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Disodium dihydrogen pyridine-2,3,5,6-tetracarboxylate trihydrate

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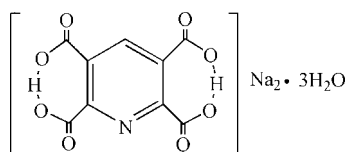
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.043; wR factor = 0.139; data-to-parameter ratio = 10.8.

In the title compound, $2\text{Na}^+\cdot\text{C}_9\text{H}_3\text{NO}_8^{2-}\cdot 3\text{H}_2\text{O}$, the asymmetric unit consists of two Na^+ cations, one dihydrogen pyridine-2,3,5,6-tetracarboxylate dianion ($\text{H}_2\text{pdtc}^{2-}$) and three water molecules coordinated to the Na^+ cations. The configuration of the anion is stabilized by intramolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonding between vicinal carboxylate/carboxy groups. The Na^+ cations are bridged by the $\text{H}_2\text{pdtc}^{2-}$ dianions, generating layers extending infinitely in sheets parallel to (001), and further pillared by the water molecule linkers to build up a three-dimensional framework.

Related literature

For related compounds involving the pyridine-2,3,5,6-tetracarboxylic acid ligand, see: Zhang *et al.* (2010); Yang *et al.* (2008); Sun, Zhou & An (2009); Sun, Zhou & Yan (2009).



Experimental

Crystal data

 $2\text{Na}^+\cdot\text{C}_9\text{H}_3\text{NO}_8^{2-}\cdot 3\text{H}_2\text{O}$ $M_r = 353.15$ Triclinic, $P\bar{1}$ $a = 5.5844$ (11) Å $b = 6.6770$ (13) Å $c = 18.631$ (4) Å $\alpha = 81.34$ (3)° $\beta = 86.77$ (3)° $\gamma = 68.36$ (3)° $V = 638.4$ (2) Å³ $Z = 2$ Mo $K\alpha$ radiation $\mu = 0.23$ mm⁻¹ $T = 293$ K

0.1 × 0.1 × 0.1 mm

Data collection

Rigaku R-AXIS RAPID diffractometer
Absorption correction: multi-scan (*ABSCOR*; Higashi, 1995)
 $T_{\min} = 0.97$, $T_{\max} = 0.98$

5075 measured reflections
2247 independent reflections
1780 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.018$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.139$
 $S = 1.14$
2247 reflections

208 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.55$ e Å⁻³
 $\Delta\rho_{\min} = -0.33$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O}3-\text{H}3\cdots\text{O}2$	0.86	1.55	2.410	175
$\text{O}7-\text{H}7\cdots\text{O}6$	0.86	1.55	2.402	175
$\text{O}9-\text{H}9\text{A}\cdots\text{O}1^{\text{i}}$	0.85	2.05	2.900	175
$\text{O}9-\text{H}9\text{B}\cdots\text{O}1^{\text{ii}}$	0.86	2.11	2.945	161
$\text{O}10-\text{H}10\text{A}\cdots\text{O}2^{\text{iii}}$	0.88	2.21	3.024	153
$\text{O}10-\text{H}10\text{B}\cdots\text{O}8^{\text{iv}}$	0.88	1.86	2.738	173
$\text{O}11-\text{H}11\text{A}\cdots\text{O}4^{\text{v}}$	0.89	2.05	2.917	167
$\text{O}11-\text{H}11\text{B}\cdots\text{O}4^{\text{v}}$	0.88	2.08	2.949	170

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

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2004); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: DS2024).

