

2-Amino-4-methylpyridinium 6-carboxy-pyridine-2-carboxylate sesquihydrate

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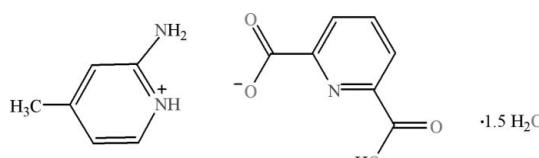
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Key indicators: single-crystal X-ray study; $T = 120\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$; disorder in main residue; R factor = 0.047; wR factor = 0.102; data-to-parameter ratio = 17.6.

In the title compound, $\text{C}_6\text{H}_9\text{N}_2^+\cdot\text{C}_7\text{H}_4\text{NO}_4^-\cdot1.5\text{H}_2\text{O}$, extensive $\text{O}-\text{H}\cdots\text{O}$, $\text{O}-\text{H}\cdots\text{N}$, $\text{N}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds, as well as ion pairing, $\pi-\pi$ stacking interactions [centroid-centroid distances = 3.4690 (8) and 3.6932 (8) \AA between aromatic rings] occur in the crystal. There are hydrogen-bonding interactions between water molecules, which result in cyclic tetrameric water clusters. One of the water O molecules has half occupancy. In the anion molecules, the $-\text{CO}_2$ and $-\text{CO}_2\text{H}$ groups make torsion angles of 1.73 (18) and -12.14 (18) $^\circ$ with respect to the ring.

Related literature

For background to hydrogen bonding involving water, see: Long *et al.* (2004); Atwood *et al.*, (2001); Miyake & Aida (2003). For related structures, see: Aghabozorg *et al.* (2008); Tabatabaei *et al.* (2009).



Experimental

Crystal data



$M_r = 302.29$

Monoclinic, $P2_1/c$

$a = 9.2373$ (6) \AA

$b = 7.1972$ (5) \AA

$c = 21.6495$ (14) \AA

$\beta = 93.951$ (1) $^\circ$

$V = 1435.90$ (17) \AA^3

$Z = 4$

Mo $K\alpha$ radiation

$\mu = 0.11\text{ mm}^{-1}$

$T = 120\text{ K}$

$0.20 \times 0.20 \times 0.10\text{ mm}$

Data collection

Bruker SMART 1000 CCD area detector diffractometer

Absorption correction: multi-scan (*SADABS*; Bruker, 1998)

$T_{\min} = 0.980$, $T_{\max} = 0.995$

15297 measured reflections

3801 independent reflections

3077 reflections with $I > 2.0\sigma(I)$

$R_{\text{int}} = 0.026$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.047$

$wR(F^2) = 0.102$

$S = 0.99$

3801 reflections

216 parameters

H atoms treated by a mixture of independent and constrained refinement

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

$\Delta\rho_{\min} = -0.25\text{ e \AA}^{-3}$

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1W—H1WA \cdots O4 ⁱ	0.85	2.01	2.846 (2)	169
O1W—H1WB \cdots O3	0.85	1.97	2.813 (2)	169
O2W—H2WB \cdots O1W ⁱⁱ	0.85	2.11	2.944 (2)	167
O2W—H2WA \cdots O1W	0.85	2.07	2.919 (2)	175
O1—H1O \cdots O1W ⁱⁱⁱ	0.93 (3)	1.78 (3)	2.661 (2)	155 (2)
N2—H2N \cdots O4	0.96 (2)	1.74 (2)	2.700 (2)	174 (2)
N3—H3NB \cdots O3	0.98 (2)	1.86 (2)	2.829 (2)	172 (2)
N3—H3NA \cdots O2 ^{iv}	0.90 (2)	2.09 (2)	2.955 (2)	160 (2)
C2—H2A \cdots O2 ^v	0.95	2.47	3.158 (2)	129
C9—H9A \cdots O1 ^{iv}	0.95	2.56	3.399 (2)	147
C11—H11A \cdots O2W ^{vi}	0.95	2.55	3.405 (2)	149

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

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT-Plus* (Bruker, 1998); data reduction: *SAINT-Plus*; 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: *SHELXTL* (Sheldrick, 2008).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2352).

