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Poly[diaqua(μ_5-pyridine-3,5-dicarboxylato)strontium]

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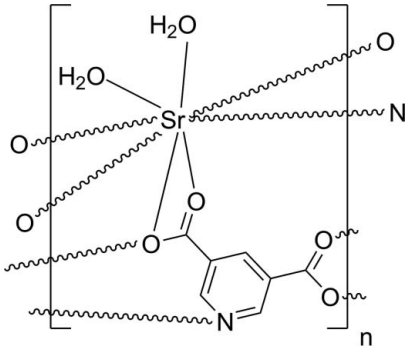
Key indicators: single-crystal X-ray study; T = 293 K; mean σ(C–C) = 0.002 Å; R factor = 0.017; wR factor = 0.046; data-to-parameter ratio = 16.9.

In the structure of the title compound, $[Sr(C_7H_3NO_4)(H_2O)_2]_n$, the Sr^{II} cation is eight-coordinated in form of a distorted dodecahedron by two water O atoms and by five O atoms and one N atom from five pyridine-3,5-dicarboxylate anions. The bridging mode of the anions leads to the formation of a layered network parallel to (100). O—H···O hydrogen bonding between the coordinating water molecules and the carboxylate groups of adjacent layers consolidates the crystal packing. Weak C—H···O interactions are also observed.

Related literature

For related structures with pyridine-3,5-dicarboxylato ligands, see: Aghabozorg *et al.* (2008); Dang *et al.* (2010); Du *et al.* (2009); Lv *et al.* (2010); Wu *et al.* (2008); Yao *et al.* (2010).



Experimental

Crystal data

$[Sr(C_7H_3NO_4)(H_2O)_2]$
M_r = 288.76
Triclinic, $P\overline{1}$
a = 7.9098 (4) Å
b = 8.0028 (4) Å
c = 8.0864 (5) Å
α = 88.620 (2)°
β = 71.270 (2)°
γ = 72.030 (2)°
V = 459.52 (4) Å^3
Z = 2
Mo $K\alpha$ radiation
μ = 5.88 mm^{-1}
T = 293 K
0.30 × 0.17 × 0.16 mm

Data collection

Bruker APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2005)
T_{min} = 0.272, T_{max} = 0.453
8322 measured reflections
2305 independent reflections
2226 reflections with $I > 2\sigma(I)$
R_{int} = 0.022

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.017
$wR(F^2)$ = 0.046
S = 1.08
2305 reflections
136 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.38 e Å^{-3}
$\Delta\rho_{min}$ = −0.51 e Å^{-3}

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
O5—H5A···O4^{i}	0.85	2.01	2.8046 (17)	155
O5—H5B···O3^{ii}	0.85	2.05	2.8718 (18)	163
O6—H6A···O2^{iii}	0.85	1.87	2.7135 (17)	172
O6—H6B···O5^{iv}	0.86	2.03	2.848 (2)	160
C1—H1···O3^{ii}	0.93	2.37	3.286 (2)	169

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

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 2005); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *XP* in *SHELXTL* (Sheldrick, 2008) and *DIAMOND* (Brandenburg, 2012); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: WM2630).

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supporting information

Acta Cryst. (2012). E**68**, m835 [doi:10.1107/S1600536812023379]

Poly[diaqua(μ_5-pyridine-3,5-dicarboxylato)strontium]

Dan Li and Chaowen Duan

S1. Comment

Pyridine-3,5-dicarboxylic acid (py-3,5-dc) is an interesting ligand because it can act in a multidentate fashion and also can participate in hydrogen bonding interactions with its N and O acceptors. Some polymeric complexes of this ligand with 3*d* metals (Dang *et al.*, 2010; Lv *et al.*, 2010; Du *et al.*, 2009) and with mixed 3*d*-4*f* metals (Yao *et al.*, 2010; Wu *et al.*, 2008) have been published. A related strontium complex with a chain-structure has been reported by Aghabozorg *et al.* (2008). Here we report the layered structure of another polymeric strontium complex, $[Sr(C_7H_3NO_4)(H_2O)_2]_n$, (I).

