (6)	0.6662 (4)	0.0489 (2)	0.0493 (16)
C41C	0.7621 (7)	0.7097 (4)	-0.0242 (2)	0.0529 (17)
C51C	0.9232 (6)	0.6511 (4)	-0.0534 (2)	0.0476 (16)
N1D	0.1735 (4)	0.0297 (2)	0.31881 (15)	0.0257 (8)
N11D	-0.0524 (4)	-0.2131 (3)	0.48368 (14)	0.0327 (10)
C2D	0.1839 (5)	-0.0236 (3)	0.37895 (18)	0.0239 (10)
C3D	0.1071 (4)	-0.1235 (3)	0.38558 (16)	0.0211 (9)
C4D	0.0196 (5)	-0.1695 (3)	0.32573 (17)	0.0256 (10)
C5D	0.0119 (5)	-0.1153 (3)	0.26338 (18)	0.0283 (11)
C6D	0.0898 (5)	-0.0152 (3)	0.26175 (19)	0.0284 (11)
C11D	-0.1872 (6)	-0.1290 (4)	0.4998 (2)	0.0430 (14)
C21D	0.1287 (5)	-0.1769 (3)	0.45539 (17)	0.0267 (10)
C31D	0.2434 (6)	-0.2778 (4)	0.4595 (2)	0.0442 (14)
C41D	0.1992 (8)	-0.3244 (4)	0.5301 (2)	0.0626 (19)

C51D	0.0103 (7)	-0.2808 (4)	0.5453 (2)	0.0564 (16)
H2A	0.30050	0.32920	0.15050	0.0510*
H3A	0.47890	0.57260	0.16810	0.0290*
H4A	0.61610	0.63970	0.27220	0.0290*
H6A	0.49510	0.37020	0.37230	0.0280*
H11A	0.26610	0.14530	0.32280	0.0520*
H2B	0.71920	0.07050	0.34230	0.0520*
H3B	0.53810	-0.17170	0.32020	0.0310*
H4B	0.39000	-0.23190	0.21620	0.0310*
H6B	0.50900	0.04360	0.12060	0.0270*
H11B	0.75890	0.26070	0.17310	0.0520*
H2C	0.79870	0.41780	0.08180	0.0300*
H4C	1.07430	0.63980	0.19820	0.0350*
H5A	1.06000	0.53000	0.29570	0.0350*
H6C	0.92630	0.36360	0.28190	0.0360*
H11C	0.90280	0.50850	-0.00850	0.0350*
H12C	1.20770	0.48160	0.03520	0.0580*
H13C	1.26050	0.59270	0.00170	0.0580*
H21C	1.02610	0.67240	0.07990	0.0350*
H31C	0.64180	0.61330	0.04910	0.0590*
H32C	0.71820	0.72360	0.08200	0.0590*
H41C	0.64530	0.69750	-0.05370	0.0630*
H42C	0.78790	0.78660	-0.02240	0.0630*
H51C	1.02880	0.69990	-0.06010	0.0570*
H52C	0.88390	0.61850	-0.09900	0.0570*
H134	1.19250	0.49640	-0.04800	0.0580*
H12D	-0.21690	-0.08640	0.45780	0.0650*
H2D	0.24650	0.00810	0.41890	0.0290*
H4D	-0.03450	-0.23790	0.32810	0.0310*
H5D	-0.04590	-0.14600	0.22220	0.0340*
H6D	0.08380	0.02270	0.21880	0.0340*
H11D	-0.11170	-0.25710	0.45020	0.0390*
H21D	0.18880	-0.12540	0.48950	0.0320*
H13D	-0.13180	-0.08360	0.53710	0.0650*
H14D	-0.30210	-0.16130	0.51510	0.0650*
H31D	0.37870	-0.26240	0.45810	0.0530*
H32D	0.20480	-0.32650	0.42080	0.0530*