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supporting information

Acta Cryst. (2010). E66, o3232 [https://doi.org/10.1107/S1600536810046866]

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S1. Comment

The presence of water is important in establishing H-bonded contributions to the total lattice energy, and is significant in establishing the stability of the hydrated crystal structures (Long *et al.*, 2004). Several water clusters found in organic or metallo-organic crystal hosts have been structurally characterized (Atwood *et al.*, 2001). A detailed understanding of the numerous possible structures and stability of isolated water clusters in diverse surroundings can help us understand the nature of water-water interactions in bulk water or ice. In this paper, we report the synthesis and crystal structure of the title proton transfer system, (I), derived from pyridine-2,6-dicarboxylic acid (pydcH₂) and 2-amino-4-methylpyridine (2a4mp).

In the title compound, the asymmetric unit contains a cation, (2a4mpH)²⁺, an anion, (pydcH)⁻ and 1.5 water molecules (Fig. 1). The bond distances and bond angles in the title compound are in agreement with the corresponding distances and angles reported in some related crystal structures (Aghabozorg *et al.*, (2008); Tabatabaei *et al.*, (2009)). In the crystal structure, the cations and the anions are linked by hydrogen bonds (Tab. 1 and Fig. 2). In the structure, water molecules form cyclic tetrameric water clusters (Tab. 1 and Fig. 3) in the most stable pattern (Miyake & Aida, 2003). The clusters play a bridging role (Fig. 2), linking the adjacent cations and anions *via* hydrogen bonds and contributing to the formation of an extensive supramolecular structure.

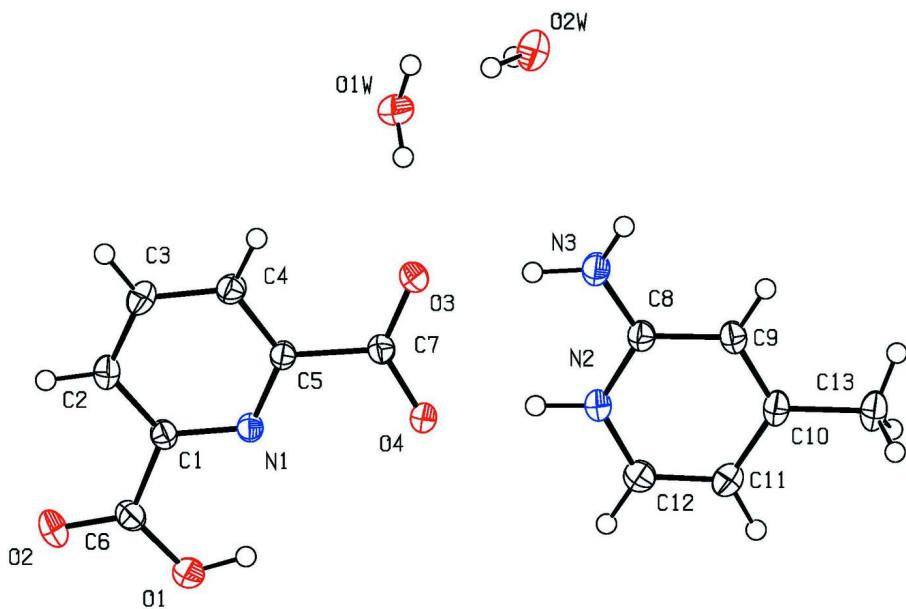
Moreover, $\pi-\pi$ stacking interactions with distances between ring centroids = 3.4690 (8) Å and 3.6931 (8) Å, (Fig. 4) together with C7=O3 \cdots π involving aromatic ring of (pydcH)⁻ (Fig. 5) seem to be effective in stabilizing the crystal structure.