The coordination number of the cation is eight, with bonds to seven O and one N atoms from five py-3,5-dc anions (Fig. 1). The corresponding geometry is distorted dodecahedral (Fig. 2). The Sr—O distances are in the range of 2.5287 (13)–2.7232 (12) Å, and the Sr—N bond is the longest of the coordination polyhedron with a length of 2.8064 (14) Å. The carboxylate O1 atom links two cations to form a rhombic binuclear Sr_2O_2 unit. These units are further connected by carboxylate groups and N atoms of symmetry-related ligands to form a two-dimensional network parallel to (100). Atoms O2, O3 and O4 from these ligands are acceptor atoms for hydrogen bonding with coordinating water molecules of adjacent ligands as donor atoms, leading to a three-dimensional set-up of the structure (Table 1 and Fig. 3). Weak C—H···O interactions are also observed.

S2. Experimental

A mixture of 3,5-H_2pdc (0.0330 g, 0.2 mmol), $SrCl_2{\cdot}6H_2O$ (0.0539 g, 0.2 mmol), imidazole (0.0320 g, 0.47 mmol), C_2H_5OH / H_2O = 1: 2 (3 ml) was placed in a Pyrex-tube (8 ml). The tube was heated for 4 days at 393 K under autogenous pressure. Colourless rod-shaped crystals were obtained. Anal. Calc. for $C_7H_7NO_6Sr$: C, 29.12; N, 4.85; H, 2.44. Found: C, 28.65; N, 4.60; H, 2.35%. IR (KBr, cm^{-1}): 3490 br, 1610 s, 1552 s, 1455 s, 1421 s, 1385 s, 1311 s, 1132 s, 1030 s, 941 s, 827 s, 782 s, 739 s, 651 s, 570 s, 439 s.

S3. Refinement

All hydrogen atoms were located in difference Fourier maps. The water H-atoms were restrained to bond lengths of O—H = 0.85–0.86 Å and with a common U_{iso} parameter of 0.028 $Å^2$ for the H atoms. The C-bound H-atoms were included in calculated positions and treated as riding atoms with C—H = 0.95 Å and U_{iso}(H) = 1.2 U_{eq}(C).

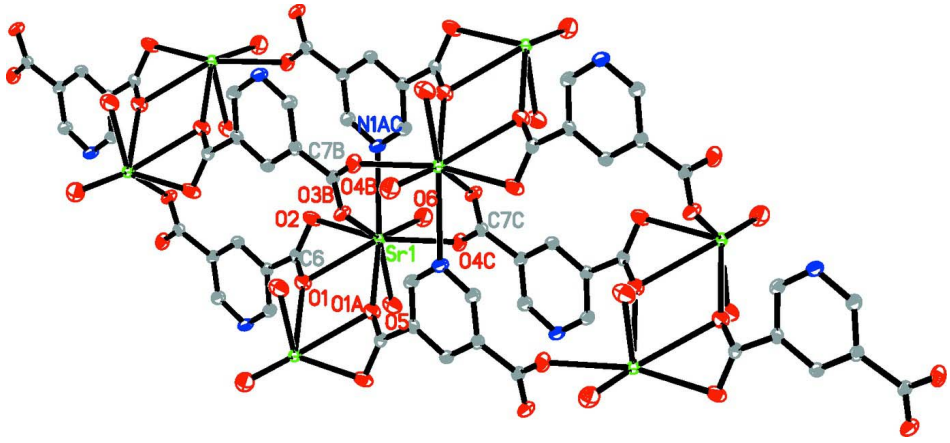

Figure 1

The molecular structure of compound (I), with displacement ellipsoids drawn at the 50% probability level [H-atoms have been omitted for clarity; symmetry codes: (A) -*x* + 1, -*y* + 1, -*z* + 1; (B) -*x* + 1, -*y* + 1, -*z*; (C) *x*, *y* + 1, *z* + 1; (AC) *x*, *y* + 1, *z*.

