

Dipotassium disulfanilamidate trihydrate

Fiona N.-F. How,^a Z. A. Rahima,^a Hamid Khaledi^{b*} and Hapipah Mohd Ali^b

^aDepartment of Biomedical Sciences, Kulliyah of Science, IIUM Kuantan, 25200 Kuantan, Malaysia, and ^bDepartment of Chemistry, University of Malaya, 50603 Kuala Lumpur, Malaysia

Correspondence e-mail: khaledi@siswa.um.edu.my

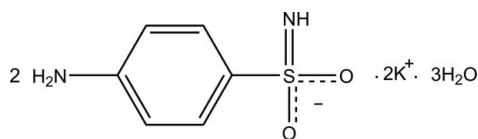
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.015; wR factor = 0.042; data-to-parameter ratio = 14.9.

The asymmetric unit of the title compound, $2\text{K}^+ \cdot 2\text{C}_6\text{H}_7\text{N}_2\text{O}_2\text{S}^- \cdot 3\text{H}_2\text{O}$, consists of two potassium cations located on mirror planes, one sulfanilamidate anion in a general position and one and a half molecules of water, one of which is also located on a mirror plane. One potassium cation is seven-coordinated by six sulfonyl O atoms and one water molecule, whereas the other is surrounded by six water O atoms and two sulfonyl O atoms. In the crystal structure, the components are connected into polymeric sheets in the bc plane. The two-dimensional structure is consolidated by $\text{N} \cdots \text{H} \cdots \text{O}$, $\text{O} \cdots \text{H} \cdots \text{O}$, $\text{O} \cdots \text{H} \cdots \text{N}$ and $\text{C} \cdots \text{H} \cdots \pi$ interactions. The layers are further linked into a three-dimensional network via $\text{N} \cdots \text{H} \cdots \text{O}$, $\text{N} \cdots \text{H} \cdots \text{N}$ and $\text{O} \cdots \text{H} \cdots \text{N}$ hydrogen bonds.

Related literature

For the structures of similar potassium salts, see: Gowda *et al.* (2011) and references cited therein; Moers *et al.* (2001). For the structure of sodium sulfanilamide monohydrate, see: Moreno & Alleaume (1968).



Experimental

Crystal data

$2\text{K}^+ \cdot 2\text{C}_6\text{H}_7\text{N}_2\text{O}_2\text{S}^- \cdot 3\text{H}_2\text{O}$
 $M_r = 474.64$
 Orthorhombic, $Cmc2_1$
 $a = 23.8174$ (4) Å
 $b = 10.9141$ (2) Å
 $c = 7.4645$ (1) Å

$V = 1940.36$ (5) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.75$ mm⁻¹
 $T = 100$ K
 $0.33 \times 0.27 \times 0.10$ mm

Data collection

Bruker APEXII CCD
 diffractometer
 Absorption correction: multi-scan
 (*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.791$, $T_{\max} = 0.929$

8594 measured reflections
 2157 independent reflections
 2149 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.015$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.015$
 $wR(F^2) = 0.042$
 $S = 1.11$
 2157 reflections
 145 parameters
 7 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.26$ e Å⁻³
 $\Delta\rho_{\min} = -0.23$ e Å⁻³
 Absolute structure: Flack (1983), 985 Friedel pairs
 Flack parameter: 0.04 (3)

Table 1

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of the C1–C6 ring.

