

Poly[μ_6 -adipato-diaquadi- μ_2 -oxalato-digadolium(III)]

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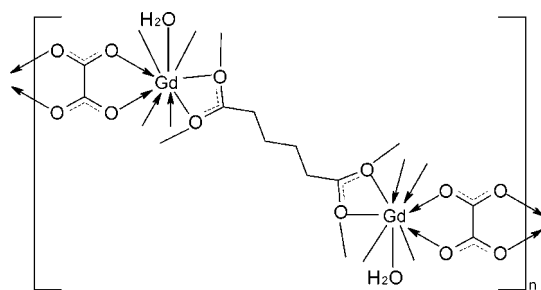
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 Key indicators: single-crystal X-ray study; $T = 295$ K; mean $\sigma(\text{C}-\text{C}) = 0.008$ Å; disorder in main residue; R factor = 0.023; wR factor = 0.059; data-to-parameter ratio = 9.7.

In the centrosymmetric title compound, $[\text{Gd}_2(\text{C}_6\text{H}_8\text{O}_4)(\text{C}_2\text{O}_4)_2(\text{H}_2\text{O})_2]_n$, the Gd^{3+} cations are each coordinated by nine O atoms, three from adipate anions, two from oxalate anions and one from an aqua ligand, completing a distorted tricapped trigonal-prismatic geometry. These tricapped trigonal prisms are bridged by the adipate ligands, generating layers lying parallel to (010). The coordination polymer layers are linked into a three-dimensional framework by the rigid oxalate ligands. The adipate and oxalate ions are all located on centers of inversion. A part of the adipate anion is disordered over two positions in a 0.75:0.25 ratio.

Related literature

For structures involving adipate ligands and lanthanide ions, see: Dimos *et al.* (2002). For structures involving oxalate ligands and lanthanide ions, see: Trombe & Mohanu (2004).



Experimental

Crystal data

$[\text{Gd}_2(\text{C}_6\text{H}_8\text{O}_4)(\text{C}_2\text{O}_4)_2(\text{H}_2\text{O})_2]$
 $M_r = 670.70$
 Triclinic, $P\bar{1}$
 $a = 6.815$ (4) Å
 $b = 6.982$ (4) Å
 $c = 8.997$ (7) Å
 $\alpha = 104.759$ (10)°
 $\beta = 108.11$ (1)°

$\gamma = 104.320$ (7)°
 $V = 367.8$ (4) Å³
 $Z = 1$
 Mo $K\alpha$ radiation
 $\mu = 9.02$ mm⁻¹
 $T = 295$ K
 $0.23 \times 0.11 \times 0.08$ mm

Data collection

Bruker SMART APEXII CCD
 area-detector diffractometer
 Absorption correction: multi-scan
 (*SADABS*; Sheldrick, 2003)
 $T_{\min} = 0.241$, $T_{\max} = 0.491$

1751 measured reflections
 1236 independent reflections
 1157 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.013$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.023$
 $wR(F^2) = 0.059$
 $S = 1.05$
 1236 reflections
 127 parameters
 2 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 1.56$ e Å⁻³
 $\Delta\rho_{\text{min}} = -1.41$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O7}-\text{H7A}\cdots\text{O3}^i$	0.84 (6)	2.00 (5)	2.799 (4)	160
$\text{O7}-\text{H7B}\cdots\text{O6}^{ii}$	0.83 (7)	2.07 (7)	2.888 (5)	168

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

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINTE* (Bruker, 2004); data reduction: *SAINTE*; 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*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SU2209).

References

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 Dimos, A., Tsaousis, D., Michaelides, A. & Skoulika, S. (2002). *Chem. Mater.* **14**, 2616–2622.
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supporting information

Acta Cryst. (2010). E66, m1263 [doi:10.1107/S1600536810036111]

Poly[μ_6 -adipato-diaquadi- μ_2 -oxalato-digadolinium(III)]**Zhi-Feng Li and Chun-Xiang Wang****S1. Comment**

