

Hexaaquabis[3,5-bis(hydroxyimino)-1-methyl-2,4,6-trioxocyclohexanido- $\kappa^2 N^3, O^4$]barium tetrahydrate

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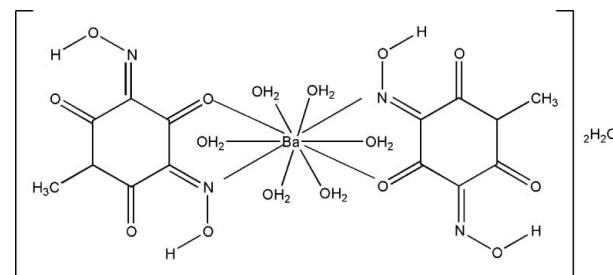
Received 3 August 2013; accepted 9 October 2013

Key indicators: single-crystal X-ray study; $T = 293\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.018; wR factor = 0.048; data-to-parameter ratio = 10.9.

In the title compound, $[\text{Ba}(\text{C}_7\text{H}_5\text{N}_2\text{O}_5)_2(\text{H}_2\text{O})_6]\cdot 4\text{H}_2\text{O}$, the Ba^{2+} cation lies on a twofold rotation axis and is ten-coordinated by two 3,5-bis(hydroxyimino)-1-methyl-2,4,6-trioxocyclohexanide oxo O atoms [$\text{Ba}-\text{O} = 2.8715(17)\text{ \AA}$], two hydroxyimino N atoms [$\text{Ba}-\text{N} = 3.036(2)\text{ \AA}$], and six water molecules [$\text{Ba}-\text{O} = 2.847(2)$, $2.848(2)$, and $2.880(2)\text{ \AA}$]. The 3,5-bis(hydroxyimino)-1-methyl-2,4,6-trioxocyclohexanide monoanions act in a bidentate chelating manner, coordinating through an N atom of the non-deprotonated hydroxyimino group and an O atom of the neighboring oxo group. Two lattice water molecules are located in the cavities of the framework and are involved in hydrogen bonding to O atoms of one of the coordinating water molecules and the O atom of a keto group of the ligand. As a result, a three-dimensional network is formed.

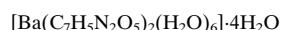
Related literature

For the synthesis and crystal structure of sodium 3,5-bis(hydroxyimino)-1-methyl-2,4,6-trioxocyclohexanide, see: Kovalchukova *et al.* (2012). For related structures of metal complexes with 1,2-benzo(naphtho)quinone-1-oximes, see: Chakravorty (1974); Charalambous *et al.* (1993, 1995, 1996); Adatia *et al.* (1996); Basu & Chakravorty (1992); McPartlin (1973); Djinovic *et al.* (1992); Liu *et al.* (2010). For applications of related complexes as catalysts, see: Gharah *et al.* (2009).



Experimental

Crystal data



$M_r = 711.76$

Monoclinic, $C2/c$

$a = 17.235(3)\text{ \AA}$

$b = 6.736(1)\text{ \AA}$

$c = 23.074(5)\text{ \AA}$

$\beta = 108.61(3)^\circ$

$V = 2538.7(9)\text{ \AA}^3$

$Z = 4$

Mo $K\alpha$ radiation

$\mu = 1.66\text{ mm}^{-1}$

$T = 293\text{ K}$

$0.33 \times 0.12 \times 0.07\text{ mm}$

Data collection

Enraf–Nonius CAD-4
diffractometer

Absorption correction: part of the

refinement model (ΔF)

(Walker & Stuart, 1983)

$T_{\min} = 0.370$, $T_{\max} = 0.780$

2429 measured reflections

2346 independent reflections

1755 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.019$

3 standard reflections every 60 min
intensity decay: none

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.018$

$wR(F^2) = 0.048$

$S = 0.98$

2346 reflections

216 parameters

17 restraints

H atoms treated by a mixture of
independent and constrained
refinement

$\Delta\rho_{\max} = 0.47\text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.90\text{ e \AA}^{-3}$