$D \cdots H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$\text{N1} \cdots \text{H1} \cdots \text{O1}^{\text{i}}$	0.88 (1)	2.17 (1)	2.9854 (15)	154 (2)
$\text{N2} \cdots \text{H2A} \cdots \text{N1}^{\text{ii}}$	0.90 (1)	2.13 (1)	3.0109 (16)	166 (1)
$\text{N2} \cdots \text{H2B} \cdots \text{O3}^{\text{iii}}$	0.91 (1)	2.12 (1)	3.0183 (15)	171 (1)
$\text{O3} \cdots \text{H3A} \cdots \text{N2}^{\text{iv}}$	0.85 (2)	1.99 (2)	2.8366 (15)	172 (2)
$\text{O3} \cdots \text{H3B} \cdots \text{O2}^{\text{v}}$	0.83 (2)	1.95 (2)	2.7561 (13)	164 (2)
$\text{O4} \cdots \text{H4} \cdots \text{N1}^{\text{vi}}$	0.79 (1)	2.05 (1)	2.8291 (14)	175 (2)
$\text{C2} \cdots \text{H2} \cdots \text{Cg1}^{\text{i}}$	0.95	2.98	3.5872 (14)	123
$\text{C5} \cdots \text{H5} \cdots \text{Cg1}^{\text{ii}}$	0.95	2.66	3.4531 (14)	141

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

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *SHELXL97* and *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: OM2446).

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supplementary materials

Acta Cryst. (2011). E67, m1071 [doi:10.1107/S1600536811026791]

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Comment

The title potassium salt (Fig. 1) was obtained through the deprotonation of sulfanilamide by KOH. The structure contains two types of potassium cation, one of which, K1, is hepta-coordinated by six sulfonyl O atoms and one water molecule while the other one, K2, is octa-coordinated by six water O atoms and two sulfonyl O atoms. Both potassium atoms and one water oxygen atom, O4, are placed on a mirror plane.

The bond distance of S—N [1.5410 (12) Å] implies its double bond character and is slightly shorter than the observed values (~ 1.58 Å) in similar potassium salts (Gowda *et al.*, 2011; Moers *et al.*, 2001), but comparable with the bond length reported for sodium sulfanilamide (Moreno & Alleaume, 1968). The S1—O1 and S1—O2 distances of 1.4686 (9) and 1.4705 (9) Å attest to delocalization of the negative charge over the O—S—O fragment. These values are similar to the corresponding values in sodium sulfanilamide (1.45–1.46 Å).

In the crystal, coordination polymeric layers are formed in the *bc* plane (Fig. 2). The two dimensional structure is supplemented by N—H···O, O—H···O, O—H···N and C—H··· π interactions (Table 1). The layers are further linked into a three dimensional network *via* N—H···O, N—H···N and O—H···N hydrogen bonds (Fig. 3)

Experimental

Sulfanilamide (2 g) in ethanol (30 ml) was mixed with an equimolar amount of KOH in 90% ethanol (10 ml). The solution was then refluxed for 2 hr and left to cool down at room temperature. The colorless crystals of the the potassium salt were obtained within a day.

Refinement

The C-bound H atoms were placed at calculated positions and were treated as riding on their parent C atoms with C—H = 0.95 Å. The N- and O-bound H atoms were located in a difference Fourier map, and refined with distance restraints of O—H = 0.84 (2) Å and N—H = 0.91 (2) Å. For all H atoms, $U_{\text{iso}}(\text{H})$ was set to 1.2(1.5 for hydroxyl) $U_{\text{eq}}(\text{carrier atom})$. An absolute structure was established using anomalous dispersion effects; 985 Friedel pairs were not merged.

Figures

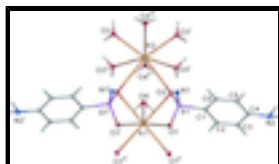


Fig. 1. Thermal ellipsoid plot of the title compound at the 50% probability level. Hydrogen atoms are drawn as spheres of arbitrary radius. Symmetry code: i = $-x + 1, y, z$; ii = $x, -y + 1, z + 1/2$; iii = $-x + 1, -y + 1, z + 1/2$; iv = $x, -y, z + 1/2$; v = $-x + 1, -y, z + 1/2$; vi = $x, y, z - 1$; vii = $-x + 1, -y, z - 1/2$.

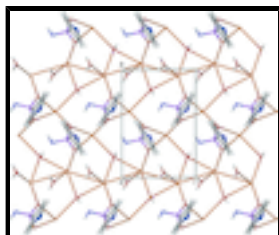


Fig. 2. Packing view looking down the crystallographic a axis showing the two dimensional coordination polymer network.