As shown in Fig. 1 the asymmetric unit of the title compound consists of one Gd^{3+} cation, half of an adipate anion, two half oxalate anions and one aqua ligand. The Gd atom is coordinated by nine oxygen atoms, in which four oxygen atoms are from three adipate anions, four from two oxalate anions and one from a water molecule, to form a SmO_9 polyhedron with a distorted tricapped trigonal-prismatic geometry. The Gd—O(adipate) distances vary in the range of 2.430 (4)–2.575 (4) Å (average 2.512 (4) Å), which is nearly identical to the value of 2.425 (8)–2.611 (8) Å (average 2.471 (8) Å) observed in $\text{Gd}_2(\text{C}_6\text{H}_8\text{O}_4)_3(\text{H}_2\text{O})_4$ (Dimos *et al.*, 2002). The Gd—O(oxalate) distances are in the range of 2.385 (4)–2.540 (4) Å, which are usual for lanthanide oxalates complexes (Trombe & Mohanu, 2004). In the title complex, the adipate anions are located on inversion centers and atom C3 is positionally disordered (C3A and C3B; occupancies 0.75/0.25). Two carboxylate oxygen atoms chelate one Gd atom with each oxygen atom additionally bonded to another Gd atom. To the best of our knowledge, this η^2, μ_3 - η^2, μ_3 -chelating-bridging octadentate coordination mode of the adipate ligand has not been reported previously.

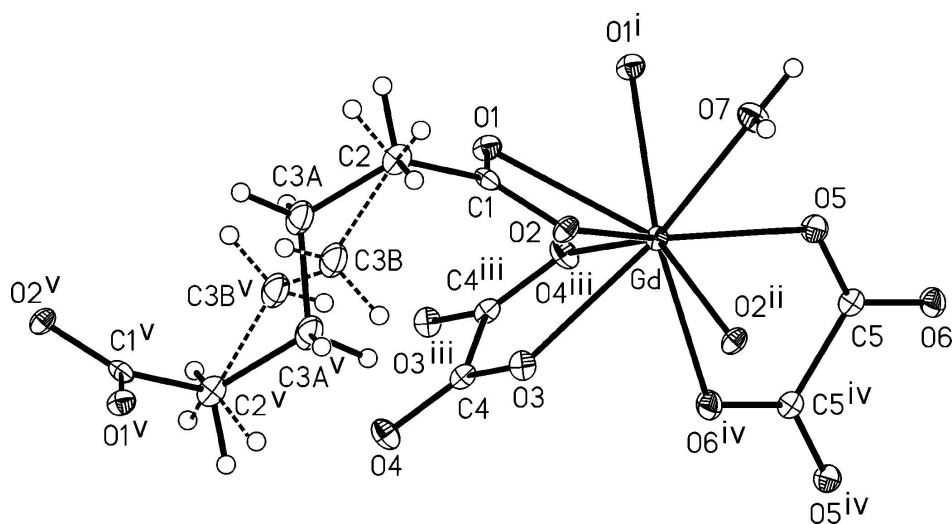
Through the terminal carboxylato bridging interactions, the GdO_9 polyhedra are edged-shared to generate metal-oxygen chains extending infinitely along [100], in which the adjacent Gd...Gd distances are 4.23 (3) Å and 4.25 (2) Å, respectively. Along [001] the chains are linked by the adipate anions into layers parallel to (010) (Fig. 2). Two symmetry independent oxalate ions are also located on centers of inversion and act as double bidentate (tetradentate) ligands in a linear chain, which connect Gd atoms to form zigzag chains along [001]. Through the oxalate and adipate ligand bridging interactions, the Gd atoms build up a three-dimensional open framework with the channels propagating in [001] (Fig. 3). The aqua ligand provides H-bond donors which participate in O—H...O hydrogen bonds with the oxalate atoms O3 and O6 (Table 1).