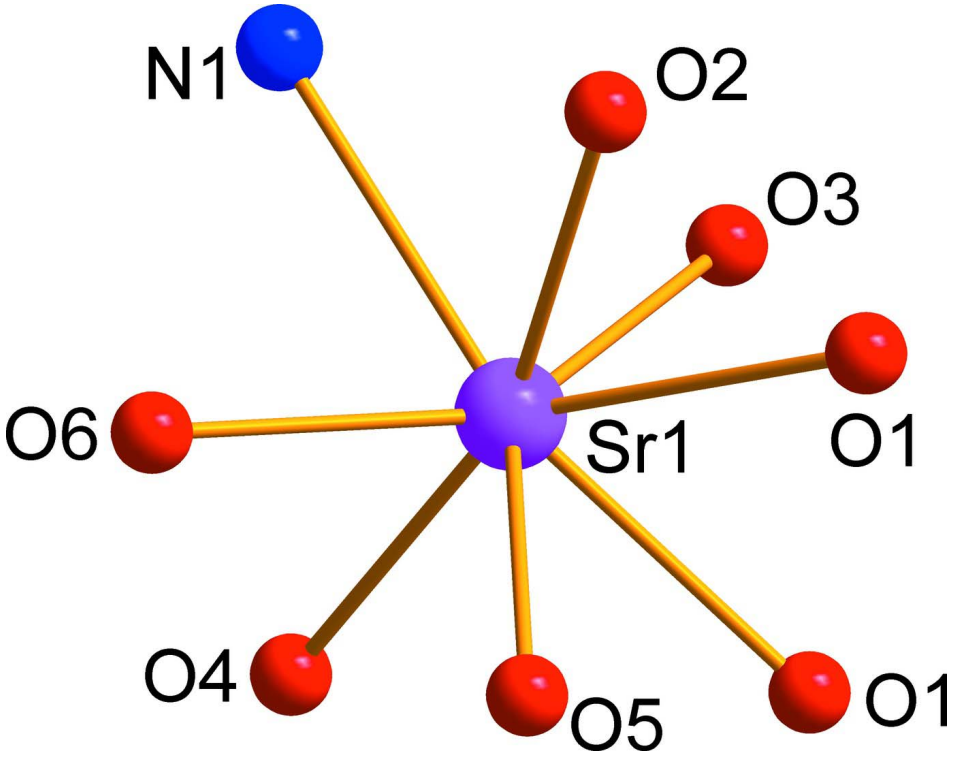

Figure 2

A view of the distorted dodecahedral environment around the Sr^{II} atom in (I).