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O21—H21 \cdots O3	0.86 (3)	1.70 (3)	2.476 (3)	150 (4)
O61—H61 \cdots O5	0.86 (3)	1.64 (3)	2.469 (2)	160 (4)
O11—H111 \cdots O3 ⁱ	0.85 (1)	1.99 (1)	2.827 (2)	167 (4)
O11—H112 \cdots O12 ⁱⁱ	0.84 (1)	2.15 (3)	2.854 (3)	142 (4)
O12—H121 \cdots O14 ⁱⁱⁱ	0.84 (1)	1.90 (1)	2.739 (3)	172 (4)
O12—H122 \cdots O11 ^{iv}	0.84 (1)	2.17 (3)	2.839 (3)	137 (3)
O13—H131 \cdots O14	0.85 (1)	2.13 (1)	2.957 (3)	165 (4)
O13—H132 \cdots O15 ^v	0.85 (1)	1.93 (1)	2.766 (3)	169 (4)
O14—H141 \cdots O13 ^{vi}	0.85 (1)	1.99 (1)	2.835 (3)	173 (4)
O15—H151 \cdots O5 ^{vii}	0.85 (1)	1.94 (1)	2.789 (3)	177 (4)

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

Data collection: *CAD-4-PC* (Enraf–Nonius, 1993); cell refinement: *CAD-4-PC*; data reduction: *CAD-4-PC*; 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: *CIFTAB97* (Sheldrick, 2008) and *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BV2225).

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supporting information

Acta Cryst. (2013). E69, m602–m603 [doi:10.1107/S1600536813027761]

Hexaaqua bis[3,5-bis(hydroxyimino)-1-methyl-2,4,6-trioxocyclohexanido- κ^2N^3,O^4]barium tetrahydrate

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S1. Comment

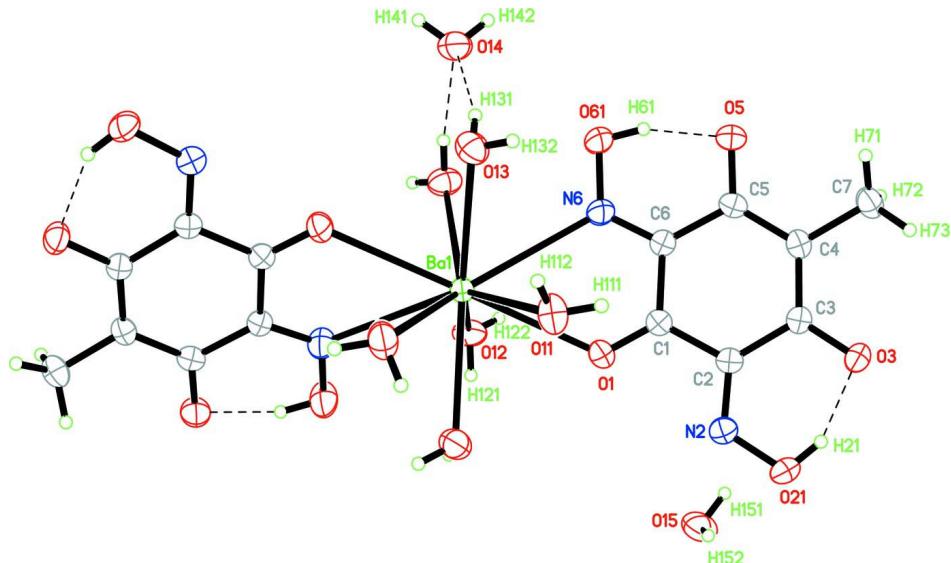
1,2-Quinone monooximes (2-nitrosophenols) are good chelating agents which form stable colored complexes with a wide range of metal ions (Chakravorty, 1974; Charalambous *et al.*, 1996; Djinovic *et al.*, 1992; Liu *et al.*, 2010). Increasing attention is also being devoted to these complexes in recent years because of their applications in organic synthesis (Gharah *et al.*, 2009). Sodium 3,5-bis(hydroxyimino)-1-methyl-2,4,6-trioxocyclohexanide has been recently isolated as the only product of the reaction of nitrosation of methylphloroglucinol, and its crystal and molecular structure was reported (Kovalchukova *et al.*, 2012). In the present paper we report the crystal and molecular structure of barium hexa-aqua bis (3,5-bis(hydroxyimino)-1-methyl-2,4,6- trioxocyclohexanide) dihydrate, $C_{14}H_{30}BaN_4O_{20}$. The Ba cation of the reported structure lies on a center of inversion, and is ten-coordinated. It is surrounded by two oxo O atoms of 3,5-bis(hydroxyimino)-1-methyl-2,4,6- trioxocyclohexanide species [$Ba—O = 2.8715$ (17) Å], two hydroxyimino N atoms [$Ba—N = 3.036$ (2) Å], and six water molecules [$Ba—O = 2.847$ (2), 2.848 (2), and 2.880 (2) Å]. The 3,5-bis(hydroxyimino)-1-methyl-2,4,6- trioxocyclohexanide mono anions act as bidentate chelating coordinated through an N-atom of the non-deprotonated hydroxyimino group and an O-atom of the neighboring oxo-group. The conjugation in the chelate ring is observed in the equalizing of the chemical bonds [$C1—O1 = 1.219$ (3), $C6—N6 = 1.299$ (3), $N6—O61 = 1.352$, $C1—C6 = 1.482$ Å]. There are two intramolecular hydrogen bonds in the 3,5-bis(hydroxyimino)-1-methyl-2,4,6- trioxocyclohexanide anion adjusting the hydroxyimino H atoms with the neighboring O oxo atoms. The coordinated water molecules are involved in O—H···O hydrogen bonds with the O atoms of the organic anion. Two lattice water molecules are located in the cavities of the framework and involved in hydrogen bonding to O atoms of one of the coordinated H_2O and the O atom of an oxo group of the ligand. As a result, a three-dimensional network is formed. The described structure is in accordance with the other known structures of complexes of 1,2-benzo(naphto) quinone -1-oximes with the alkaline (Charalambous *et al.*, 1993, 1995; Adatia *et al.*, 1996) and transition metals (Basu and Chakravorty, 1992; Charalambous *et al.*, 1996; McPartlin, 1973) where the formation of monomeric complexes with the bidentate chelating coordination mode of the ligands was observed. The only one difference is in the nature of the hydroxyimino group which is not ionized in the structure reported here. From the other side, the 3,5-bis(hydroxyimino)-1-methyl-2,4,6-trioxocyclohexanide anion in the sodium salt (Kovalchukova *et al.*, 2012) acts as m-1,2,2,3 polydentate bridging ligands were both N and O hydroxyimino atoms take part in coordination, and one of the O oxo coordinated atoms forms a bifurcated bond.