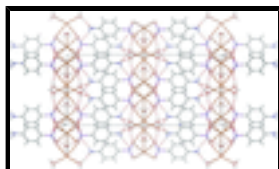
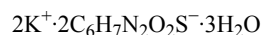


Fig. 3. Packing view looking down the crystallographic c axis. H-bonds are shown as red dashed lines.

Dipotassium disulfanilamidate trihydrate

Crystal data



$$M_r = 474.64$$

Orthorhombic, $Cmc2_1$

Hall symbol: C 2c -2

$$a = 23.8174 (4) \text{ \AA}$$

$$b = 10.9141 (2) \text{ \AA}$$

$$c = 7.4645 (1) \text{ \AA}$$

$$V = 1940.36 (5) \text{ \AA}^3$$

$$Z = 4$$

$$F(000) = 984$$

$$D_x = 1.625 \text{ Mg m}^{-3}$$

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

Cell parameters from 9001 reflections

$$\theta = 2.7\text{--}30.5^\circ$$

$$\mu = 0.75 \text{ mm}^{-1}$$

$$T = 100 \text{ K}$$

Blade, colorless

$$0.33 \times 0.27 \times 0.10 \text{ mm}$$

Data collection

Bruker APEXII CCD
diffractometer

Radiation source: fine-focus sealed tube
graphite

ϕ and ω scans

Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)