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supporting information

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Disodium dihydrogen pyridine-2,3,5,6-tetracarboxylate trihydrate

Fang-Hong Hu and Jian-Li Lin

S1. Comment

A great many metal-polycarboxylate compounds with benzene-1,2,4,5-tetracarboxylic acid (H_4btec) as ligands have been designed and characterized (Zhang *et al.*, 2010). Analogous in structure to H_4btec , only four complexes $[Ni(H_2pdtc)(H_2O)_2].3H_2O$ (Yang *et al.*, 2008), $[Zn_4(pdtc)_2(phen)_2(H_2O)_2].20H_2O$ (Yang *et al.*, 2008), $[Cd(H_2pdtc)(H_2O)_3].3H_2O$ (Sun, Zhou & Yan, 2009) and $[Ni_2(pdtc)(H_2O)_3(2,2'-bpy)].4H_2O$ (Sun, Zhou & An, 2009) about pyridine-2,3,5,6-tetracarboxylic acid (H_4pdtc) have been reported. The four coordination polymers are constructed by linking transition metal centres through the ligands. In this context, we represent a disodium salt of dihydrogen pyridine-2,3,5,6-tetracarboxylate dianion $[Na_2(H_2pdtc)(H_2O)_3]$.

The asymmetric unit of the title compound consists of two crystallographically independent Na cations, one dihydrogen pyridine-2,3,5,6-tetracarboxylate dianion (H_2pdtc^{2-}) and three water molecules (Fig. 1). The ligand is deprotonated at 2,5-positioned carboxylate groups. The values of the dihedral angle between the planes of the carboxylic groups and the planar pyridine ring are 3.7 (4)°, 2.0 (5)°, 10.5 (4)°, 2.4 (5)°, respectively. Both Na1 and Na2 ions are seven-coordinated with one N atom, two carboxylate O atoms and four water molecules at Na1 and five carboxylate O atoms and two water molecules at Na2, building highly distorted pentagonal-bipyramidal environment. The average value of the Na—O(water) length [2.494 (2) Å] is closer to the average Na—O (carboxylate) distance [2.498 (2) Å], and both are slightly smaller than the Na—N value [2.654 (2) Å]. The sodium cations are bridged by the (H_2pdtc^{2-}) to generate layers extending infinitely in sheets parallel to (001) (Fig. 2), and further pillared by the water molecule linkers (Fig. 3) to build up 3D framework, which is found to be stabilized by hydrogen bonds from water molecules to carboxylate O atoms (Table 1).

S2. Experimental

0.0766 g (0.3 mmol) pyridine-2,3,5,6-tetracarboxylic acid and 0.024 g (0.6 mmol) NaOH were successively added to 10.0 ml H_2O and stirred at room temperature for 2 h, and the resulting colorless solution (pH = 3.48) was then transferred to a 50 ml beaker for slow evaporation at room temperature for several months, affording colorless block crystals (yield: 0.05 g).

S3. Refinement

H atoms bonded to C atoms were placed in geometrically calculated positions and were refined using a riding model, with $U_{iso}(H) = 1.2U_{eq}(C)$. H atoms attached to O atoms were found in a difference Fourier synthesis and were refined using a riding model, with $U_{iso}(H)$ values set at $1.2U_{eq}(O)$.