S2. Experimental

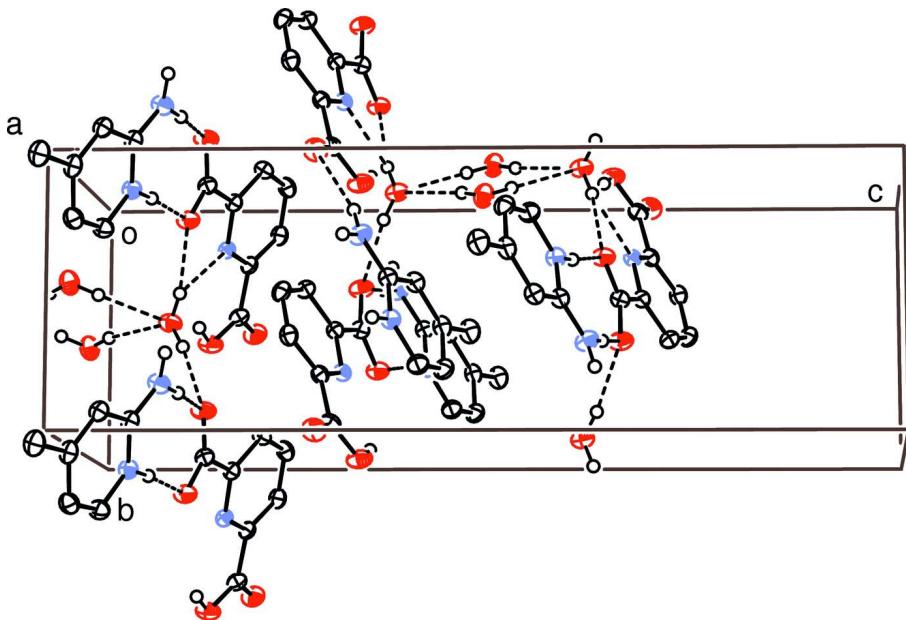
An aqueous solution of 2a4mp (324 mg, 3 mmol) in water (10 ml) was added to a stirring solution of pydcH₂ (501 mg, 3 mmol) in water (10 ml). The reaction mixture was stirred at 298 K for 2 h. Colorless crystals of the title compound were obtained by slow concentration of the solution at room temperature.

S3. Refinement

One of the water molecules (O2W) has 0.5 occupancy factor. The hydrogen atoms of OH, NH and NH₂ groups and water molecules were found in difference Fourier synthesis. The H-atoms of OH, NH and NH₂ groups were refined in isotropic approximation. The rest of the H-atoms were refined in riding model with C–H = 0.95 and 0.98 Å for aryl and methyl H-atoms and O–H = 0.85 Å for water molecules. The $U_{\text{iso}}(\text{H})$ parameters were 1.2 $U_{\text{eq}}(\text{C}_{\text{aryl}}/\text{O})$ and 1.5 $U_{\text{eq}}(\text{C}_{\text{methyl}})$.

**Figure 1**

The asymmetric unit of (I), showing the atom numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

**Figure 2**

A packing diagram of (I) showing hydrogen bonds as dashed lines. Hydrogen atoms not involved in H-bonds have been excluded for clarity.

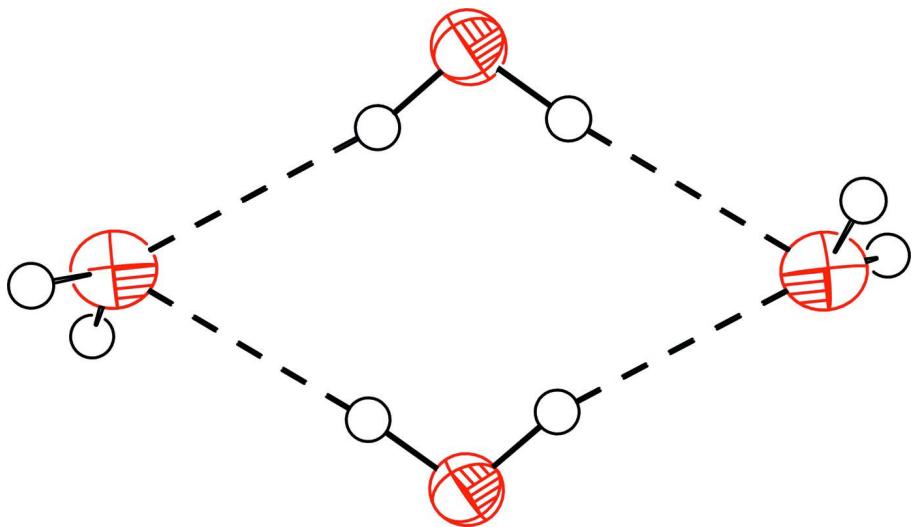


Figure 3

Tetrameric water cluster formed by H-bonds between water molecules in the title compound.