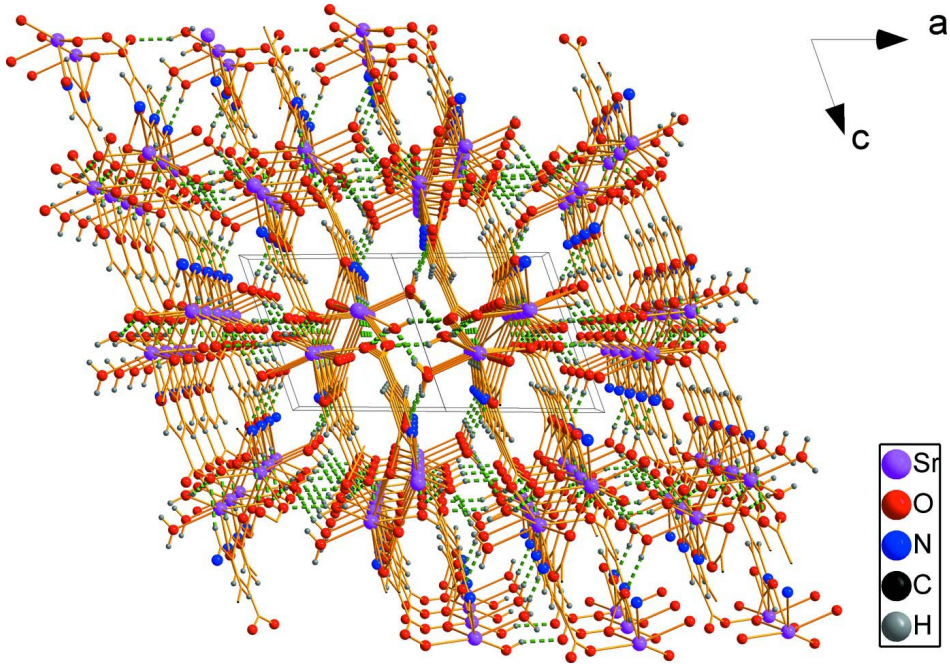

Figure 3

A view down the *b* axis of the crystal packing of compound (I). Hydrogen bonds are shown as dashed lines.

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Crystal data

$[Sr(C_7H_3NO_4)(H_2O)_2]$
$M_r = 288.76$
Triclinic, $P\overline{1}$
Hall symbol: -P 1
$a = 7.9098$ (4) Å
$b = 8.0028$ (4) Å
$c = 8.0864$ (5) Å
$\alpha = 88.620$ (2)°
$\beta = 71.270$ (2)°
$\gamma = 72.030$ (2)°
$V = 459.52$ (4) Å^3
$Z = 2$
$F(000) = 284$
$D_x = 2.087$ Mg m^{-3}
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 3510 reflections
$\theta = 2.7$–$25.3°$
$\mu = 5.88$ mm^{-1}
$T = 293$ K
Block, colourless
0.30 × 0.17 × 0.16 mm

Data collection

Bruker APEXII CCD diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
φ and ω scans
Absorption correction: multi-scan (*SADABS*; Bruker, 2005)
$T_{min} = 0.272$, $T_{max} = 0.453$
8322 measured reflections
2305 independent reflections
2226 reflections with $I > 2\sigma(I)$
$R_{int} = 0.022$
$\theta_{max} = 28.4°$, $\theta_{min} = 2.7°$
$h = -10 \rightarrow 10$
$k = -10 \rightarrow 10$
$l = -9 \rightarrow 10$

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)] = 0.017$
$wR(F^2) = 0.046$
$S = 1.08$
2305 reflections
136 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
$w = 1/[\sigma^2(F_o^2) + (0.0269P)^2 + 0.1887P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} = 0.001$
$\Delta\rho_{max} = 0.38$ e Å^{-3}
$\Delta\rho_{min} = -0.51$ e Å^{-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 (Å²)