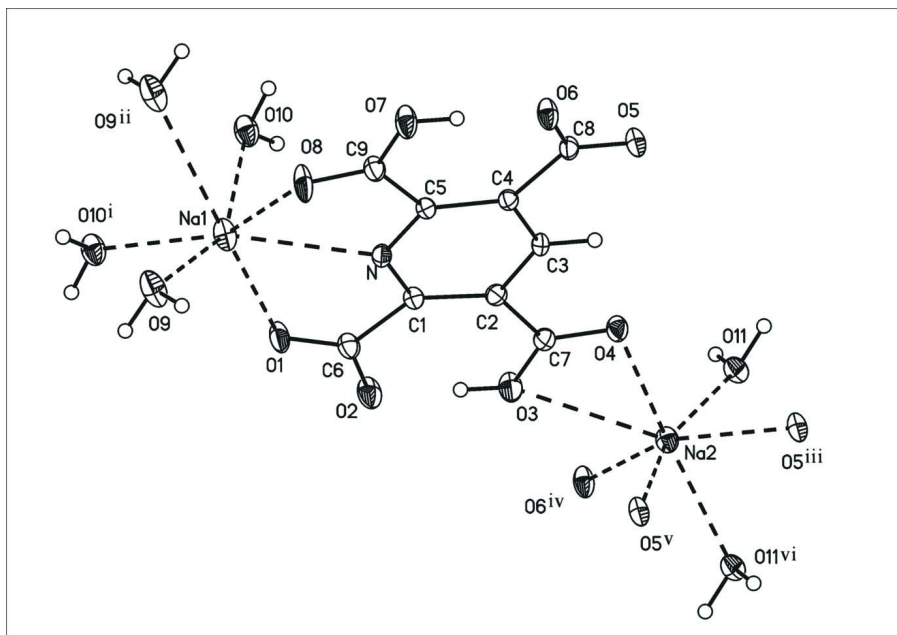


Figure 1

ORTEP view of the title compound, Displacement ellipsoids are drawn at the 30% probability level. [Symmetry codes: (i) $2 - x, 1 - y, 1 - z$; (ii) $3 - x, 1 - y, 1 - z$; (iii) $2 - x, 1 - y, -z$; (iv) $x - 1, y - 1, z$; (v) $x, y - 1, z$; (vi) $1 - x, -y, -z$.]

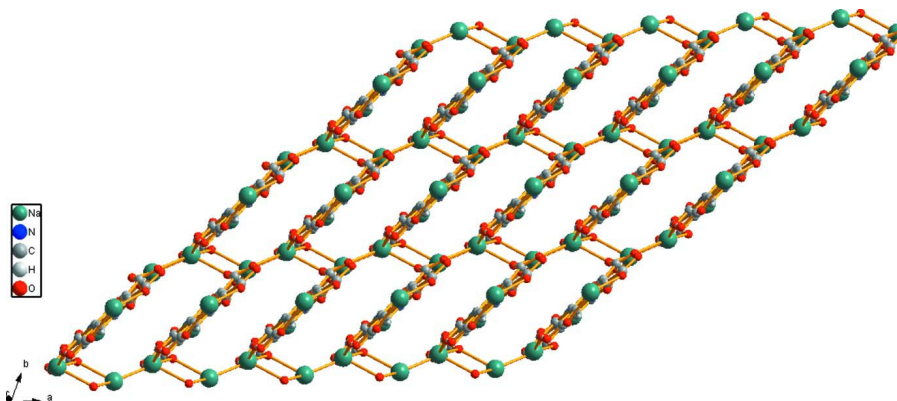


Figure 2

2D layer $\infty[\text{Na}_2(\text{H}_2\text{pdtc})]$ parallel to (001) generated from bridging sodium cations and $(\text{H}_2\text{pdtc})^{2-}$.