S2. Experimental

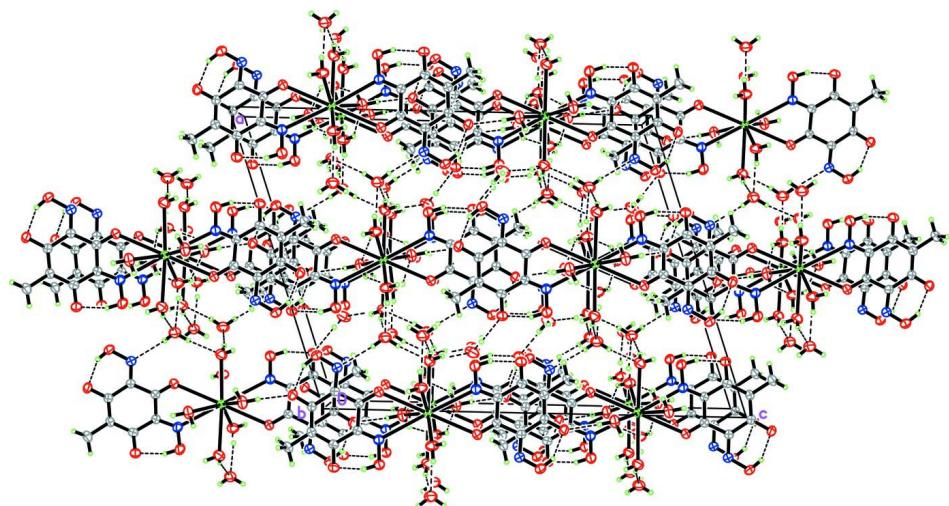
Single crystals of $C_{14}H_{30}BaN_4O_{20}$ were grown by the slow evaporation of the ethanol solution of the 1-to-1 molar mixture of barium nitrate and sodium 3,5-bis(hydroxyimino)-1-methyl-2,4,6-trioxocyclohexanide.