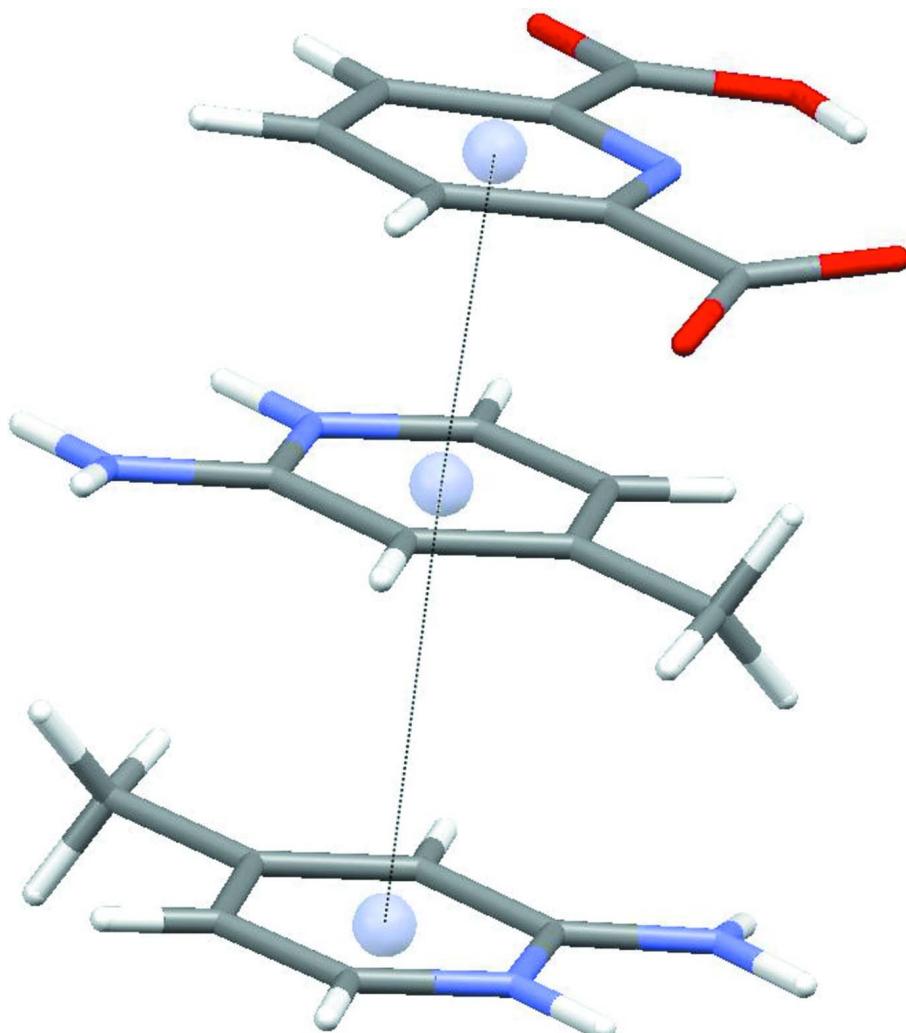
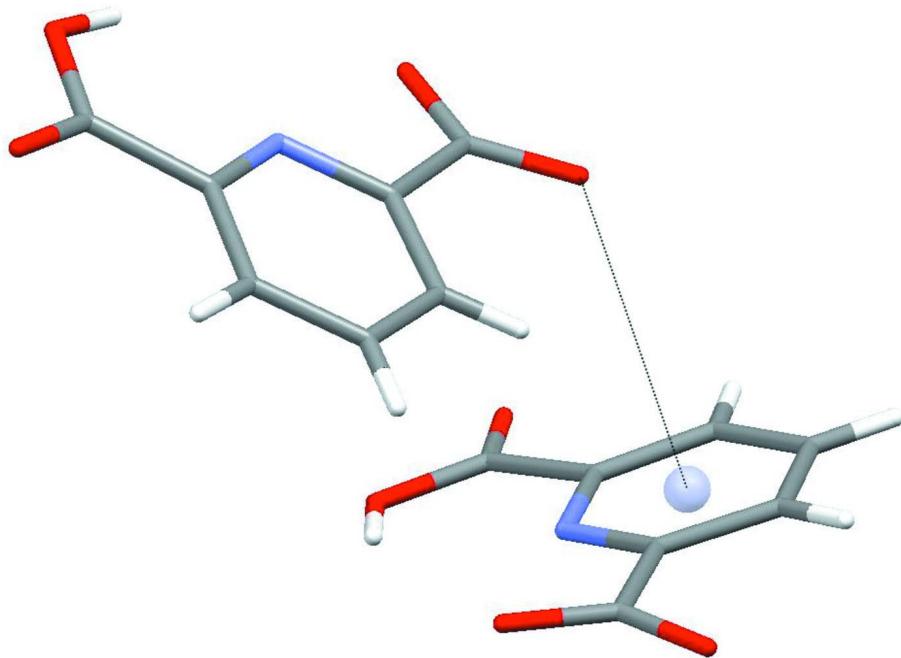