H41D	0.19550	-0.40260	0.52770	0.0750*
H42D	0.29520	-0.30300	0.56690	0.0750*
H51D	-0.07990	-0.33910	0.55110	0.0680*
H52D	0.01960	-0.23820	0.58870	0.0680*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S5A	0.0295 (4)	0.0253 (5)	0.0232 (4)	-0.0031 (4)	-0.0062 (3)	0.0026 (4)
O2A	0.0467 (15)	0.0280 (15)	0.0261 (13)	-0.0022 (13)	-0.0058 (11)	-0.0012 (11)
O11A	0.0456 (15)	0.0234 (14)	0.0352 (14)	-0.0115 (13)	0.0018 (11)	0.0042 (12)

O12A	0.0397 (14)	0.0249 (14)	0.0360 (14)	-0.0065 (13)	-0.0013 (11)	-0.0045 (12)
O51A	0.0347 (14)	0.073 (2)	0.0424 (16)	0.0124 (17)	-0.0146 (12)	0.0011 (17)
O52A	0.0580 (18)	0.0242 (14)	0.0417 (16)	-0.0116 (14)	-0.0180 (13)	0.0052 (13)
O53A	0.0496 (15)	0.0460 (17)	0.0219 (12)	-0.0115 (15)	-0.0002 (10)	0.0003 (12)
C1A	0.0201 (16)	0.0212 (17)	0.0249 (16)	0.0028 (16)	0.0010 (12)	0.0024 (15)
C2A	0.0238 (17)	0.0227 (18)	0.0197 (16)	0.0035 (15)	-0.0009 (13)	0.0010 (14)
C3A	0.0271 (17)	0.0256 (18)	0.0193 (16)	0.0050 (16)	0.0008 (13)	0.0056 (14)
C4A	0.0213 (15)	0.0197 (17)	0.0305 (17)	-0.0002 (16)	0.0009 (13)	0.0055 (16)
C5A	0.0221 (16)	0.0196 (17)	0.0210 (16)	-0.0011 (16)	-0.0025 (12)	0.0008 (14)
C6A	0.0264 (18)	0.0203 (18)	0.0233 (17)	0.0010 (16)	0.0002 (13)	0.0055 (14)
C11A	0.0231 (17)	0.0221 (19)	0.0326 (19)	0.0001 (15)	0.0022 (14)	-0.0005 (16)
S5B	0.0413 (5)	0.0300 (5)	0.0305 (5)	-0.0046 (5)	-0.0110 (4)	-0.0026 (4)
O2B	0.0499 (15)	0.0292 (14)	0.0227 (12)	-0.0051 (14)	-0.0078 (11)	-0.0009 (11)
O11B	0.0440 (15)	0.0269 (15)	0.0330 (14)	-0.0111 (13)	0.0007 (11)	0.0065 (12)
O12B	0.0442 (15)	0.0270 (14)	0.0286 (14)	-0.0052 (13)	-0.0029 (11)	-0.0020 (12)
O51B	0.0379 (16)	0.070 (2)	0.075 (2)	0.0237 (18)	-0.0305 (15)	-0.029 (2)
O52B	0.0402 (16)	0.0333 (17)	0.068 (2)	0.0016 (14)	-0.0145 (14)	-0.0229 (16)
O53B	0.126 (3)	0.084 (3)	0.0272 (16)	-0.057 (3)	-0.0197 (18)	0.0087 (18)
C1B	0.0202 (15)	0.0139 (17)	0.0277 (16)	0.0026 (14)	0.0020 (12)	-0.0005 (13)
C2B	0.0234 (17)	0.0248 (19)	0.0218 (17)	0.0048 (16)	0.0013 (13)	-0.0008 (15)
C3B	0.0310 (18)	0.0187 (17)	0.0263 (17)	0.0035 (17)	-0.0021 (13)	0.0067 (15)
C4B	0.0242 (17)	0.0175 (18)	0.0358 (19)	0.0016 (15)	-0.0052 (14)	0.0029 (15)