$$T_{\min} = 0.791, T_{\max} = 0.929$$

8594 measured reflections

2157 independent reflections

2149 reflections with $I > 2\sigma(I)$

$$R_{\text{int}} = 0.015$$

$$\theta_{\max} = 27.0^\circ, \theta_{\min} = 2.1^\circ$$

$$h = -30 \rightarrow 30$$

$$k = -13 \rightarrow 13$$

$$l = -9 \rightarrow 9$$

Refinement

Refinement on F^2

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.015$$

$$wR(F^2) = 0.042$$

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

$$w = 1/[\sigma^2(F_o^2) + (0.0201P)^2 + 1.0371P]$$

$S = 1.11$	where $P = (F_o^2 + 2F_c^2)/3$
2157 reflections	$(\Delta/\sigma)_{\max} < 0.001$
145 parameters	$\Delta\rho_{\max} = 0.26 \text{ e } \text{\AA}^{-3}$
7 restraints	$\Delta\rho_{\min} = -0.23 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	Absolute structure: Flack (1983), 985 Friedel pairs Flack parameter: 0.04 (3)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
K1	0.5000	0.39021 (4)	0.59975 (5)	0.01535 (8)
K2	0.5000	0.05477 (3)	0.26065 (6)	0.01442 (8)
S1	0.404982 (11)	0.35192 (2)	0.25404 (4)	0.01158 (7)
O1	0.42926 (4)	0.47094 (8)	0.30427 (12)	0.01535 (19)
O2	0.42708 (4)	0.25122 (8)	0.36386 (12)	0.01634 (19)
N1	0.41122 (4)	0.31915 (11)	0.05418 (15)	0.0147 (2)
H1	0.4057 (7)	0.3869 (13)	−0.007 (2)	0.018*
N2	0.15806 (5)	0.38097 (11)	0.38621 (15)	0.0148 (2)
H2A	0.1423 (6)	0.3121 (13)	0.430 (2)	0.018*
H2B	0.1402 (6)	0.4057 (14)	0.285 (2)	0.018*
C1	0.33256 (5)	0.36379 (11)	0.30816 (17)	0.0124 (2)
C2	0.30175 (5)	0.46150 (10)	0.23864 (18)	0.0138 (2)
H2	0.3202	0.5243	0.1732	0.017*
C3	0.24417 (5)	0.46705 (10)	0.26498 (19)	0.0134 (2)
H3	0.2234	0.5342	0.2183	0.016*
C4	0.21636 (5)	0.37450 (11)	0.35982 (16)	0.0126 (2)
C5	0.24773 (5)	0.27905 (12)	0.43346 (18)	0.0144 (2)
H5	0.2295	0.2175	0.5023	0.017*
C6	0.30546 (5)	0.27342 (11)	0.40674 (17)	0.0138 (2)
H6	0.3265	0.2075	0.4561	0.017*
O3	0.58605 (4)	−0.04452 (9)	0.07181 (13)	0.0173 (2)
H3A	0.6103 (6)	0.0020 (18)	0.023 (3)	0.026*
H3B	0.5880 (8)	−0.1110 (15)	0.019 (3)	0.026*
O4	0.5000	0.18900 (13)	0.8893 (2)	0.0207 (3)
H4	0.4747 (6)	0.2267 (16)	0.929 (3)	0.031*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
K1	0.01301 (16)	0.01830 (18)	0.01474 (17)	0.000	0.000	−0.00331 (15)
K2	0.01360 (16)	0.01577 (17)	0.01391 (16)	0.000	0.000	−0.00172 (17)
S1	0.01032 (12)	0.01235 (13)	0.01206 (13)	−0.00085 (9)	0.00042 (13)	0.00013 (13)
O1	0.0144 (4)	0.0149 (4)	0.0168 (5)	−0.0035 (3)	0.0009 (3)	−0.0022 (3)
O2	0.0134 (4)	0.0169 (4)	0.0188 (5)	0.0009 (3)	−0.0002 (4)	0.0048 (4)
N1	0.0158 (5)	0.0161 (5)	0.0123 (5)	−0.0002 (4)	0.0024 (4)	−0.0005 (4)
N2	0.0118 (5)	0.0178 (5)	0.0148 (5)	0.0007 (4)	0.0004 (4)	0.0019 (4)
C1	0.0113 (6)	0.0142 (5)	0.0118 (5)	−0.0010 (5)	0.0009 (4)	−0.0027 (4)
C2	0.0163 (5)	0.0129 (5)	0.0120 (6)	−0.0015 (4)	0.0019 (5)	0.0017 (5)
C3	0.0157 (5)	0.0123 (5)	0.0121 (5)	0.0017 (4)	−0.0004 (5)	0.0008 (5)
C4	0.0121 (6)	0.0156 (6)	0.0102 (6)	−0.0001 (5)	0.0009 (4)	−0.0028 (5)
C5	0.0161 (6)	0.0136 (5)	0.0136 (5)	−0.0019 (5)	0.0017 (5)	0.0016 (5)
C6	0.0139 (6)	0.0134 (5)	0.0142 (6)	0.0010 (4)	−0.0007 (5)	0.0016 (5)
O3	0.0177 (4)	0.0155 (4)	0.0187 (5)	−0.0009 (3)	0.0033 (4)	−0.0010 (4)
O4	0.0135 (7)	0.0205 (7)	0.0280 (8)	0.000	0.000	−0.0097 (6)

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

K1—O1 ⁱ	2.7325 (9)	N1—H1	0.878 (14)
K1—O1 ⁱⁱ	2.7325 (9)	N2—C4	1.4044 (16)
K1—O2 ⁱⁱⁱ	2.9014 (10)	N2—H2A	0.903 (13)
K1—O2	2.9014 (10)	N2—H2B	0.906 (14)
K1—O1	2.9121 (10)	C1—C6	1.3895 (17)
K1—O1 ⁱⁱⁱ	2.9121 (10)	C1—C2	1.3947 (17)
K1—O4	3.0810 (16)	C2—C3	1.3868 (17)
K2—O3	2.7133 (10)	C2—H2	0.9500
K2—O3 ⁱⁱⁱ	2.7133 (10)	C3—C4	1.4000 (17)
K2—O4 ^{iv}	2.8285 (15)	C3—H3	0.9500
K2—O2 ⁱⁱⁱ	2.8648 (10)	C4—C5	1.3948 (18)
K2—O2	2.8648 (10)	C5—C6	1.3908 (17)
K2—O3 ^v	3.0996 (10)	C5—H5	0.9500
K2—O3 ^{vi}	3.0996 (10)	C6—H6	0.9500
K2—O4 ^{vii}	3.1355 (17)	O3—K2 ^{iv}	3.0996 (10)
S1—O2	1.4686 (9)	O3—H3A	0.852 (15)
S1—O1	1.4705 (9)	O3—H3B	0.828 (15)
S1—N1	1.5413 (12)	O4—K2 ^{vi}	2.8284 (15)
S1—C1	1.7763 (13)	O4—K2 ^{ix}	3.1355 (17)
O1—K1 ^{viii}	2.7324 (9)	O4—H4	0.787 (13)
O1 ⁱ —K1—O1 ⁱⁱ	76.15 (4)	O3 ^{vi} —K2—O4 ^{vii}	132.79 (2)
O1 ⁱ —K1—O2 ⁱⁱⁱ	105.06 (3)	O2—S1—O1	112.21 (6)
O1 ⁱⁱ —K1—O2 ⁱⁱⁱ	176.54 (3)	O2—S1—N1	109.40 (6)