S3. Refinement

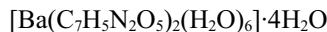
The structure of $C_{14}H_{30}BaN_4O_{20}$ was solved by direct methods and all non-hydrogen atoms were located and refined in anisotropically. All the hydrogen atoms were located in difference electron density syntheses and included in refinement with fixed parameters. The O-H distances in the hydroxy groups were constrained at 0.84 Å. The O-H distances and H-O-H angles in the water molecules were constrained. A riding model was used in the refinement of H atoms of the methyl group.

**Figure 1**

ORTEP view of $C_{14}H_{30}BaN_4O_{20}$ with atom labeling scheme (displacement ellipsoids are drawn at the 50% probability level for non-hydrogen atoms). Hydrogen bonds shown as dashed lines.

**Figure 2**

Molecular packing in the crystal of the complex along the crystallographic axis b . Hydrogen bonds shown as dashed lines.

Hexaaquabis[3,5-bis(hydroxyimino)-1-methyl-2,4,6-trioxocyclohexanido- κ^2N^3,O^4]barium tetrahydrate*Crystal data*

$M_r = 711.76$

Monoclinic, $C2/c$

Hall symbol: -C 2yc

$a = 17.235 (3) \text{ \AA}$

$b = 6.736 (1) \text{ \AA}$

$c = 23.074 (5) \text{ \AA}$

$\beta = 108.61 (3)^\circ$

$V = 2538.7 (9) \text{ \AA}^3$

$Z = 4$

$F(000) = 1432$

$D_x = 1.862 \text{ Mg m}^{-3}$

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

Cell parameters from 25 reflections

$\theta = 9.2\text{--}11.7^\circ$

$\mu = 1.66 \text{ mm}^{-1}$

$T = 293 \text{ K}$

Plate, red

$0.33 \times 0.12 \times 0.07 \text{ mm}$

Data collection

Enraf–Nonius CAD-4

diffractometer

Radiation source: fine-focus sealed tube

β -filter monochromator

$\omega/2\theta$ scans

Absorption correction: part of the refinement
model (ΔF)

(Walker & Stuart, 1983)

$T_{\min} = 0.370$, $T_{\max} = 0.780$

2429 measured reflections

2346 independent reflections

1755 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.019$

$\theta_{\max} = 25.5^\circ$, $\theta_{\min} = 1.9^\circ$

$h = 0 \rightarrow 20$

$k = -8 \rightarrow 0$

$l = -27 \rightarrow 26$

3 standard reflections every 60 min

intensity decay: none

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.018$

$wR(F^2) = 0.048$

$S = 0.98$

2346 reflections

216 parameters

17 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: difference Fourier map

H atoms treated by a mixture of independent
and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0317P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$

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

$\Delta\rho_{\max} = 0.47 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.90 \text{ e \AA}^{-3}$

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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Ba1	0.5000	0.24104 (3)	0.7500	0.02124 (7)
O1	0.56648 (10)	0.2550 (3)	0.65071 (7)	0.0332 (4)
O3	0.54847 (10)	0.3322 (3)	0.44397 (7)	0.0317 (4)