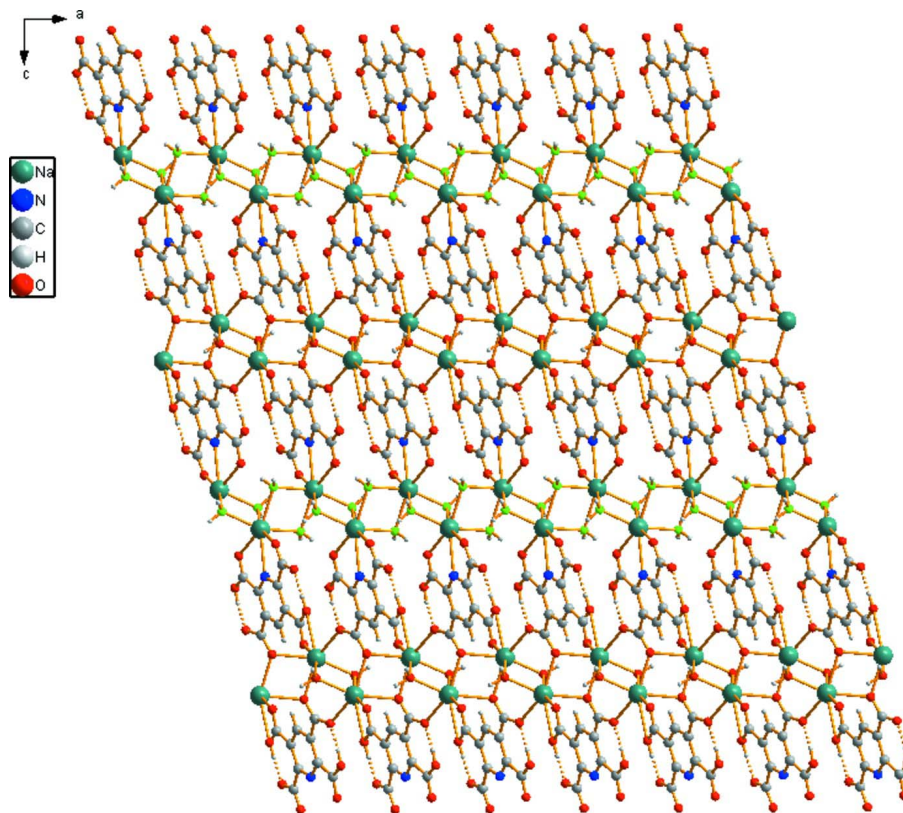


Figure 3

3D framework built from linking $\infty[\text{Na}_2(\text{H}_2\text{pdtc})]$ layers by the water molecules (the water molecule linkers are marked in green)

Disodium dihydrogen pyridine-2,3,5,6-tetracarboxylate trihydrate

Crystal data

$2\text{Na}^+\cdot\text{C}_9\text{H}_3\text{NO}_8^{2-}\cdot 3\text{H}_2\text{O}$

$M_r = 353.15$

Triclinic, $P\bar{1}$

Hall symbol: $-P\ 1$

$a = 5.5844$ (11) Å

$b = 6.6770$ (13) Å

$c = 18.631$ (4) Å

$\alpha = 81.34$ (3)°

$\beta = 86.77$ (3)°

$\gamma = 68.36$ (3)°

$V = 638.4$ (2) Å³

$Z = 2$

$F(000) = 360$

$D_x = 1.837$ Mg m⁻³

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

Cell parameters from 4730 reflections

$\theta = 3.3\text{--}27.5^\circ$

$\mu = 0.23$ mm⁻¹

$T = 293$ K

Block, colourless

$0.1 \times 0.1 \times 0.1$ mm

Data collection

Rigaku R-AXIS RAPID
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 0 pixels mm⁻¹

ω scans

Absorption correction: multi-scan
(*ABSCOR*; Higashi, 1995)

$T_{\min} = 0.97$, $T_{\max} = 0.98$

5075 measured reflections

2247 independent reflections

1780 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.018$

$\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 3.3^\circ$
 $h = -6 \rightarrow 6$

$k = -7 \rightarrow 7$
 $l = -22 \rightarrow 22$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.139$
 $S = 1.14$
 2247 reflections