C5B	0.0211 (16)	0.0219 (18)	0.0260 (17)	0.0044 (15)	-0.0002 (13)	0.0007 (15)
C6B	0.0222 (16)	0.0218 (18)	0.0224 (16)	0.0028 (15)	-0.0001 (13)	0.0025 (14)
C11B	0.0249 (17)	0.0193 (18)	0.034 (2)	0.0029 (15)	0.0047 (14)	0.0001 (16)
N1C	0.0256 (14)	0.0258 (17)	0.0294 (15)	0.0004 (14)	0.0003 (11)	0.0076 (14)
N11C	0.0363 (15)	0.0269 (16)	0.0251 (14)	-0.0033 (15)	0.0003 (11)	0.0064 (14)
C2C	0.0250 (18)	0.0242 (19)	0.0263 (17)	0.0007 (16)	-0.0012 (13)	0.0008 (15)
C3C	0.0225 (16)	0.0238 (19)	0.0312 (18)	0.0015 (15)	0.0041 (13)	0.0038 (15)
C4C	0.0239 (16)	0.0269 (19)	0.0375 (19)	-0.0031 (16)	0.0036 (14)	-0.0039 (17)
C5C	0.0255 (18)	0.038 (2)	0.0249 (18)	0.0011 (17)	-0.0003 (14)	-0.0009 (17)
C6C	0.0263 (18)	0.035 (2)	0.0278 (18)	0.0032 (17)	0.0025 (14)	0.0108 (17)
C11C	0.038 (2)	0.044 (3)	0.036 (2)	-0.003 (2)	0.0049 (16)	-0.006 (2)
C21C	0.0339 (19)	0.0202 (17)	0.0333 (19)	0.0005 (16)	0.0064 (14)	0.0042 (16)
C31C	0.048 (2)	0.043 (3)	0.058 (3)	0.020 (2)	0.013 (2)	0.024 (2)
C41C	0.064 (3)	0.038 (3)	0.055 (3)	0.005 (2)	-0.009 (2)	0.020 (2)
C51C	0.060 (3)	0.048 (3)	0.035 (2)	0.003 (2)	0.0038 (19)	0.024 (2)
N1D	0.0265 (14)	0.0213 (15)	0.0299 (15)	-0.0009 (13)	0.0067 (11)	0.0047 (13)
N11D	0.0428 (18)	0.0322 (17)	0.0230 (15)	-0.0165 (15)	0.0009 (12)	-0.0011 (14)
C2D	0.0264 (17)	0.0190 (17)	0.0265 (17)	0.0002 (15)	0.0033 (13)	0.0007 (14)
C3D	0.0215 (15)	0.0182 (17)	0.0239 (16)	0.0011 (15)	0.0033 (12)	-0.0009 (14)
C4D	0.0266 (17)	0.0226 (18)	0.0278 (18)	-0.0038 (15)	0.0038 (13)	0.0005 (15)
C5D	0.0253 (18)	0.035 (2)	0.0241 (17)	-0.0020 (17)	-0.0020 (13)	0.0008 (16)
C6D	0.0281 (18)	0.033 (2)	0.0245 (18)	0.0016 (17)	0.0058 (14)	0.0052 (16)
C11D	0.040 (2)	0.052 (3)	0.038 (2)	-0.010 (2)	0.0113 (17)	-0.012 (2)
C21D	0.0310 (18)	0.0252 (18)	0.0231 (17)	-0.0095 (16)	-0.0031 (13)	-0.0007 (14)
C31D	0.058 (3)	0.037 (2)	0.036 (2)	0.012 (2)	-0.0081 (19)	0.005 (2)
C41D	0.099 (4)	0.039 (3)	0.048 (3)	0.011 (3)	-0.008 (3)	0.019 (2)

C51D	0.077 (3)	0.061 (3)	0.030 (2)	-0.022 (3)	-0.006 (2)	0.023 (2)
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Geometric parameters (\AA , $\text{^{\circ}}$)