O1 ⁱ —K1—O2	176.54 (3)	O1—S1—N1	114.44 (6)
O1 ⁱⁱ —K1—O2	105.06 (3)	O2—S1—C1	106.00 (6)
O2 ⁱⁱⁱ —K1—O2	73.54 (4)	O1—S1—C1	105.03 (6)
O1 ⁱ —K1—O1	127.75 (2)	N1—S1—C1	109.31 (6)
O1 ⁱⁱ —K1—O1	84.18 (2)	S1—O1—K1 ^{viii}	126.15 (5)
O2 ⁱⁱⁱ —K1—O1	92.57 (3)	S1—O1—K1	98.82 (4)
O2—K1—O1	49.62 (3)	K1 ^{viii} —O1—K1	103.54 (3)
O1 ⁱ —K1—O1 ⁱⁱⁱ	84.18 (2)	S1—O2—K2	128.84 (5)
O1 ⁱⁱ —K1—O1 ⁱⁱⁱ	127.75 (2)	S1—O2—K1	99.32 (5)
O2 ⁱⁱⁱ —K1—O1 ⁱⁱⁱ	49.62 (3)	K2—O2—K1	101.04 (3)
O2—K1—O1 ⁱⁱⁱ	92.58 (3)	S1—N1—H1	106.9 (11)
O1—K1—O1 ⁱⁱⁱ	70.71 (4)	C4—N2—H2A	114.8 (10)
O1 ⁱ —K1—O4	90.20 (3)	C4—N2—H2B	111.3 (10)
O1 ⁱⁱ —K1—O4	90.20 (3)	H2A—N2—H2B	110.9 (15)
O2 ⁱⁱⁱ —K1—O4	93.04 (3)	C6—C1—C2	119.69 (11)
O2—K1—O4	93.04 (3)	C6—C1—S1	121.31 (10)
O1—K1—O4	138.32 (2)	C2—C1—S1	118.83 (10)
O1 ⁱⁱⁱ —K1—O4	138.32 (2)	C3—C2—C1	120.07 (11)
O3—K2—O3 ⁱⁱⁱ	98.12 (4)	C3—C2—H2	120.0
O3—K2—O4 ^{iv}	78.50 (3)	C1—C2—H2	120.0
O3 ⁱⁱⁱ —K2—O4 ^{iv}	78.50 (3)	C2—C3—C4	120.53 (11)
O3—K2—O2 ⁱⁱⁱ	88.89 (3)	C2—C3—H3	119.7
O3 ⁱⁱⁱ —K2—O2 ⁱⁱⁱ	153.71 (3)	C4—C3—H3	119.7
O4 ^{iv} —K2—O2 ⁱⁱⁱ	127.79 (3)	C5—C4—C3	119.00 (11)
O3—K2—O2	153.71 (3)	C5—C4—N2	120.79 (11)
O3 ⁱⁱⁱ —K2—O2	88.89 (3)	C3—C4—N2	120.15 (11)
O4 ^{iv} —K2—O2	127.78 (3)	C6—C5—C4	120.40 (12)
O2 ⁱⁱⁱ —K2—O2	74.64 (4)	C6—C5—H5	119.8
O3—K2—O3 ^v	82.84 (3)	C4—C5—H5	119.8
O3 ⁱⁱⁱ —K2—O3 ^v	150.97 (2)	C1—C6—C5	120.25 (11)
O4 ^{iv} —K2—O3 ^v	73.24 (3)	C1—C6—H6	119.9
O2 ⁱⁱⁱ —K2—O3 ^v	54.88 (3)	C5—C6—H6	119.9
O2—K2—O3 ^v	103.08 (3)	K2—O3—K2 ^{iv}	84.51 (3)
O3—K2—O3 ^{vi}	150.97 (2)	K2—O3—H3A	119.9 (14)
O3 ⁱⁱⁱ —K2—O3 ^{vi}	82.84 (3)	K2 ^{iv} —O3—H3A	98.5 (13)
O4 ^{iv} —K2—O3 ^{vi}	73.24 (3)	K2—O3—H3B	129.9 (13)
O2 ⁱⁱⁱ —K2—O3 ^{vi}	103.08 (3)	K2 ^{iv} —O3—H3B	69.3 (13)
O2—K2—O3 ^{vi}	54.87 (3)	H3A—O3—H3B	106 (2)
O3 ^v —K2—O3 ^{vi}	82.79 (4)	K2 ^{vi} —O4—K1	115.62 (5)
O3—K2—O4 ^{vii}	74.17 (3)	K2 ^{vi} —O4—K2 ^{ix}	81.99 (4)
O3 ⁱⁱⁱ —K2—O4 ^{vii}	74.17 (3)	K1—O4—K2 ^{ix}	162.39 (5)