	x	*y*	*z*	U_{iso}*/U_{eq}
Sr1	0.685514 (18)	0.654700 (16)	0.373999 (17)	0.01302 (5)
O1	0.53412 (17)	0.40183 (16)	0.33347 (15)	0.0220 (2)
O3	0.63315 (17)	0.12206 (15)	−0.40559 (15)	0.0206 (2)
O2	0.7094 (2)	0.47815 (16)	0.09101 (16)	0.0260 (3)
O4	0.76724 (18)	−0.16816 (16)	−0.41598 (16)	0.0215 (2)
O5	0.85222 (17)	0.35498 (17)	0.47402 (19)	0.0288 (3)
O6	1.03912 (18)	0.5746 (2)	0.23839 (17)	0.0303 (3)
N1	0.7457 (2)	−0.11960 (18)	0.10565 (18)	0.0204 (3)
C7	0.7063 (2)	−0.0153 (2)	−0.34150 (19)	0.0143 (3)
C4	0.7184 (2)	0.0071 (2)	−0.16144 (19)	0.0146 (3)
C3	0.6819 (2)	0.1745 (2)	−0.0860 (2)	0.0150 (3)
H3	0.6607	0.2725	−0.1498	0.018*
C2	0.6775 (2)	0.1931 (2)	0.0855 (2)	0.0148 (3)
C5	0.7511 (2)	−0.1354 (2)	−0.0614 (2)	0.0182 (3)
H5	0.7781	−0.2475	−0.1125	0.022*
C6	0.6365 (2)	0.3698 (2)	0.1760 (2)	0.0155 (3)
C1	0.7093 (2)	0.0429 (2)	0.1755 (2)	0.0189 (3)
H1	0.7051	0.0559	0.2909	0.023*
H6A	1.1093	0.5659	0.1327	0.028*
H5A	0.9661	0.2895	0.4265	0.028*
H5B	0.7950	0.2793	0.4868	0.028*
H6B	1.0903	0.5704	0.3176	0.028*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Sr1	0.01678 (8)	0.01181 (8)	0.01109 (8)	−0.00511 (5)	−0.00483 (5)	0.00017 (5)
O1	0.0290 (6)	0.0226 (6)	0.0127 (5)	−0.0120 (5)	−0.0006 (5)	−0.0034 (4)
O3	0.0265 (6)	0.0195 (6)	0.0153 (5)	−0.0035 (5)	−0.0100 (5)	0.0024 (4)
O2	0.0417 (8)	0.0208 (6)	0.0155 (6)	−0.0187 (5)	−0.0006 (5)	−0.0019 (5)
O4	0.0266 (6)	0.0181 (6)	0.0196 (6)	−0.0042 (5)	−0.0097 (5)	−0.0059 (5)
O5	0.0194 (6)	0.0245 (6)	0.0418 (8)	−0.0066 (5)	−0.0101 (6)	0.0104 (6)
O6	0.0222 (6)	0.0457 (8)	0.0198 (6)	−0.0126 (6)	−0.0007 (5)	0.0007 (6)
N1	0.0317 (8)	0.0162 (6)	0.0153 (6)	−0.0076 (5)	−0.0104 (6)	0.0037 (5)
C7	0.0158 (7)	0.0163 (7)	0.0110 (7)	−0.0058 (5)	−0.0040 (5)	−0.0006 (5)
C4	0.0177 (7)	0.0141 (7)	0.0120 (7)	−0.0048 (5)	−0.0052 (5)	−0.0009 (5)
C3	0.0202 (7)	0.0128 (7)	0.0131 (7)	−0.0065 (5)	−0.0056 (5)	0.0013 (5)
C2	0.0195 (7)	0.0147 (7)	0.0117 (7)	−0.0077 (5)	−0.0044 (5)	−0.0016 (5)
C5	0.0268 (8)	0.0127 (7)	0.0161 (7)	−0.0060 (6)	−0.0084 (6)	−0.0004 (6)
C6	0.0201 (7)	0.0156 (7)	0.0125 (7)	−0.0070 (6)	−0.0063 (6)	−0.0014 (6)
C1	0.0273 (8)	0.0196 (8)	0.0121 (7)	−0.0093 (6)	−0.0079 (6)	0.0016 (6)

Geometric parameters (Å, º)