S5A—O51A	1.445 (3)	C3B—H3B	0.9500
S5A—O52A	1.464 (3)	C4B—H4B	0.9500
S5A—O53A	1.441 (3)	C6B—H6B	0.9500
S5A—C5A	1.770 (3)	C2C—C3C	1.374 (5)
S5B—O51B	1.457 (3)	C3C—C21C	1.506 (5)
S5B—O52B	1.434 (3)	C3C—C4C	1.392 (5)
S5B—O53B	1.436 (4)	C4C—C5C	1.374 (5)
S5B—C5B	1.771 (4)	C5C—C6C	1.375 (5)
O2A—C2A	1.343 (4)	C21C—C31C	1.529 (6)
O11A—C11A	1.291 (4)	C31C—C41C	1.518 (6)
O12A—C11A	1.242 (4)	C41C—C51C	1.508 (7)
O2A—H2A	0.8400	C2C—H2C	0.9500
O11A—H11A	0.9500	C4C—H4C	0.9500
O2B—C2B	1.352 (4)	C5C—H5A	0.9500
O11B—C11B	1.297 (4)	C6C—H6C	0.9500
O12B—C11B	1.241 (4)	C11C—H134	0.9800
O2B—H2B	0.8400	C11C—H12C	0.9800
O11B—H11B	0.9500	C11C—H13C	0.9800
N1C—C6C	1.337 (5)	C21C—H21C	1.0000
N1C—C2C	1.343 (5)	C31C—H32C	0.9900
N11C—C11C	1.483 (5)	C31C—H31C	0.9900
N11C—C51C	1.514 (6)	C41C—H41C	0.9900
N11C—C21C	1.497 (5)	C41C—H42C	0.9900
N11C—H11C	0.9300	C51C—H52C	0.9900
N1D—C2D	1.332 (4)	C51C—H51C	0.9900
N1D—C6D	1.338 (5)	C2D—C3D	1.386 (5)
N11D—C51D	1.503 (5)	C3D—C21D	1.497 (5)
N11D—C11D	1.482 (6)	C3D—C4D	1.397 (5)
N11D—C21D	1.508 (5)	C4D—C5D	1.375 (5)
N11D—H11D	0.9300	C5D—C6D	1.384 (5)
C1A—C2A	1.416 (5)	C21D—C31D	1.516 (6)
C1A—C11A	1.484 (5)	C31D—C41D	1.527 (6)
C1A—C6A	1.388 (5)	C41D—C51D	1.506 (7)
C2A—C3A	1.389 (5)	C2D—H2D	0.9500
C3A—C4A	1.381 (5)	C4D—H4D	0.9500
C4A—C5A	1.389 (5)	C5D—H5D	0.9500
C5A—C6A	1.380 (5)	C6D—H6D	0.9500
C3A—H3A	0.9500	C11D—H12D	0.9800
C4A—H4A	0.9500	C11D—H13D	0.9800
C6A—H6A	0.9500	C11D—H14D	0.9800
C1B—C6B	1.399 (5)	C21D—H21D	1.0000
C1B—C11B	1.484 (5)	C31D—H31D	0.9900
C1B—C2B	1.397 (5)	C31D—H32D	0.9900
C2B—C3B	1.397 (5)	C41D—H41D	0.9900

C3B—C4B	1.373 (5)	C41D—H42D	0.9900
C4B—C5B	1.407 (5)	C51D—H51D	0.9900
C5B—C6B	1.375 (5)	C51D—H52D	0.9900
O51A—S5A—O52A	111.65 (19)	N11C—C21C—C31C	101.9 (3)
O51A—S5A—O53A	112.86 (17)	C21C—C31C—C41C	104.4 (3)
O51A—S5A—C5A	106.66 (17)	C31C—C41C—C51C	106.1 (4)
O52A—S5A—O53A	112.94 (16)	N11C—C51C—C41C	105.9 (3)
O52A—S5A—C5A	105.07 (17)	N1C—C2C—H2C	119.00
O53A—S5A—C5A	107.03 (15)	C3C—C2C—H2C	119.00
O51B—S5B—C5B	105.99 (18)	C5C—C4C—H4C	120.00
O51B—S5B—O52B	111.8 (2)	C3C—C4C—H4C	120.00
O51B—S5B—O53B	109.1 (2)	C4C—C5C—H5A	121.00
O53B—S5B—C5B	106.1 (2)	C6C—C5C—H5A	121.00