O5	0.32241 (10)	0.0962 (3)	0.49495 (7)	0.0349 (4)
O11	0.48141 (12)	0.6073 (3)	0.68255 (8)	0.0377 (4)
O12	0.59225 (10)	-0.1143 (3)	0.75770 (9)	0.0368 (4)
O13	0.32986 (11)	0.3326 (3)	0.70959 (8)	0.0371 (4)
O14	0.24224 (11)	-0.0492 (3)	0.70127 (9)	0.0447 (5)
O15	0.80288 (12)	0.0550 (3)	0.60419 (10)	0.0486 (5)
O21	0.67935 (10)	0.3795 (3)	0.52748 (8)	0.0335 (4)
O61	0.33722 (12)	0.0432 (3)	0.60383 (8)	0.0443 (5)
N2	0.64131 (11)	0.3306 (3)	0.56837 (9)	0.0259 (4)
N6	0.41475 (12)	0.1133 (3)	0.61886 (9)	0.0317 (4)
C1	0.52814 (12)	0.2383 (3)	0.59644 (9)	0.0226 (4)
C2	0.56427 (13)	0.2839 (3)	0.54772 (10)	0.0212 (4)
C3	0.51345 (14)	0.2780 (3)	0.48228 (10)	0.0214 (4)
C4	0.43167 (14)	0.2147 (3)	0.46508 (9)	0.0230 (4)
C5	0.39533 (13)	0.1576 (3)	0.50912 (10)	0.0235 (4)
C6	0.44222 (13)	0.1663 (3)	0.57513 (10)	0.0222 (4)
C7	0.38032 (15)	0.2124 (4)	0.39871 (10)	0.0314 (5)
H71	0.3245	0.2422	0.3951	0.038*
H72	0.3831	0.0834	0.3818	0.038*
H73	0.4005	0.3101	0.3769	0.038*
H21	0.646 (2)	0.360 (7)	0.4916 (14)	0.077 (4)*
H61	0.321 (2)	0.050 (6)	0.5645 (14)	0.077 (4)*
H111	0.470 (2)	0.606 (6)	0.6438 (5)	0.077 (4)*
H112	0.4404 (16)	0.664 (6)	0.6874 (16)	0.077 (4)*
H121	0.6431 (8)	-0.096 (6)	0.7668 (16)	0.077 (4)*
H122	0.582 (2)	-0.198 (5)	0.7293 (14)	0.077 (4)*
H131	0.302 (2)	0.228 (4)	0.7002 (17)	0.077 (4)*
H132	0.315 (2)	0.399 (5)	0.6766 (10)	0.077 (4)*
H141	0.217 (2)	-0.083 (6)	0.7256 (12)	0.077 (4)*
H142	0.2137 (19)	-0.102 (6)	0.6680 (9)	0.077 (4)*
H151	0.7660 (19)	0.008 (5)	0.5737 (12)	0.077 (4)*
H152	0.794 (3)	0.1784 (18)	0.6008 (18)	0.077 (4)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Ba1	0.02168 (10)	0.02131 (10)	0.02071 (9)	0.000	0.00676 (6)	0.000
O1	0.0286 (8)	0.0467 (10)	0.0227 (7)	-0.0042 (8)	0.0059 (6)	-0.0036 (8)
O3	0.0325 (9)	0.0387 (9)	0.0265 (8)	-0.0059 (8)	0.0131 (7)	0.0014 (7)
O5	0.0256 (8)	0.0448 (10)	0.0327 (9)	-0.0101 (8)	0.0070 (7)	-0.0014 (8)
O11	0.0491 (11)	0.0352 (10)	0.0281 (8)	0.0063 (8)	0.0116 (8)	0.0013 (7)
O12	0.0292 (8)	0.0334 (9)	0.0473 (10)	-0.0028 (8)	0.0114 (8)	-0.0080 (8)
O13	0.0344 (9)	0.0388 (10)	0.0347 (9)	0.0039 (8)	0.0062 (8)	0.0012 (8)
O14	0.0308 (9)	0.0582 (13)	0.0432 (10)	-0.0059 (9)	0.0092 (8)	0.0087 (10)
O15	0.0353 (10)	0.0522 (12)	0.0471 (11)	-0.0040 (10)	-0.0025 (8)	0.0100 (10)
O21	0.0273 (8)	0.0412 (10)	0.0351 (9)	-0.0068 (8)	0.0145 (7)	0.0002 (8)
O61	0.0343 (9)	0.0687 (14)	0.0310 (9)	-0.0242 (10)	0.0120 (8)	-0.0031 (9)
N2	0.0267 (10)	0.0223 (9)	0.0292 (10)	-0.0008 (8)	0.0097 (8)	-0.0012 (8)