S2. Experimental

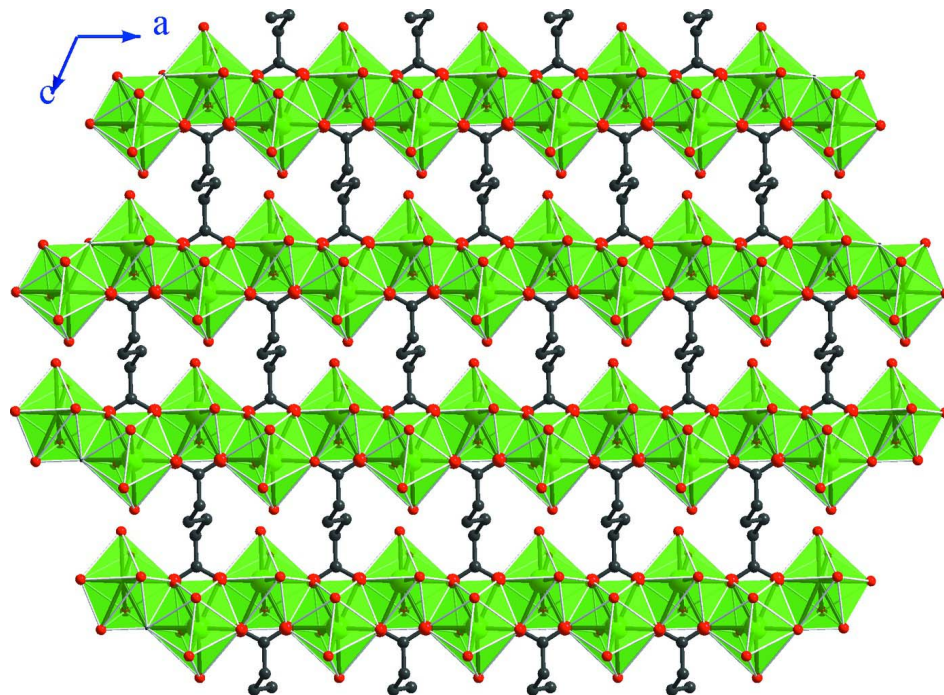
A mixture of $\text{GdCl}_3 \cdot 6\text{H}_2\text{O}$ (1.00 mmol, 0.36 g), oxalic acid (0.50 mmol, 0.05 g), adipic acid (0.50 mmol, 0.07 g), NaOH (2.00 mmol, 0.08 g) and H_2O (10.0 ml) was heated in a 23 ml stainless steel reactor with a Teflon liner at 443 K for 48 h. On cooling a small amount of colorless plate-like crystals were obtained. They were filtered off and washed with water and acetone.

S3. Refinement

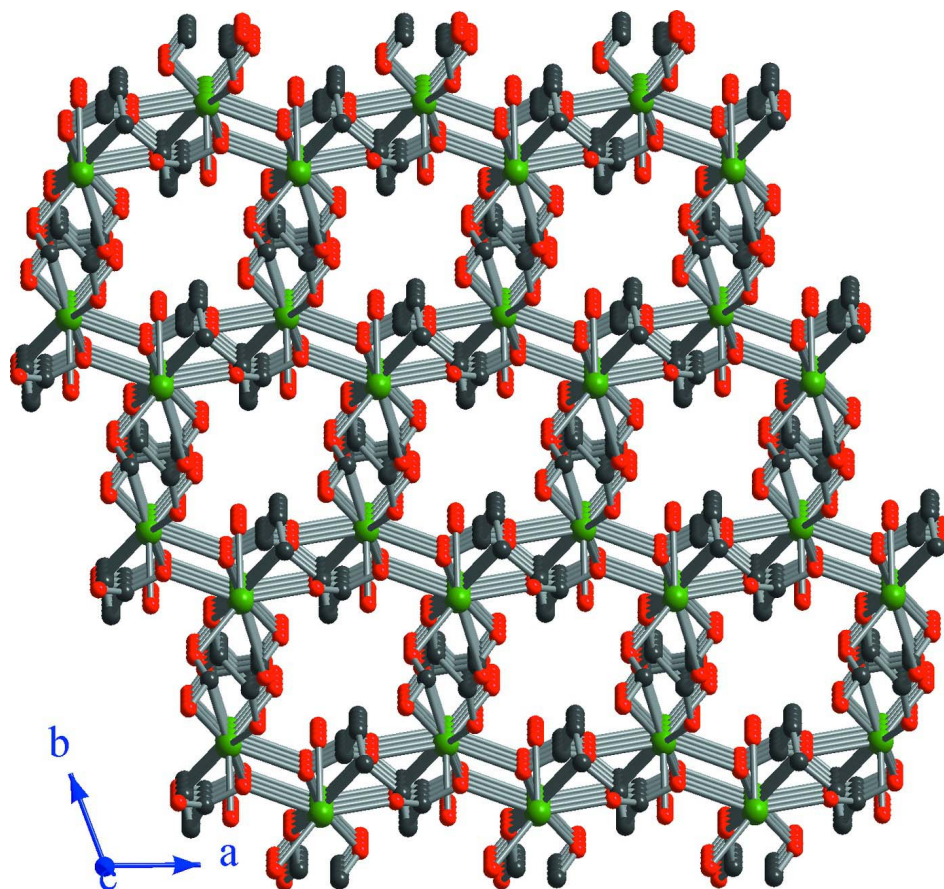
Atom C3 of the adipate anion is positionally disordered (C3A and C3B) and these atoms were refined with occupancies of 0.75/0.25. The water H-atoms were located in difference Fourier maps and were refined with distance restraints: O—H distance of 0.84 (2) Å and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$. The C-bound atoms were included in calculated positions and treated as riding atoms: C—H = 0.97 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. The highest density peak and deepest hole are located at 0.97 Å and 0.89 Å, respectively, from the Gd atom.