O52B—S5B—O53B	116.4 (2)	C5C—C6C—H6C	119.00
O52B—S5B—C5B	106.70 (18)	N1C—C6C—H6C	119.00
C2A—O2A—H2A	109.00	N11C—C11C—H13C	109.00
C11A—O11A—H11A	113.00	H12C—C11C—H13C	109.00
C2B—O2B—H2B	109.00	N11C—C11C—H12C	109.00
C11B—O11B—H11B	112.00	H13C—C11C—H134	109.00
C2C—N1C—C6C	118.8 (4)	N11C—C11C—H134	109.00
C11C—N11C—C21C	114.9 (3)	H12C—C11C—H134	109.00
C11C—N11C—C51C	113.6 (3)	C31C—C21C—H21C	108.00
C21C—N11C—C51C	103.8 (3)	C3C—C21C—H21C	108.00
C51C—N11C—H11C	108.00	N11C—C21C—H21C	108.00
C21C—N11C—H11C	108.00	C41C—C31C—H32C	111.00
C11C—N11C—H11C	108.00	C21C—C31C—H31C	111.00
C2D—N1D—C6D	119.2 (3)	C41C—C31C—H31C	111.00
C11D—N11D—C51D	114.2 (3)	H31C—C31C—H32C	109.00
C11D—N11D—C21D	116.3 (3)	C21C—C31C—H32C	111.00
C21D—N11D—C51D	103.6 (3)	C51C—C41C—H41C	110.00
C51D—N11D—H11D	107.00	H41C—C41C—H42C	109.00
C11D—N11D—H11D	107.00	C31C—C41C—H42C	111.00
C21D—N11D—H11D	107.00	C31C—C41C—H41C	110.00
C6A—C1A—C11A	121.9 (3)	C51C—C41C—H42C	110.00
C2A—C1A—C11A	119.4 (3)	H51C—C51C—H52C	109.00
C2A—C1A—C6A	118.7 (3)	N11C—C51C—H51C	111.00
O2A—C2A—C1A	121.8 (3)	C41C—C51C—H52C	111.00
O2A—C2A—C3A	118.5 (3)	C41C—C51C—H51C	111.00
C1A—C2A—C3A	119.7 (3)	N11C—C51C—H52C	111.00
C2A—C3A—C4A	120.1 (3)	N1D—C2D—C3D	122.7 (3)
C3A—C4A—C5A	120.7 (3)	C2D—C3D—C21D	118.3 (3)
C4A—C5A—C6A	119.3 (3)	C2D—C3D—C4D	117.6 (3)
S5A—C5A—C4A	120.7 (3)	C4D—C3D—C21D	124.1 (3)
S5A—C5A—C6A	119.9 (3)	C3D—C4D—C5D	119.7 (3)
C1A—C6A—C5A	121.4 (3)	C4D—C5D—C6D	118.8 (3)
O11A—C11A—C1A	115.6 (3)	N1D—C6D—C5D	121.9 (3)
O12A—C11A—C1A	120.5 (3)	C3D—C21D—C31D	116.7 (3)

O11A—C11A—O12A	123.9 (3)	N11D—C21D—C3D	114.6 (3)
C2A—C3A—H3A	120.00	N11D—C21D—C31D	101.6 (3)
C4A—C3A—H3A	120.00	C21D—C31D—C41D	103.2 (3)
C3A—C4A—H4A	120.00	C31D—C41D—C51D	105.7 (4)
C5A—C4A—H4A	120.00	N11D—C51D—C41D	106.4 (3)
C5A—C6A—H6A	119.00	N1D—C2D—H2D	119.00
C1A—C6A—H6A	119.00	C3D—C2D—H2D	119.00
C6B—C1B—C11B	121.0 (3)	C3D—C4D—H4D	120.00
C2B—C1B—C11B	120.1 (3)	C5D—C4D—H4D	120.00
C2B—C1B—C6B	118.9 (3)	C4D—C5D—H5D	121.00
C1B—C2B—C3B	120.6 (3)	C6D—C5D—H5D	121.00
O2B—C2B—C3B	117.5 (3)	N1D—C6D—H6D	119.00
O2B—C2B—C1B	121.9 (3)	C5D—C6D—H6D	119.00
C2B—C3B—C4B	119.6 (3)	N11D—C11D—H12D	109.00