 208 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0916P)^2 + 0.0526P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.55 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -0.33 \text{ e } \text{Å}^{-3}$

Special details

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 $F^2 > \sigma(F^2)$ 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 (Å^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Na1	1.28244 (19)	0.41119 (18)	0.44136 (5)	0.0360 (3)
Na2	0.70103 (18)	0.05082 (15)	0.05556 (5)	0.0275 (3)
N	1.2379 (4)	0.4267 (3)	0.29938 (10)	0.0234 (5)
C1	1.1039 (4)	0.3240 (4)	0.27451 (12)	0.0222 (5)
C2	1.0250 (4)	0.3696 (4)	0.20111 (12)	0.0227 (5)
C3	1.0921 (4)	0.5287 (4)	0.15822 (12)	0.0225 (5)
H3A	1.0392	0.5653	0.1100	0.027*
C4	1.2339 (4)	0.6372 (4)	0.18300 (12)	0.0205 (5)
C5	1.3080 (4)	0.5774 (4)	0.25638 (12)	0.0213 (5)
C6	1.0527 (5)	0.1626 (4)	0.33584 (13)	0.0280 (6)
O1	1.1267 (4)	0.1553 (3)	0.39697 (9)	0.0364 (5)
O2	0.9322 (4)	0.0429 (3)	0.32087 (10)	0.0408 (5)
C7	0.8762 (5)	0.2689 (4)	0.16184 (13)	0.0252 (5)
O3	0.7999 (4)	0.1219 (3)	0.19559 (9)	0.0361 (5)
H3	0.8543	0.0954	0.2396	0.043*
O4	0.8325 (3)	0.3309 (3)	0.09639 (9)	0.0299 (4)
C8	1.2832 (4)	0.8074 (4)	0.12553 (12)	0.0231 (5)
O5	1.1680 (3)	0.8518 (3)	0.06780 (9)	0.0295 (4)
O6	1.4448 (3)	0.8925 (3)	0.13981 (9)	0.0335 (5)
C9	1.4680 (5)	0.6635 (4)	0.29836 (13)	0.0270 (6)
O7	1.5562 (4)	0.8069 (3)	0.26636 (10)	0.0385 (5)
H7	1.5260	0.8359	0.2205	0.046*
O8	1.5129 (4)	0.5911 (3)	0.36238 (9)	0.0437 (5)