Figure 4

A view of the $\pi\text{-}\pi$ stacking interaction between aromatic rings of the pyridine-2-carboxylate-6-carbonic acid and 2-amino-4-picolinium.

**Figure 5**

A view of the $\text{C}=\text{O}\cdots\pi$ interaction between $\text{C}7=\text{O}3$ group and the centroid of the N1/C1-C5 aromatic ring of the anion, $(\text{pydcH})^-$.

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Crystal data



$M_r = 302.29$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 9.2373 (6)$ Å

$b = 7.1972 (5)$ Å

$c = 21.6495 (14)$ Å

$\beta = 93.951 (1)^\circ$

$V = 1435.90 (17)$ Å³

$Z = 4$

$F(000) = 636$

$D_x = 1.398 \text{ Mg m}^{-3}$

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

Cell parameters from 1125 reflections

$\theta = 2-25^\circ$

$\mu = 0.11 \text{ mm}^{-1}$

$T = 120$ K

Rhombic, colorless

$0.20 \times 0.20 \times 0.10$ mm

Data collection

Bruker SMART 1000 CCD area detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(SADABS; Bruker, 1998)

$T_{\min} = 0.980$, $T_{\max} = 0.995$

15297 measured reflections

3801 independent reflections

3077 reflections with $I > 2.0\sigma(I)$

$R_{\text{int}} = 0.026$

$\theta_{\max} = 29.0^\circ$, $\theta_{\min} = 1.9^\circ$

$h = -12 \rightarrow 12$

$k = -9 \rightarrow 9$

$l = -29 \rightarrow 29$

*Refinement*Refinement on F^2

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.047$$

$$wR(F^2) = 0.102$$

$$S = 0.99$$

3801 reflections

216 parameters

0 restraints

Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map

Hydrogen site location: mixed

H atoms treated by a mixture of independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0244P)^2 + 1.3516P]$$