N6	0.0294 (10)	0.0372 (12)	0.0293 (10)	-0.0115 (9)	0.0105 (8)	-0.0031 (9)
C1	0.0246 (10)	0.0187 (9)	0.0241 (9)	0.0008 (9)	0.0073 (8)	-0.0011 (9)
C2	0.0235 (10)	0.0137 (11)	0.0263 (10)	0.0013 (8)	0.0078 (8)	-0.0025 (8)
C3	0.0269 (10)	0.0138 (11)	0.0242 (9)	0.0021 (8)	0.0092 (8)	0.0000 (8)
C4	0.0293 (11)	0.0156 (11)	0.0226 (10)	0.0022 (8)	0.0064 (8)	0.0005 (8)
C5	0.0229 (11)	0.0180 (10)	0.0277 (11)	-0.0004 (9)	0.0054 (8)	-0.0018 (9)
C6	0.0250 (11)	0.0174 (9)	0.0244 (10)	0.0001 (9)	0.0082 (9)	-0.0011 (8)
C7	0.0352 (12)	0.0301 (13)	0.0245 (10)	-0.0031 (9)	0.0033 (9)	0.0003 (9)

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

Ba1—O13	2.8467 (19)	O15—H151	0.846 (10)
Ba1—O13 ⁱ	2.8467 (19)	O15—H152	0.845 (10)
Ba1—O12	2.8475 (19)	O21—N2	1.350 (3)
Ba1—O12 ⁱ	2.8475 (19)	O21—H21	0.86 (3)
Ba1—O1 ⁱ	2.8715 (17)	O61—N6	1.354 (3)
Ba1—O1	2.8715 (17)	O61—H61	0.86 (3)
Ba1—O11 ⁱ	2.8799 (19)	N2—C2	1.298 (3)
Ba1—O11	2.8799 (19)	N6—C6	1.294 (3)
Ba1—N6	3.036 (2)	C1—C2	1.481 (3)
Ba1—N6 ⁱ	3.036 (2)	C1—C6	1.485 (3)
O1—C1	1.220 (3)	C2—C3	1.485 (3)
O3—C3	1.273 (3)	C3—O3	1.273 (3)
O5—C5	1.263 (3)	C3—C4	1.403 (3)
O11—H111	0.851 (10)	C4—C5	1.407 (3)
O11—H112	0.841 (10)	C4—C7	1.504 (3)
O12—H121	0.842 (10)	C5—O5	1.263 (3)
O12—H122	0.841 (10)	C5—C6	1.480 (3)
O13—H131	0.847 (10)	C7—H71	0.9600
O13—H132	0.847 (10)	C7—H72	0.9600
O14—H141	0.851 (10)	C7—H73	0.9600
O14—H142	0.849 (10)		
O13—Ba1—O13 ⁱ	154.98 (9)	Ba1—O11—H111	121 (3)
O13—Ba1—O12	134.33 (6)	Ba1—O11—H112	106 (3)
O13 ⁱ —Ba1—O12	70.45 (6)	H111—O11—H112	103 (2)
O13—Ba1—O12 ⁱ	70.45 (6)	Ba1—O12—H121	114 (3)
O13 ⁱ —Ba1—O12 ⁱ	134.33 (5)	Ba1—O12—H122	123 (3)
O12—Ba1—O12 ⁱ	65.59 (7)	H121—O12—H122	104 (2)
O13—Ba1—O1 ⁱ	67.83 (5)	Ba1—O13—H131	111 (3)
O13 ⁱ —Ba1—O1 ⁱ	111.29 (5)	Ba1—O13—H132	113 (3)
O12—Ba1—O1 ⁱ	109.55 (5)	H131—O13—H132	103 (2)
O12 ⁱ —Ba1—O1 ⁱ	73.76 (5)	H141—O14—H142	102 (2)
O13—Ba1—O1	111.30 (5)	H151—O15—H152	103 (2)
O13 ⁱ —Ba1—O1	67.83 (5)	N2—O21—H21	108 (3)
O12—Ba1—O1	73.76 (5)	N6—O61—H61	102 (3)
O12 ⁱ —Ba1—O1	109.55 (5)	C2—N2—O21	118.06 (19)
O1 ⁱ —Ba1—O1	176.25 (8)	C6—N6—O61	118.25 (19)