O9	1.6891 (4)	0.2384 (3)	0.49492 (11)	0.0466 (5)
H9A	1.8156	0.2081	0.4655	0.056*
H9B	1.7251	0.1117	0.5202	0.056*
O10	0.8582 (4)	0.6942 (3)	0.43369 (10)	0.0363 (5)
H10A	0.8616	0.8205	0.4131	0.044*
H10B	0.7414	0.6720	0.4094	0.044*
O11	0.2943 (3)	0.2647 (3)	0.00399 (9)	0.0304 (4)
H11A	0.2605	0.3963	-0.0208	0.037*
H11B	0.1455	0.2949	0.0272	0.037*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Na1	0.0340 (6)	0.0478 (6)	0.0265 (6)	-0.0181 (5)	-0.0043 (4)	0.0044 (4)
Na2	0.0285 (5)	0.0318 (5)	0.0256 (5)	-0.0155 (4)	-0.0043 (4)	-0.0008 (4)
N	0.0272 (10)	0.0255 (10)	0.0180 (10)	-0.0103 (9)	-0.0006 (8)	-0.0027 (8)
C1	0.0233 (11)	0.0233 (12)	0.0200 (12)	-0.0090 (10)	-0.0002 (9)	-0.0012 (9)
C2	0.0238 (11)	0.0224 (11)	0.0227 (12)	-0.0092 (10)	-0.0015 (9)	-0.0036 (9)
C3	0.0277 (12)	0.0245 (12)	0.0150 (11)	-0.0092 (10)	-0.0040 (9)	-0.0018 (9)
C4	0.0223 (11)	0.0206 (11)	0.0190 (11)	-0.0087 (10)	0.0001 (9)	-0.0020 (9)
C5	0.0239 (11)	0.0210 (11)	0.0189 (11)	-0.0087 (10)	0.0008 (9)	-0.0012 (9)
C6	0.0314 (13)	0.0300 (13)	0.0237 (13)	-0.0146 (11)	0.0001 (10)	0.0013 (10)
O1	0.0510 (12)	0.0438 (11)	0.0195 (10)	-0.0264 (10)	-0.0049 (8)	0.0057 (8)
O2	0.0608 (13)	0.0453 (11)	0.0292 (10)	-0.0379 (11)	-0.0054 (9)	0.0056 (8)
C7	0.0299 (13)	0.0243 (12)	0.0228 (13)	-0.0113 (11)	-0.0018 (10)	-0.0031 (10)
O3	0.0517 (12)	0.0412 (10)	0.0275 (10)	-0.0324 (10)	-0.0060 (8)	0.0008 (8)
O4	0.0415 (10)	0.0320 (9)	0.0227 (9)	-0.0210 (8)	-0.0068 (7)	-0.0009 (7)
C8	0.0273 (12)	0.0261 (12)	0.0169 (11)	-0.0116 (10)	0.0006 (9)	-0.0017 (9)
O5	0.0327 (9)	0.0376 (10)	0.0201 (9)	-0.0176 (8)	-0.0049 (7)	0.0046 (7)
O6	0.0455 (11)	0.0461 (11)	0.0213 (9)	-0.0337 (10)	-0.0057 (8)	0.0040 (8)
C9	0.0331 (13)	0.0292 (13)	0.0226 (13)	-0.0171 (11)	-0.0036 (10)	0.0007 (10)
O7	0.0557 (12)	0.0519 (12)	0.0234 (9)	-0.0394 (11)	-0.0088 (8)	0.0027 (8)
O8	0.0642 (14)	0.0651 (13)	0.0189 (10)	-0.0469 (12)	-0.0146 (9)	0.0090 (9)
O9	0.0342 (11)	0.0582 (13)	0.0355 (11)	-0.0105 (10)	0.0001 (9)	0.0132 (10)
O10	0.0387 (10)	0.0452 (11)	0.0304 (10)	-0.0236 (9)	-0.0077 (8)	0.0033 (8)
O11	0.0286 (9)	0.0302 (9)	0.0296 (9)	-0.0098 (8)	0.0007 (7)	0.0019 (7)