**Figure 1**

The fragment of the molecular structure of the title compound, with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii. Symmetry codes: (i) $-x, 1-y, 1-z$; (ii) $1-x, 1-y, 1-z$; (iii) $-x, -y, 1-z$; (iv) $-x, -y, -z$; (v) $1-x, 1-y, 2-z$.

**Figure 2**

A view along the *b*-axis of the two-dimensional layer structure formed by the connectivity between the Gd atoms and the adipate moieties present in the title compound [the C-bound H-atoms and atom C3B have been omitted for clarity].

**Figure 3**

A view along the *c*-axis of the three-dimensional framework of the title compound [the C-bound H-atoms and atom C3B have been omitted for clarity].

Poly[μ_6 -adipato-diaquadi- μ_2 -oxalato-digadolinium(III)]

Crystal data

[Gd₂(C₆H₈O₄)(C₂O₄)₂(H₂O)₂]

$M_r = 670.70$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 6.815$ (4) Å

$b = 6.982$ (4) Å

$c = 8.997$ (7) Å

$\alpha = 104.759$ (10)°

$\beta = 108.11$ (1)°

$\gamma = 104.320$ (7)°

$V = 367.8$ (4) Å³

$Z = 1$

$F(000) = 312$

$D_x = 3.028$ Mg m⁻³

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

Cell parameters from 238 reflections

$\theta = 2.6$ – 28.3 °

$\mu = 9.02$ mm⁻¹

$T = 295$ K

Plate, colorless

$0.23 \times 0.11 \times 0.08$ mm

Data collection

Bruker SMART APEXII CCD area-detector diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan (SADABS; Sheldrick, 2003)

$T_{\min} = 0.241$, $T_{\max} = 0.491$

1751 measured reflections

1236 independent reflections

1157 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.013$
 $\theta_{\text{max}} = 25.0^\circ$, $\theta_{\text{min}} = 2.6^\circ$

$h = -7 \rightarrow 8$
 $k = -7 \rightarrow 8$
 $l = -9 \rightarrow 10$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.023$
 $wR(F^2) = 0.059$
 $S = 1.05$
 1236 reflections
 127 parameters
 2 restraints
 Primary atom site location: structure-invariant
 direct methods

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
 $w = 1/[\sigma^2(F_o^2) + (0.0451P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 1.56 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\text{min}} = -1.41 \text{ e } \text{Å}^{-3}$