supplementary materials

O4 ^{iv} —K2—O4 ^{vii}	137.70 (5)	K2 ^{vi} —O4—H4	128.2 (13)
O2 ⁱⁱⁱ —K2—O4 ^{vii}	83.57 (3)	K1—O4—H4	83.7 (15)
O2—K2—O4 ^{vii}	83.57 (3)	K2 ^{ix} —O4—H4	85.1 (16)
O3 ^v —K2—O4 ^{vii}	132.79 (2)		

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

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

Cg1 is the centroid of the C1–C6 ring.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1 \cdots O1 ^x	0.88 (1)	2.17 (1)	2.9854 (15)	154.(2)
N2—H2A \cdots N1 ^{xi}	0.90 (1)	2.13 (1)	3.0109 (16)	166.(1)
N2—H2B \cdots O3 ^{xii}	0.91 (1)	2.12 (1)	3.0183 (15)	171.(1)
O3—H3A \cdots N2 ^{xiii}	0.85 (2)	1.99 (2)	2.8366 (15)	172.(2)
O3—H3B \cdots O2 ^{iv}	0.83 (2)	1.95 (2)	2.7561 (13)	164.(2)
O4—H4 \cdots N1 ^{ix}	0.79 (1)	2.05 (1)	2.8291 (14)	175 (2)
C2—H2 \cdots Cg1 ^x	0.95	2.98	3.5872 (14)	123
C5—H5 \cdots Cg1 ^{xi}	0.95	2.66	3.4531 (14)	141

Symmetry codes: (x) $x, -y+1, z-1/2$; (xi) $-x+1/2, -y+1/2, z+1/2$; (xii) $x-1/2, y+1/2, z$; (xiii) $x+1/2, -y+1/2, z-1/2$; (iv) $-x+1, -y, z-1/2$; (ix) $x, y, z+1$.