where $P = (F_o^2 + 2F_c^2)/3$

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

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

$$\Delta\rho_{\min} = -0.25 \text{ e } \text{\AA}^{-3}$$

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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
O1	0.26470 (12)	-0.06479 (14)	0.64545 (5)	0.0309 (2)	
H1O	0.356 (3)	-0.019 (3)	0.6373 (11)	0.066 (7)*	
O2	0.06351 (11)	0.00989 (15)	0.68870 (5)	0.0325 (2)	
O3	0.64297 (10)	0.62489 (14)	0.66239 (5)	0.0288 (2)	
O4	0.65846 (10)	0.32081 (14)	0.64256 (5)	0.0264 (2)	
O1W	0.53707 (11)	0.96510 (15)	0.61389 (5)	0.0305 (2)	
H1WA	0.5713	1.0680	0.6275	0.037*	
H1WB	0.5756	0.8707	0.6320	0.037*	
O2W	0.6530 (2)	1.0929 (3)	0.49926 (10)	0.0310 (5)	0.50
H2WB	0.5892	1.0666	0.4704	0.037*	0.50
H2WA	0.6142	1.0552	0.5315	0.037*	0.50
N1	0.38249 (11)	0.26733 (15)	0.67222 (5)	0.0192 (2)	
N2	0.91543 (12)	0.37469 (16)	0.59293 (5)	0.0223 (2)	
H2N	0.822 (2)	0.364 (3)	0.6099 (9)	0.043 (5)*	
N3	0.93113 (13)	0.67564 (18)	0.62910 (6)	0.0270 (3)	
H3NB	0.835 (2)	0.658 (3)	0.6447 (8)	0.037 (5)*	
H3NA	0.986 (2)	0.776 (3)	0.6395 (9)	0.044 (5)*	
C1	0.25047 (13)	0.23786 (18)	0.69207 (6)	0.0199 (2)	
C2	0.17514 (14)	0.3666 (2)	0.72513 (6)	0.0232 (3)	
H2A	0.0818	0.3388	0.7386	0.028*	
C3	0.23996 (14)	0.5369 (2)	0.73793 (6)	0.0247 (3)	
H3A	0.1922	0.6286	0.7607	0.030*	
C4	0.37662 (14)	0.57143 (19)	0.71688 (6)	0.0223 (3)	
H4A	0.4232	0.6876	0.7247	0.027*	
C5	0.44366 (13)	0.43305 (18)	0.68430 (6)	0.0188 (2)	

C6	0.18551 (15)	0.05222 (19)	0.67584 (7)	0.0246 (3)
C7	0.59368 (14)	0.46266 (18)	0.66116 (6)	0.0210 (3)
C8	0.99015 (14)	0.53539 (19)	0.59952 (6)	0.0213 (3)
C9	1.12726 (14)	0.5468 (2)	0.57398 (6)	0.0225 (3)
H9A	1.1815	0.6589	0.5776	0.027*
C10	1.18187 (14)	0.3971 (2)	0.54415 (6)	0.0236 (3)
C11	1.09980 (15)	0.2309 (2)	0.53908 (7)	0.0262 (3)
H11A	1.1361	0.1252	0.5189	0.031*
C12	0.96820 (15)	0.2250 (2)	0.56356 (7)	0.0260 (3)
H12A	0.9122	0.1143	0.5601	0.031*
C13	1.32713 (15)	0.4067 (2)	0.51713 (7)	0.0304 (3)
H13A	1.3607	0.5359	0.5172	0.046*
H13B	1.3184	0.3600	0.4745	0.046*
H13C	1.3971	0.3305	0.5420	0.046*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0254 (5)	0.0214 (5)	0.0463 (6)	-0.0024 (4)	0.0069 (4)	-0.0036 (4)
O2	0.0241 (5)	0.0321 (6)	0.0420 (6)	-0.0091 (4)	0.0083 (4)	0.0018 (5)
O3	0.0212 (5)	0.0213 (5)	0.0446 (6)	-0.0042 (4)	0.0074 (4)	-0.0023 (4)
O4	0.0197 (4)	0.0216 (5)	0.0390 (6)	0.0005 (4)	0.0097 (4)	-0.0001 (4)
O1W	0.0315 (5)	0.0227 (5)	0.0377 (6)	0.0012 (4)	0.0043 (4)	-0.0029 (4)
O2W	0.0253 (10)	0.0408 (12)	0.0271 (10)	0.0025 (9)	0.0032 (8)	-0.0036 (9)
N1	0.0168 (5)	0.0204 (5)	0.0208 (5)	0.0003 (4)	0.0027 (4)	0.0018 (4)
N2	0.0180 (5)	0.0242 (6)	0.0251 (5)	-0.0012 (4)	0.0040 (4)	0.0003 (4)
N3	0.0209 (6)	0.0264 (6)	0.0345 (7)	-0.0019 (5)	0.0076 (5)	-0.0054 (5)
C1	0.0181 (6)	0.0217 (6)	0.0201 (6)	-0.0010 (5)	0.0022 (4)	0.0027 (5)
C2	0.0188 (6)	0.0295 (7)	0.0219 (6)	0.0004 (5)	0.0046 (5)	0.0015 (5)
C3	0.0220 (6)	0.0287 (7)	0.0236 (6)	0.0040 (5)	0.0039 (5)	-0.0045 (5)
C4	0.0202 (6)	0.0216 (6)	0.0249 (6)	0.0000 (5)	0.0011 (5)	-0.0032 (5)
C5	0.0163 (5)	0.0207 (6)	0.0196 (6)	0.0008 (5)	0.0026 (4)	0.0010 (5)
C6	0.0235 (6)	0.0224 (6)	0.0280 (7)	-0.0027 (5)	0.0030 (5)	0.0041 (5)
C7	0.0177 (6)	0.0212 (6)	0.0242 (6)	-0.0007 (5)	0.0022 (5)	0.0015 (5)
C8	0.0182 (6)	0.0239 (6)	0.0219 (6)	0.0003 (5)	0.0019 (5)	0.0011 (5)
C9	0.0175 (6)	0.0263 (7)	0.0239 (6)	-0.0020 (5)	0.0025 (5)	0.0022 (5)
C10	0.0169 (6)	0.0325 (7)	0.0217 (6)	0.0024 (5)	0.0032 (5)	0.0034 (5)
C11	0.0248 (6)	0.0275 (7)	0.0266 (7)	0.0038 (5)	0.0038 (5)	-0.0028 (5)
C12	0.0248 (6)	0.0244 (7)	0.0288 (7)	-0.0012 (5)	0.0019 (5)	-0.0018 (5)
C13	0.0207 (6)	0.0409 (8)	0.0304 (7)	0.0028 (6)	0.0082 (5)	0.0023 (6)