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 (Å^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
Gd	0.15528 (4)	0.34583 (3)	0.36187 (3)	0.01206 (12)	
O1	0.1523 (6)	0.5226 (6)	0.6494 (4)	0.0175 (8)	
O2	0.4788 (6)	0.5964 (5)	0.6393 (4)	0.0156 (8)	
O3	0.2680 (6)	0.1539 (6)	0.5394 (4)	0.0168 (8)	
O4	0.1538 (6)	-0.0817 (6)	0.6500 (5)	0.0191 (8)	
O5	-0.0130 (7)	0.2517 (6)	0.0669 (4)	0.0191 (8)	
O6	-0.1324 (7)	0.0169 (6)	-0.1924 (5)	0.0210 (9)	
O7	0.2703 (7)	0.6881 (6)	0.3356 (5)	0.0218 (9)	
H7A	0.403 (5)	0.763 (10)	0.370 (8)	0.033*	
H7B	0.218 (12)	0.757 (10)	0.281 (8)	0.033*	
C1	0.3614 (9)	0.6107 (8)	0.7255 (7)	0.0144 (11)	
C2	0.4651 (10)	0.7225 (9)	0.9107 (7)	0.0193 (12)	
H2A	0.4012	0.8285	0.9405	0.023*	0.75
H2B	0.6218	0.7953	0.9448	0.023*	0.75
H2C	0.3517	0.7421	0.9503	0.023*	0.25
H2D	0.5696	0.8616	0.9373	0.023*	0.25
C3A	0.4343 (15)	0.5720 (14)	1.0070 (10)	0.0262 (17)	0.75
H3A1	0.4800	0.6550	1.1242	0.031*	0.75
H3A2	0.2790	0.4864	0.9633	0.031*	0.75
C3B	0.588 (4)	0.598 (4)	1.005 (3)	0.0262 (17)	0.25
H3B1	0.6830	0.5544	0.9531	0.031*	0.25

H3B2	0.6791	0.6879	1.1213	0.031*	0.25
C4	0.1222 (8)	0.0203 (8)	0.5547 (6)	0.0140 (11)	
C5	-0.0433 (9)	0.0774 (8)	-0.0367 (7)	0.0143 (11)	

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Gd	0.00971 (18)	0.01274 (17)	0.00989 (17)	0.00183 (12)	0.00225 (12)	0.00217 (11)
O1	0.016 (2)	0.0173 (19)	0.0148 (19)	0.0059 (16)	0.0033 (16)	0.0017 (16)
O2	0.013 (2)	0.0160 (19)	0.0132 (19)	0.0034 (16)	0.0050 (16)	0.0007 (15)
O3	0.011 (2)	0.0182 (19)	0.0161 (19)	0.0011 (16)	0.0029 (16)	0.0050 (16)
O4	0.013 (2)	0.021 (2)	0.020 (2)	0.0031 (17)	0.0015 (16)	0.0105 (17)
O5	0.021 (2)	0.018 (2)	0.0131 (19)	0.0066 (17)	0.0035 (16)	0.0023 (16)
O6	0.027 (2)	0.021 (2)	0.014 (2)	0.0086 (18)	0.0061 (18)	0.0053 (17)
O7	0.017 (2)	0.020 (2)	0.029 (2)	0.0045 (18)	0.0081 (19)	0.0134 (18)
C1	0.016 (3)	0.011 (2)	0.013 (3)	0.006 (2)	0.002 (2)	0.005 (2)
C2	0.015 (3)	0.020 (3)	0.018 (3)	0.003 (2)	0.006 (2)	0.003 (2)
C3A	0.026 (4)	0.037 (4)	0.016 (4)	0.013 (4)	0.009 (3)	0.007 (4)
C3B	0.026 (4)	0.037 (4)	0.016 (4)	0.013 (4)	0.009 (3)	0.007 (4)
C4	0.009 (3)	0.016 (3)	0.012 (3)	0.003 (2)	0.002 (2)	0.001 (2)
C5	0.008 (3)	0.016 (3)	0.016 (3)	0.001 (2)	0.004 (2)	0.003 (2)

Geometric parameters (Å, °)