C3B—C4B—C5B	120.3 (3)	N11D—C11D—H13D	110.00
C4B—C5B—C6B	120.0 (3)	N11D—C11D—H14D	109.00
S5B—C5B—C4B	119.0 (3)	H12D—C11D—H13D	109.00
S5B—C5B—C6B	120.9 (3)	H12D—C11D—H14D	109.00
C1B—C6B—C5B	120.5 (3)	H13D—C11D—H14D	109.00
O12B—C11B—C1B	120.8 (3)	N11D—C21D—H21D	108.00
O11B—C11B—C1B	116.1 (3)	C3D—C21D—H21D	108.00
O11B—C11B—O12B	123.1 (3)	C31D—C21D—H21D	108.00
C2B—C3B—H3B	120.00	C21D—C31D—H31D	111.00
C4B—C3B—H3B	120.00	C21D—C31D—H32D	111.00
C3B—C4B—H4B	120.00	C41D—C31D—H31D	111.00
C5B—C4B—H4B	120.00	C41D—C31D—H32D	111.00
C5B—C6B—H6B	120.00	H31D—C31D—H32D	109.00
C1B—C6B—H6B	120.00	C31D—C41D—H41D	111.00
N1C—C2C—C3C	122.7 (3)	C31D—C41D—H42D	111.00
C2C—C3C—C4C	117.7 (3)	C51D—C41D—H41D	111.00
C4C—C3C—C21C	118.9 (3)	C51D—C41D—H42D	111.00
C2C—C3C—C21C	123.3 (3)	H41D—C41D—H42D	109.00
C3C—C4C—C5C	119.9 (3)	N11D—C51D—H51D	110.00
C4C—C5C—C6C	118.8 (3)	N11D—C51D—H52D	111.00
N1C—C6C—C5C	122.1 (3)	C41D—C51D—H51D	110.00
N11C—C21C—C3C	115.2 (3)	C41D—C51D—H52D	110.00
C3C—C21C—C31C	115.2 (3)	H51D—C51D—H52D	109.00
O51A—S5A—C5A—C4A	-95.6 (3)	C6B—C1B—C2B—O2B	-176.9 (3)
O51A—S5A—C5A—C6A	80.1 (3)	C6B—C1B—C2B—C3B	5.0 (5)
O52A—S5A—C5A—C4A	23.0 (3)	C11B—C1B—C2B—C3B	-175.9 (3)
O52A—S5A—C5A—C6A	-161.3 (3)	C2B—C1B—C6B—C5B	-2.3 (5)
O53A—S5A—C5A—C4A	143.4 (3)	C11B—C1B—C6B—C5B	178.6 (3)
O53A—S5A—C5A—C6A	-41.0 (3)	C2B—C1B—C11B—O11B	175.1 (3)
O53B—S5B—C5B—C6B	19.4 (4)	C2B—C1B—C11B—O12B	-3.5 (5)
O52B—S5B—C5B—C4B	-40.0 (3)	C6B—C1B—C11B—O11B	-5.8 (5)
O51B—S5B—C5B—C4B	79.3 (3)	C11B—C1B—C2B—O2B	2.2 (5)
O51B—S5B—C5B—C6B	-96.5 (3)	C6B—C1B—C11B—O12B	175.6 (3)

O52B—S5B—C5B—C6B	144.1 (3)	O2B—C2B—C3B—C4B	177.5 (3)
O53B—S5B—C5B—C4B	-164.8 (3)	C1B—C2B—C3B—C4B	-4.2 (5)
C2C—N1C—C6C—C5C	-0.6 (5)	C2B—C3B—C4B—C5B	0.8 (5)
C6C—N1C—C2C—C3C	1.9 (5)	C3B—C4B—C5B—C6B	1.7 (5)
C21C—N11C—C51C—C41C	-30.3 (4)	C3B—C4B—C5B—S5B	-174.1 (3)
C11C—N11C—C21C—C3C	-67.9 (4)	C4B—C5B—C6B—C1B	-1.0 (5)
C11C—N11C—C51C—C41C	-155.8 (4)	S5B—C5B—C6B—C1B	174.8 (2)
C51C—N11C—C21C—C31C	41.9 (4)	N1C—C2C—C3C—C4C	-1.4 (5)
C11C—N11C—C21C—C31C	166.6 (4)	N1C—C2C—C3C—C21C	-177.4 (3)
C51C—N11C—C21C—C3C	167.4 (3)	C2C—C3C—C4C—C5C	-0.5 (5)
C2D—N1D—C6D—C5D	0.6 (5)	C21C—C3C—C4C—C5C	175.7 (3)