Fig. 1

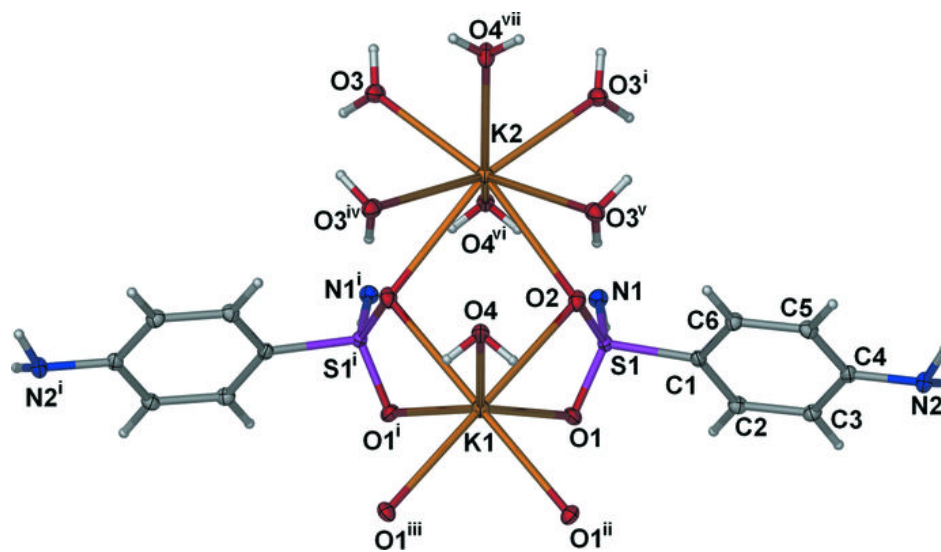


Fig. 2

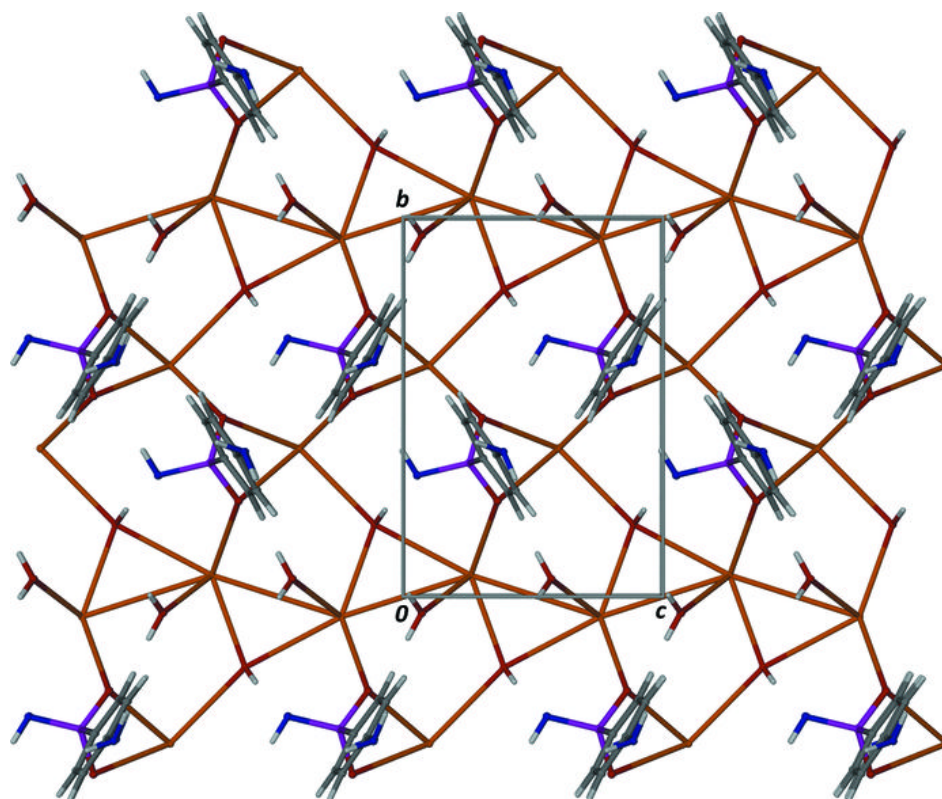


Fig. 3