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

O1—C6	1.3204 (17)	C2—C3	1.384 (2)
O1—H1O	0.93 (2)	C2—H2A	0.9500
O2—C6	1.2179 (17)	C3—C4	1.3937 (18)
O3—C7	1.2528 (16)	C3—H3A	0.9500
O4—C7	1.2632 (16)	C4—C5	1.3904 (18)

O1W—H1WA	0.8500	C4—H4A	0.9500
O1W—H1WB	0.8500	C5—C7	1.5206 (17)
O2W—H2WB	0.8500	C8—C9	1.4188 (17)
O2W—H2WA	0.8501	C9—C10	1.3701 (19)
N1—C1	1.3375 (16)	C9—H9A	0.9500
N1—C5	1.3379 (17)	C10—C11	1.416 (2)
N2—C8	1.3494 (17)	C10—C13	1.5019 (18)
N2—C12	1.3583 (18)	C11—C12	1.3598 (19)
N2—H2N	0.96 (2)	C11—H11A	0.9500
N3—C8	1.3320 (18)	C12—H12A	0.9500
N3—H3NB	0.978 (19)	C13—H13A	0.9800
N3—H3NA	0.90 (2)	C13—H13B	0.9800
C1—C2	1.3872 (18)	C13—H13C	0.9800
C1—C6	1.4966 (19)		
C6—O1—H1O	114.2 (15)	O2—C6—C1	122.14 (13)
H1WA—O1W—H1WB	113.8	O1—C6—C1	117.35 (12)
H2WB—O2W—H2WA	102.8	O3—C7—O4	125.49 (12)
C1—N1—C5	117.44 (11)	O3—C7—C5	117.50 (12)
C8—N2—C12	122.13 (12)	O4—C7—C5	117.01 (11)
C8—N2—H2N	119.4 (12)	N3—C8—N2	118.49 (12)
C12—N2—H2N	118.5 (12)	N3—C8—C9	123.32 (13)
C8—N3—H3NB	118.6 (11)	N2—C8—C9	118.18 (12)
C8—N3—H3NA	119.0 (13)	C10—C9—C8	120.37 (13)
H3NB—N3—H3NA	121.7 (17)	C10—C9—H9A	119.8
N1—C1—C2	124.10 (12)	C8—C9—H9A	119.8
N1—C1—C6	115.21 (12)	C9—C10—C11	119.22 (12)
C2—C1—C6	120.69 (12)	C9—C10—C13	121.01 (13)
C3—C2—C1	118.01 (12)	C11—C10—C13	119.77 (13)
C3—C2—H2A	121.0	C12—C11—C10	118.97 (13)
C1—C2—H2A	121.0	C12—C11—H11A	120.5
C2—C3—C4	118.79 (12)	C10—C11—H11A	120.5
C2—C3—H3A	120.6	N2—C12—C11	121.12 (13)
C4—C3—H3A	120.6	N2—C12—H12A	119.4
C5—C4—C3	118.91 (13)	C11—C12—H12A	119.4
C5—C4—H4A	120.5	C10—C13—H13A	109.5
C3—C4—H4A	120.5	C10—C13—H13B	109.5
N1—C5—C4	122.74 (11)	H13A—C13—H13B	109.5
N1—C5—C7	116.28 (11)	C10—C13—H13C	109.5
C4—C5—C7	120.97 (12)	H13A—C13—H13C	109.5
O2—C6—O1	120.49 (13)	H13B—C13—H13C	109.5
C5—N1—C1—C2	-1.24 (19)	N1—C5—C7—O3	168.20 (12)
C5—N1—C1—C6	178.42 (11)	C4—C5—C7—O3	-12.68 (19)
N1—C1—C2—C3	0.6 (2)	N1—C5—C7—O4	-12.13 (17)
C6—C1—C2—C3	-179.05 (12)	C4—C5—C7—O4	166.98 (12)
C1—C2—C3—C4	0.4 (2)	C12—N2—C8—N3	-179.79 (13)
C2—C3—C4—C5	-0.6 (2)	C12—N2—C8—C9	0.82 (19)