Geometric parameters (Å, °)

Na1—O9	2.337 (2)	C3—H3A	0.9300
Na1—O8	2.379 (2)	C4—C5	1.406 (3)
Na1—O10	2.420 (2)	C4—C8	1.531 (3)
Na1—O1	2.442 (2)	C5—C9	1.528 (3)
Na1—O10 ⁱ	2.498 (2)	C6—O1	1.221 (3)
Na1—N	2.654 (2)	C6—O2	1.287 (3)
Na1—O9 ⁱⁱ	2.835 (3)	C7—O4	1.235 (3)
Na1—Na1 ⁱ	3.657 (2)	C7—O3	1.278 (3)
Na1—Na1 ⁱⁱ	3.948 (2)	O3—H3	0.8603

Na2—O11	2.358 (2)	C8—O5	1.225 (3)
Na2—O5 ⁱⁱⁱ	2.4395 (18)	C8—O6	1.288 (3)
Na2—O6 ^{iv}	2.4477 (19)	O5—Na2 ⁱⁱⁱ	2.4395 (18)
Na2—O5 ^v	2.457 (2)	O5—Na2 ^{viii}	2.457 (2)
Na2—O4	2.4738 (18)	O6—Na2 ^{ix}	2.4477 (19)
Na2—O11 ^{vi}	2.516 (2)	C9—O8	1.221 (3)
Na2—O3	2.840 (2)	C9—O7	1.287 (3)
Na2—C7	3.022 (3)	O7—H7	0.8596
Na2—Na2 ^{vi}	3.444 (2)	O9—Na1 ⁱⁱ	2.835 (3)
Na2—Na2 ^{vii}	3.724 (2)	O9—H9A	0.8516
N—C1	1.325 (3)	O9—H9B	0.8625
N—C5	1.349 (3)	O10—Na1 ⁱ	2.498 (2)
C1—C2	1.414 (3)	O10—H10A	0.8790
C1—C6	1.539 (3)	O10—H10B	0.8816
C2—C3	1.382 (3)	O11—Na2 ^{vi}	2.516 (2)
C2—C7	1.519 (3)	O11—H11A	0.8872
C3—C4	1.389 (3)	O11—H11B	0.8833
O9—Na1—O8	81.23 (8)	O11—Na2—Na2 ^{vii}	120.83 (6)
O9—Na1—O10	151.17 (9)	O5 ⁱⁱⁱ —Na2—Na2 ^{vii}	40.67 (5)
O8—Na1—O10	99.80 (8)	O6 ^{iv} —Na2—Na2 ^{vii}	146.74 (7)
O9—Na1—O1	112.23 (9)	O5 ^v —Na2—Na2 ^{vii}	40.31 (4)
O8—Na1—O1	120.07 (7)	O4—Na2—Na2 ^{vii}	83.00 (6)
O10—Na1—O1	92.37 (7)	O11 ^{vi} —Na2—Na2 ^{vii}	79.10 (6)
O9—Na1—O10 ⁱ	81.74 (8)	O3—Na2—Na2 ^{vii}	108.88 (6)
O8—Na1—O10 ⁱ	150.12 (7)	C7—Na2—Na2 ^{vii}	93.42 (6)
O10—Na1—O10 ⁱ	83.92 (8)	Na2 ^{vi} —Na2—Na2 ^{vii}	102.29 (5)
O1—Na1—O10 ⁱ	89.14 (7)	C1—N—C5	122.15 (19)
O9—Na1—N	119.02 (8)	C1—N—Na1	120.01 (14)
O8—Na1—N	62.02 (6)	C5—N—Na1	116.33 (15)
O10—Na1—N	85.55 (8)	N—C1—C2	121.20 (19)
O1—Na1—N	60.81 (6)	N—C1—C6	110.34 (19)
O10 ⁱ —Na1—N	147.64 (7)	C2—C1—C6	128.5 (2)
O9—Na1—O9 ⁱⁱ	80.93 (8)	C3—C2—C1	116.0 (2)
O8—Na1—O9 ⁱⁱ	70.80 (7)	C3—C2—C7	114.6 (2)
O10—Na1—O9 ⁱⁱ	72.45 (7)	C1—C2—C7	129.44 (19)
O1—Na1—O9 ⁱⁱ	163.26 (7)	C2—C3—C4	123.8 (2)
O10 ⁱ —Na1—O9 ⁱⁱ	82.40 (7)	C2—C3—H3A	118.1
N—Na1—O9 ⁱⁱ	123.02 (7)	C4—C3—H3A	118.1
O9—Na1—Na1 ⁱ	118.79 (7)	C3—C4—C5	116.01 (19)
O8—Na1—Na1 ⁱ	134.81 (8)	C3—C4—C8	114.27 (19)
O10—Na1—Na1 ⁱ	42.77 (5)	C5—C4—C8	129.72 (19)
O1—Na1—Na1 ⁱ	90.98 (6)	N—C5—C4	120.8 (2)
O10 ⁱ —Na1—Na1 ⁱ	41.15 (5)	N—C5—C9	111.11 (19)
N—Na1—Na1 ⁱ	121.70 (7)	C4—C5—C9	128.05 (19)
O9 ⁱⁱ —Na1—Na1 ⁱ	73.15 (6)	O1—C6—O2	122.8 (2)
O9—Na1—Na1 ⁱⁱ	45.17 (6)	O1—C6—C1	118.3 (2)
O8—Na1—Na1 ⁱⁱ	70.94 (5)	O2—C6—C1	118.9 (2)