O13—Ba1—O11 ⁱ	85.23 (6)	C6—N6—Ba1	121.19 (14)
O13 ⁱ —Ba1—O11 ⁱ	73.27 (6)	O61—N6—Ba1	118.58 (13)
O12—Ba1—O11 ⁱ	136.29 (6)	O1—C1—C2	122.60 (19)
O12 ⁱ —Ba1—O11 ⁱ	135.85 (5)	O1—C1—C6	121.73 (19)
O1 ⁱ —Ba1—O11 ⁱ	62.92 (5)	C2—C1—C6	115.65 (18)
O1—Ba1—O11 ⁱ	113.53 (5)	N2—C2—C1	113.59 (19)
O13—Ba1—O11	73.27 (6)	N2—C2—C3	125.6 (2)
O13 ⁱ —Ba1—O11	85.23 (6)	C1—C2—C3	120.79 (19)
O12—Ba1—O11	135.85 (5)	O3—C3—C4	123.2 (2)
O12 ⁱ —Ba1—O11	136.29 (5)	O3—C3—C4	123.2 (2)
O1 ⁱ —Ba1—O11	113.53 (5)	O3—C3—C2	116.23 (19)
O1—Ba1—O11	62.92 (5)	O3—C3—C2	116.23 (19)
O11 ⁱ —Ba1—O11	62.11 (7)	C4—C3—C2	120.62 (19)
O13—Ba1—N6	67.33 (6)	C3—C4—C5	121.15 (19)
O13 ⁱ —Ba1—N6	120.54 (6)	C3—C4—C7	120.2 (2)
O12—Ba1—N6	84.66 (6)	C5—C4—C7	118.7 (2)
O12 ⁱ —Ba1—N6	67.46 (6)	O5—C5—C4	122.6 (2)
O1 ⁱ —Ba1—N6	127.95 (5)	O5—C5—C4	122.6 (2)
O1—Ba1—N6	53.38 (5)	O5—C5—C6	116.8 (2)
O11 ⁱ —Ba1—N6	135.75 (6)	O5—C5—C6	116.8 (2)
O11—Ba1—N6	76.65 (6)	C4—C5—C6	120.64 (19)
O13—Ba1—N6 ⁱ	120.54 (6)	N6—C6—C5	125.2 (2)
O13 ⁱ —Ba1—N6 ⁱ	67.33 (6)	N6—C6—C1	113.97 (19)
O12—Ba1—N6 ⁱ	67.46 (6)	C5—C6—C1	120.87 (19)
O12 ⁱ —Ba1—N6 ⁱ	84.66 (6)	C4—C7—H71	109.5
O1 ⁱ —Ba1—N6 ⁱ	53.38 (5)	C4—C7—H72	109.5
O1—Ba1—N6 ⁱ	127.96 (5)	H71—C7—H72	109.5
O11 ⁱ —Ba1—N6 ⁱ	76.65 (6)	C4—C7—H73	109.5
O11—Ba1—N6 ⁱ	135.76 (6)	H71—C7—H73	109.5
N6—Ba1—N6 ⁱ	147.08 (8)	H72—C7—H73	109.5
C1—O1—Ba1	126.33 (14)		

Symmetry code: (i) $-x+1, y, -z+3/2$.

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

$D\text{—H}\cdots A$	$D\text{—H}$	$H\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
O21—H21 \cdots O3	0.86 (3)	1.70 (3)	2.476 (3)	150 (4)
O61—H61 \cdots O5	0.86 (3)	1.64 (3)	2.469 (2)	160 (4)
O11—H111 \cdots O3 ⁱⁱ	0.85 (1)	1.99 (1)	2.827 (2)	167 (4)
O11—H112 \cdots O12 ⁱⁱⁱ	0.84 (1)	2.15 (3)	2.854 (3)	142 (4)
O12—H121 \cdots O14 ⁱ	0.84 (1)	1.90 (1)	2.739 (3)	172 (4)
O12—H122 \cdots O11 ^{iv}	0.84 (1)	2.17 (3)	2.839 (3)	137 (3)
O13—H131 \cdots O14	0.85 (1)	2.13 (1)	2.957 (3)	165 (4)
O13—H132 \cdots O15 ^v	0.85 (1)	1.93 (1)	2.766 (3)	169 (4)

O14—H141···O13 ^{vi}	0.85 (1)	1.99 (1)	2.835 (3)	173 (4)
O15—H151···O5 ^{vii}	0.85 (1)	1.94 (1)	2.789 (3)	177 (4)

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