C1—N1—C5—C4	0.94 (18)	N3—C8—C9—C10	179.92 (13)
C1—N1—C5—C7	-179.97 (11)	N2—C8—C9—C10	-0.72 (19)
C3—C4—C5—N1	0.0 (2)	C8—C9—C10—C11	0.0 (2)
C3—C4—C5—C7	-179.08 (12)	C8—C9—C10—C13	-179.71 (13)
N1—C1—C6—O2	-176.83 (13)	C9—C10—C11—C12	0.5 (2)
C2—C1—C6—O2	2.8 (2)	C13—C10—C11—C12	-179.69 (13)
N1—C1—C6—O1	1.73 (18)	C8—N2—C12—C11	-0.2 (2)
C2—C1—C6—O1	-178.60 (12)	C10—C11—C12—N2	-0.5 (2)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
O1W—H1WA···O4 ⁱ	0.85	2.01	2.846 (2)	169
O1W—H1WA···N1 ⁱ	0.85	2.50	2.935 (2)	112
O1W—H1WB···O3	0.85	1.97	2.813 (2)	169
O2W—H2WB···O1W ⁱⁱ	0.85	2.11	2.944 (2)	167
O2W—H2WA···O1W	0.85	2.07	2.919 (2)	175
O1—H1O···O1W ⁱⁱⁱ	0.93 (3)	1.78 (3)	2.661 (2)	155 (2)
O1—H1O···N1	0.93 (3)	2.20 (2)	2.673 (2)	110 (2)
N2—H2N···O4	0.96 (2)	1.74 (2)	2.700 (2)	174 (2)
N3—H3NB···O3	0.98 (2)	1.86 (2)	2.829 (2)	172 (2)
N3—H3NA···O2 ^{iv}	0.90 (2)	2.09 (2)	2.955 (2)	160 (2)
C2—H2A···O2 ^v	0.95	2.47	3.158 (2)	129
C9—H9A···O1 ^{iv}	0.95	2.56	3.399 (2)	147
C11—H11A···O2W ^{vi}	0.95	2.55	3.405 (2)	149
Cg(1) _{N2/C8—C12} ···Cg(1) ^{vi}			3.4690 (8)	
Cg(1)···Cg(2) _{N1/C1—C5} ^{vii}			3.6932 (8)	
C7—O3···Cg(2) ^{viii}		3.50 (1)		125 (1)

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