Sr1—O6	2.5287 (13)	O5—H5A	0.8542
Sr1—O3^{i}	2.5377 (12)	O5—H5B	0.8477
Sr1—O1^{ii}	2.5715 (12)	O6—H6A	0.8468
Sr1—O4^{iii}	2.5881 (12)	O6—H6B	0.8563
Sr1—O5	2.6050 (13)	N1—C1	1.339 (2)
Sr1—O2	2.6423 (12)	N1—C5	1.345 (2)
Sr1—O1	2.7232 (12)	N1—Sr1^{vi}	2.8065 (14)
Sr1—N1^{iv}	2.8064 (14)	C7—C4	1.508 (2)
Sr1—C6	3.0051 (15)	C4—C5	1.390 (2)
Sr1—H6B	2.9429	C4—C3	1.391 (2)
O1—C6	1.2520 (19)	C3—C2	1.387 (2)
O1—Sr1^{ii}	2.5715 (12)	C3—H3	0.9300
O3—C7	1.2590 (19)	C2—C1	1.388 (2)
O3—Sr1^{i}	2.5377 (12)	C2—C6	1.502 (2)
O2—C6	1.258 (2)	C5—H5	0.9300
O4—C7	1.2554 (19)	C1—H1	0.9300
O4—Sr1^{v}	2.5881 (12)		
O6—Sr1—O3^{i}	146.80 (4)	O1^{ii}—Sr1—H6B	119.9
O6—Sr1—O1^{ii}	133.77 (4)	O4^{iii}—Sr1—H6B	65.0
O3^{i}—Sr1—O1^{ii}	75.71 (4)	O5—Sr1—H6B	63.0
O6—Sr1—O4^{iii}	77.95 (4)	O2—Sr1—H6B	98.6
O3^{i}—Sr1—O4^{iii}	95.67 (4)	O1—Sr1—H6B	121.6
O1^{ii}—Sr1—O4^{iii}	81.28 (4)	N1^{iv}—Sr1—H6B	85.2
O6—Sr1—O5	69.37 (4)	C6—Sr1—H6B	107.7
O3^{i}—Sr1—O5	143.83 (4)	Sr1^{ii}—Sr1—H6B	128.1
O1^{ii}—Sr1—O5	70.70 (4)	C6—O1—Sr1^{ii}	156.53 (11)

O4^{iii}—Sr1—O5	92.21 (4)	C6—O1—Sr1	90.40 (9)
O6—Sr1—O2	84.52 (4)	Sr1^{ii}—O1—Sr1	112.09 (4)
O3^{i}—Sr1—O2	95.32 (4)	C7—O3—Sr1^{i}	140.72 (10)
O1^{ii}—Sr1—O2	116.27 (4)	C6—O2—Sr1	94.03 (9)
O4^{iii}—Sr1—O2	161.21 (4)	C7—O4—Sr1^{v}	137.91 (10)
O5—Sr1—O2	87.86 (4)	Sr1—O5—H5A	127.0
O6—Sr1—O1	115.98 (4)	Sr1—O5—H5B	115.1
O3^{i}—Sr1—O1	86.90 (4)	H5A—O5—H5B	100.7