O10—Na1—Na1 ⁱⁱ	107.58 (7)	C6—O1—Na1	127.72 (16)
O1—Na1—Na1 ⁱⁱ	155.73 (7)	O4—C7—O3	120.7 (2)
O10 ⁱ —Na1—Na1 ⁱⁱ	79.63 (5)	O4—C7—C2	118.50 (19)
N—Na1—Na1 ⁱⁱ	132.72 (6)	O3—C7—C2	120.8 (2)
O9 ⁱⁱ —Na1—Na1 ⁱⁱ	35.76 (5)	O4—C7—Na2	52.54 (11)
Na1 ⁱ —Na1—Na1 ⁱⁱ	94.41 (5)	O3—C7—Na2	69.52 (13)
O11—Na2—O5 ⁱⁱⁱ	80.18 (7)	C2—C7—Na2	164.48 (15)
O11—Na2—O6 ^{iv}	82.71 (7)	C7—O3—Na2	85.55 (14)
O5 ⁱⁱⁱ —Na2—O6 ^{iv}	150.56 (7)	C7—O3—H3	107.5
O11—Na2—O5 ^v	161.10 (7)	Na2—O3—H3	160.1
O5 ⁱⁱⁱ —Na2—O5 ^v	80.98 (7)	C7—O4—Na2	104.12 (14)
O6 ^{iv} —Na2—O5 ^v	113.75 (7)	O5—C8—O6	123.9 (2)
O11—Na2—O4	101.38 (7)	O5—C8—C4	117.35 (19)
O5 ⁱⁱⁱ —Na2—O4	89.54 (6)	O6—C8—C4	118.8 (2)
O6 ^{iv} —Na2—O4	117.34 (7)	C8—O5—Na2 ⁱⁱⁱ	130.76 (15)
O5 ^v —Na2—O4	79.82 (7)	C8—O5—Na2 ^{viii}	123.43 (15)
O11—Na2—O11 ^{vi}	90.16 (7)	Na2 ⁱⁱⁱ —O5—Na2 ^{viii}	99.02 (7)
O5 ⁱⁱⁱ —Na2—O11 ^{vi}	78.64 (7)	C8—O6—Na2 ^{ix}	128.80 (14)
O6 ^{iv} —Na2—O11 ^{vi}	77.63 (6)	O8—C9—O7	121.0 (2)
O5 ^v —Na2—O11 ^{vi}	84.77 (7)	O8—C9—C5	118.5 (2)
O4—Na2—O11 ^{vi}	161.90 (7)	O7—C9—C5	120.5 (2)
O11—Na2—O3	117.19 (7)	C9—O7—H7	112.8
O5 ⁱⁱⁱ —Na2—O3	134.97 (6)	C9—O8—Na1	126.87 (16)
O6 ^{iv} —Na2—O3	74.36 (6)	Na1—O9—Na1 ⁱⁱ	99.07 (8)
O5 ^v —Na2—O3	77.80 (7)	Na1—O9—H9A	115.5
O4—Na2—O3	47.92 (6)	Na1 ⁱⁱ —O9—H9A	104.4
O11 ^{vi} —Na2—O3	137.20 (7)	Na1—O9—H9B	118.8
O11—Na2—C7	113.48 (7)	Na1 ⁱⁱ —O9—H9B	120.3
O5 ⁱⁱⁱ —Na2—C7	110.79 (7)	H9A—O9—H9B	98.8
O6 ^{iv} —Na2—C7	97.96 (7)	Na1—O10—Na1 ⁱ	96.08 (8)
O5 ^v —Na2—C7	74.82 (7)	Na1—O10—H10A	112.2
O4—Na2—C7	23.34 (6)	Na1 ⁱ —O10—H10A	129.4
O11 ^{vi} —Na2—C7	155.41 (7)	Na1—O10—H10B	116.3
O3—Na2—C7	24.93 (6)	Na1 ⁱ —O10—H10B	97.7
O11—Na2—Na2 ^{vi}	46.94 (5)	H10A—O10—H10B	105.2
O5 ⁱⁱⁱ —Na2—Na2 ^{vi}	74.89 (5)	Na2—O11—Na2 ^{vi}	89.84 (7)
O6 ^{iv} —Na2—Na2 ^{vi}	75.92 (5)	Na2—O11—H11A	122.9
O5 ^v —Na2—Na2 ^{vi}	125.55 (7)	Na2 ^{vi} —O11—H11A	121.8
O4—Na2—Na2 ^{vi}	146.06 (7)	Na2—O11—H11B	125.4
O11 ^{vi} —Na2—Na2 ^{vi}	43.23 (5)	Na2 ^{vi} —O11—H11B	100.3
O3—Na2—Na2 ^{vi}	148.10 (6)	H11A—O11—H11B	96.5
C7—Na2—Na2 ^{vi}	159.59 (7)		

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

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
O3—H3 \cdots O2	0.86	1.55	2.410	175
O7—H7 \cdots O6	0.86	1.55	2.402	175
O9—H9 <i>A</i> \cdots O1 ^x	0.85	2.05	2.900	175
O9—H9 <i>B</i> \cdots O1 ^{xi}	0.86	2.11	2.945	161
O10—H10 <i>A</i> \cdots O2 ^{viii}	0.88	2.21	3.024	153
O10—H10 <i>B</i> \cdots O8 ^{xii}	0.88	1.86	2.738	173
O11—H11 <i>A</i> \cdots O4 ^{xiii}	0.89	2.05	2.917	167
O11—H11 <i>B</i> \cdots O4 ^{xii}	0.88	2.08	2.949	170

Symmetry codes: (viii) $x, y+1, z$; (x) $x+1, y, z$; (xi) $-x+3, -y, -z+1$; (xii) $x-1, y, z$; (xiii) $-x+1, -y+1, -z$.