C6D—N1D—C2D—C3D	-1.6 (5)	C2C—C3C—C21C—N11C	-45.7 (4)
C51D—N11D—C21D—C3D	169.5 (3)	C2C—C3C—C21C—C31C	72.5 (5)
C11D—N11D—C21D—C31D	168.9 (3)	C4C—C3C—C21C—C31C	-103.4 (4)
C21D—N11D—C51D—C41D	-27.3 (4)	C4C—C3C—C21C—N11C	138.4 (3)
C51D—N11D—C21D—C31D	42.6 (3)	C3C—C4C—C5C—C6C	1.7 (5)
C11D—N11D—C51D—C41D	-154.9 (4)	C4C—C5C—C6C—N1C	-1.2 (6)
C11D—N11D—C21D—C3D	-64.3 (4)	N11C—C21C—C31C—C41C	-38.1 (4)
C2A—C1A—C11A—O12A	3.6 (5)	C3C—C21C—C31C—C41C	-163.5 (3)
C2A—C1A—C11A—O11A	-175.9 (3)	C21C—C31C—C41C—C51C	19.6 (5)
C6A—C1A—C11A—O11A	2.6 (5)	C31C—C41C—C51C—N11C	6.2 (5)
C6A—C1A—C11A—O12A	-178.0 (3)	N1D—C2D—C3D—C4D	1.4 (5)
C6A—C1A—C2A—C3A	-2.6 (5)	N1D—C2D—C3D—C21D	179.3 (3)
C11A—C1A—C2A—O2A	-2.4 (5)	C2D—C3D—C4D—C5D	-0.2 (5)
C11A—C1A—C2A—C3A	176.0 (3)	C21D—C3D—C4D—C5D	-177.9 (3)
C2A—C1A—C6A—C5A	0.6 (5)	C2D—C3D—C21D—N11D	125.7 (3)
C11A—C1A—C6A—C5A	-177.9 (3)	C2D—C3D—C21D—C31D	-115.7 (4)
C6A—C1A—C2A—O2A	179.1 (3)	C4D—C3D—C21D—N11D	-56.6 (5)
C1A—C2A—C3A—C4A	3.1 (5)	C4D—C3D—C21D—C31D	62.0 (5)
O2A—C2A—C3A—C4A	-178.5 (3)	C3D—C4D—C5D—C6D	-0.8 (5)
C2A—C3A—C4A—C5A	-1.6 (5)	C4D—C5D—C6D—N1D	0.6 (5)
C3A—C4A—C5A—C6A	-0.4 (5)	N11D—C21D—C31D—C41D	-41.6 (4)
C3A—C4A—C5A—S5A	175.3 (3)	C3D—C21D—C31D—C41D	-167.0 (3)
C4A—C5A—C6A—C1A	0.9 (5)	C21D—C31D—C41D—C51D	25.0 (5)
S5A—C5A—C6A—C1A	-174.9 (3)	C31D—C41D—C51D—N11D	1.3 (5)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
O2A—H2A···O12A	0.84	1.80	2.549 (4)	147
O2B—H2B···O12B	0.84	1.82	2.561 (4)	146
O11A—H11A···N1D	0.95	1.60	2.555 (4)	179
O11B—H11B···N1C	0.95	1.61	2.558 (4)	179
N11C—H11C···O51B ⁱ	0.93	2.32	3.022 (5)	132
N11C—H11C···O53B ⁱ	0.93	2.15	3.029 (5)	157
N11D—H11D···O52A ⁱⁱ	0.93	1.85	2.735 (4)	158
C11D—H12D···O2B ⁱⁱⁱ	0.98	2.51	3.491 (5)	174
C2C—H2C···O53B ⁱ	0.95	2.29	3.201 (5)	160

C2D—H2D···O53A ^{iv}	0.95	2.45	3.359 (4)	160
C4A—H4A···O52A	0.95	2.54	2.914 (4)	103
C6B—H6B···O53B	0.95	2.54	2.913 (5)	103
C11C—H12C···O2A ^v	0.98	2.52	3.481 (5)	165
C11C—H13C···O52B ^{vi}	0.98	2.46	3.290 (5)	142
C11D—H13D···O51A ^{iv}	0.98	2.37	3.251 (5)	150
C21C—H21C···O52B ^{vi}	1.00	2.42	3.331 (5)	151

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