O1^{ii}—Sr1—O1	67.91 (4)	Sr1—O6—H6A	131.4
O4^{iii}—Sr1—O1	147.46 (4)	Sr1—O6—H6B	110.7
O5—Sr1—O1	68.48 (4)	H6A—O6—H6B	117.3
O2—Sr1—O1	48.48 (4)	C1—N1—C5	117.16 (14)
O6—Sr1—N1^{iv}	74.58 (5)	C1—N1—Sr1^{vi}	109.60 (10)
O3^{i}—Sr1—N1^{iv}	73.32 (4)	C5—N1—Sr1^{vi}	130.26 (11)
O1^{ii}—Sr1—N1^{iv}	147.89 (4)	O4—C7—O3	124.65 (14)
O4^{iii}—Sr1—N1^{iv}	93.47 (4)	O4—C7—C4	118.35 (14)
O5—Sr1—N1^{iv}	141.38 (4)	O3—C7—C4	117.00 (13)
O2—Sr1—N1^{iv}	75.20 (4)	C5—C4—C3	118.26 (14)
O1—Sr1—N1^{iv}	118.12 (4)	C5—C4—C7	121.27 (14)
O6—Sr1—C6	97.37 (4)	C3—C4—C7	120.27 (13)
O3^{i}—Sr1—C6	95.75 (4)	C2—C3—C4	119.12 (14)
O1^{ii}—Sr1—C6	92.37 (4)	C2—C3—H3	120.4
O4^{iii}—Sr1—C6	165.14 (4)	C4—C3—H3	120.4
O5—Sr1—C6	72.98 (4)	C3—C2—C1	118.28 (14)
O2—Sr1—C6	24.68 (4)	C3—C2—C6	122.02 (14)
O1—Sr1—C6	24.62 (4)	C1—C2—C6	119.69 (13)
N1^{iv}—Sr1—C6	98.90 (4)	N1—C5—C4	123.41 (14)
O6—Sr1—Sr1^{ii}	132.57 (3)	N1—C5—H5	118.3
O3^{i}—Sr1—Sr1^{ii}	79.75 (3)	C4—C5—H5	118.3
O1^{ii}—Sr1—Sr1^{ii}	35.06 (3)	O1—C6—O2	122.85 (14)
O4^{iii}—Sr1—Sr1^{ii}	115.72 (3)	O1—C6—C2	118.69 (14)
O5—Sr1—Sr1^{ii}	65.11 (3)	O2—C6—C2	118.45 (14)
O2—Sr1—Sr1^{ii}	81.27 (3)	O1—C6—Sr1	64.98 (8)
O1—Sr1—Sr1^{ii}	32.85 (2)	O2—C6—Sr1	61.29 (8)
N1^{iv}—Sr1—Sr1^{ii}	142.01 (3)	C2—C6—Sr1	159.61 (11)
C6—Sr1—Sr1^{ii}	57.36 (3)	N1—C1—C2	123.75 (14)
O6—Sr1—H6B	15.8	N1—C1—H1	118.1
O3^{i}—Sr1—H6B	150.4	C2—C1—H1	118.1

O6—Sr1—O1—C6	−43.90 (11)	Sr1—O1—C6—O2	−21.19 (17)
O3^{i}—Sr1—O1—C6	111.03 (10)	Sr1^{ii}—O1—C6—C2	−6.4 (4)
O1^{ii}—Sr1—O1—C6	−173.14 (12)	Sr1—O1—C6—C2	157.48 (13)
O4^{iii}—Sr1—O1—C6	−153.28 (9)	Sr1^{ii}—O1—C6—Sr1	−163.9 (3)
O5—Sr1—O1—C6	−96.05 (10)	Sr1—O2—C6—O1	21.92 (17)
O2—Sr1—O1—C6	11.13 (9)	Sr1—O2—C6—C2	−156.75 (12)
N1^{iv}—Sr1—O1—C6	41.81 (11)	C3—C2—C6—O1	139.97 (16)
Sr1^{ii}—Sr1—O1—C6	−173.14 (12)	C1—C2—C6—O1	−39.0 (2)
O6—Sr1—O1—Sr1^{ii}	129.25 (5)	C3—C2—C6—O2	−41.3 (2)

O3^{i}—Sr1—O1—Sr1^{ii}	−75.83 (5)	C1—C2—C6—O2	139.73 (16)
O1^{ii}—Sr1—O1—Sr1^{ii}	0.0	C3—C2—C6—Sr1	−125.0 (3)
O4^{iii}—Sr1—O1—Sr1^{ii}	19.86 (10)	C1—C2—C6—Sr1	56.1 (4)
O5—Sr1—O1—Sr1^{ii}	77.09 (5)	O6—Sr1—C6—O1	141.06 (10)
O2—Sr1—O1—Sr1^{ii}	−175.73 (8)	O3^{i}—Sr1—C6—O1	−69.51 (10)
N1^{iv}—Sr1—O1—Sr1^{ii}	−145.05 (5)	O1^{ii}—Sr1—C6—O1	6.36 (11)
C6—Sr1—O1—Sr1^{ii}	173.14 (12)	O4^{iii}—Sr1—C6—O1	70.52 (19)
O6—Sr1—O2—C6	121.16 (11)	O5—Sr1—C6—O1	75.34 (10)
O3^{i}—Sr1—O2—C6	−92.19 (10)	O2—Sr1—C6—O1	−159.75 (16)
O1^{ii}—Sr1—O2—C6	−15.52 (11)	N1^{iv}—Sr1—C6—O1	−143.48 (9)
O4^{iii}—Sr1—O2—C6	142.24 (12)	Sr1^{ii}—Sr1—C6—O1	4.41 (8)
O5—Sr1—O2—C6	51.69 (10)	O6—Sr1—C6—O2	−59.19 (11)
O1—Sr1—O2—C6	−11.10 (9)	O3^{i}—Sr1—C6—O2	90.24 (10)
N1^{iv}—Sr1—O2—C6	−163.36 (11)	O1^{ii}—Sr1—C6—O2	166.11 (10)
Sr1^{ii}—Sr1—O2—C6	−13.45 (10)	O4^{iii}—Sr1—C6—O2	−129.73 (16)
Sr1^{v}—O4—C7—O3	−33.4 (3)	O5—Sr1—C6—O2	−124.91 (11)
Sr1^{v}—O4—C7—C4	145.77 (12)	O1—Sr1—C6—O2	159.75 (16)
Sr1^{i}—O3—C7—O4	105.41 (18)	N1^{iv}—Sr1—C6—O2	16.27 (11)
Sr1^{i}—O3—C7—C4	−73.8 (2)	Sr1^{ii}—Sr1—C6—O2	164.16 (11)
O4—C7—C4—C5	−15.8 (2)	O6—Sr1—C6—C2	35.7 (3)
O3—C7—C4—C5	163.49 (15)	O3^{i}—Sr1—C6—C2	−174.9 (3)
O4—C7—C4—C3	169.42 (14)	O1^{ii}—Sr1—C6—C2	−99.0 (3)
O3—C7—C4—C3	−11.3 (2)	O4^{iii}—Sr1—C6—C2	−34.8 (4)
C5—C4—C3—C2	−0.7 (2)	O5—Sr1—C6—C2	−30.0 (3)
C7—C4—C3—C2	174.30 (14)	O2—Sr1—C6—C2	94.9 (3)
C4—C3—C2—C1	−0.2 (2)	O1—Sr1—C6—C2	−105.4 (3)
C4—C3—C2—C6	−179.19 (14)	N1^{iv}—Sr1—C6—C2	111.2 (3)
C1—N1—C5—C4	−1.0 (3)	Sr1^{ii}—Sr1—C6—C2	−100.9 (3)
Sr1^{vi}—N1—C5—C4	157.24 (12)	C5—N1—C1—C2	0.0 (3)
C3—C4—C5—N1	1.3 (2)	Sr1^{vi}—N1—C1—C2	−162.51 (13)
C7—C4—C5—N1	−173.61 (15)	C3—C2—C1—N1	0.6 (2)
Sr1^{ii}—O1—C6—O2	174.94 (19)	C6—C2—C1—N1	179.60 (15)

Symmetry codes: (i) −*x*+1, −*y*+1, −*z*; (ii) −*x*+1, −*y*+1, −*z*+1; (iii) *x*, *y*+1, *z*+1; (iv) *x*, *y*+1, *z*; (v) *x*, *y*−1, *z*−1; (vi) *x*, *y*−1, *z*.

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
O5—H5*A*···O4^{vii}	0.85	2.01	2.8046 (17)	155
O5—H5*B*···O3^{viii}	0.85	2.05	2.8718 (18)	163
O6—H6*A*···O2^{ix}	0.85	1.87	2.7135 (17)	172
O6—H6*B*···O5^{x}	0.86	2.03	2.848 (2)	160
C1—H1···O3^{viii}	0.93	2.37	3.286 (2)	169

Symmetry codes: (vii) −*x*+2, −*y*, −*z*; (viii) *x*, *y*, *z*+1; (ix) −*x*+2, −*y*+1, −*z*; (x) −*x*+2, −*y*+1, −*z*+1.