Gd—O1	2.575 (4)	O7—H7B	0.83 (7)
Gd—O1 ⁱ	2.473 (4)	C1—C2	1.489 (8)
Gd—O2	2.569 (4)	C2—C3A	1.538 (10)
Gd—O2 ⁱⁱ	2.430 (4)	C2—C3B	1.57 (3)
Gd—O3	2.403 (4)	C2—H2A	0.9700
Gd—O4 ⁱⁱⁱ	2.385 (4)	C2—H2B	0.9700
Gd—O5	2.374 (4)	C2—H2C	0.9700
Gd—O6 ^{iv}	2.540 (4)	C2—H2D	0.9700
Gd—O7	2.420 (4)	C3A—C3A ^v	1.510 (16)
O1—C1	1.274 (7)	C3A—H3A1	0.9700
O2—C1	1.279 (6)	C3A—H3A2	0.9700
O3—C4	1.252 (7)	C3B—C3B ^v	1.54 (5)
O4—C4	1.249 (6)	C3B—H3B1	0.9700
O5—C5	1.253 (6)	C3B—H3B2	0.9700
O6—C5	1.246 (7)	C4—C4 ⁱⁱⁱ	1.558 (10)
O7—H7A	0.84 (6)	C5—C5 ^{iv}	1.546 (10)
O5—Gd—O4 ⁱⁱⁱ	89.95 (13)	C4—O3—Gd	118.6 (3)
O5—Gd—O3	133.53 (13)	C4—O4—Gd ⁱⁱⁱ	119.1 (3)
O4 ⁱⁱⁱ —Gd—O3	68.29 (13)	C5—O5—Gd	123.7 (3)
O5—Gd—O7	78.95 (14)	C5—O6—Gd ^{iv}	117.9 (3)
O4 ⁱⁱⁱ —Gd—O7	142.92 (13)	Gd—O7—H7A	122 (5)
O3—Gd—O7	141.60 (14)	Gd—O7—H7B	139 (5)
O5—Gd—O2 ⁱⁱ	92.50 (13)	H7A—O7—H7B	97 (7)

O4 ⁱⁱⁱ —Gd—O2 ⁱⁱ	142.01 (12)	O1—C1—O2	118.3 (5)
O3—Gd—O2 ⁱⁱ	83.07 (12)	O1—C1—C2	120.4 (5)
O7—Gd—O2 ⁱⁱ	74.35 (13)	O2—C1—C2	121.2 (5)
O5—Gd—O1 ⁱ	81.91 (12)	O1—C1—Gd	59.4 (3)
O4 ⁱⁱⁱ —Gd—O1 ⁱ	69.18 (13)	O2—C1—Gd	59.1 (3)
O3—Gd—O1 ⁱ	122.76 (12)	C2—C1—Gd	174.0 (4)
O7—Gd—O1 ⁱ	74.25 (14)	C1—C2—C3A	112.9 (5)
O2 ⁱⁱ —Gd—O1 ⁱ	148.60 (13)	C1—C2—C3B	112.6 (10)
O5—Gd—O6 ^{iv}	65.44 (13)	C1—C2—H2A	109.0
O4 ⁱⁱⁱ —Gd—O6 ^{iv}	70.90 (14)	C3A—C2—H2A	109.0
O3—Gd—O6 ^{iv}	68.66 (13)	C3B—C2—H2A	135.3
O7—Gd—O6 ^{iv}	131.77 (14)	C1—C2—H2B	109.0
O2 ⁱⁱ —Gd—O6 ^{iv}	75.79 (13)	C3A—C2—H2B	109.0
O1 ⁱ —Gd—O6 ^{iv}	127.57 (13)	C3B—C2—H2B	73.4
O5—Gd—O2	147.43 (13)	H2A—C2—H2B	107.8
O4 ⁱⁱⁱ —Gd—O2	122.50 (13)	C1—C2—H2C	109.1
O3—Gd—O2	69.27 (13)	C3A—C2—H2C	73.3
O7—Gd—O2	73.02 (13)	C3B—C2—H2C	109.1
O2 ⁱⁱ —Gd—O2	64.38 (14)	H2B—C2—H2C	136.9
O1 ⁱ —Gd—O2	105.56 (12)	C1—C2—H2D	109.1
O6 ^{iv} —Gd—O2	124.28 (12)	C3A—C2—H2D	134.9
O5—Gd—O1	147.09 (13)	C3B—C2—H2D	109.1
O4 ⁱⁱⁱ —Gd—O1	80.01 (13)	H2C—C2—H2D	107.8
O3—Gd—O1	71.21 (13)	C3A ^v —C3A—C2	112.2 (8)
O7—Gd—O1	90.34 (13)	C3A ^v —C3A—H2C	148.3
O2 ⁱⁱ —Gd—O1	114.63 (12)	C3A ^v —C3A—H3A1	109.2
O1 ⁱ —Gd—O1	65.20 (14)	C2—C3A—H3A1	109.2
O6 ^{iv} —Gd—O1	136.78 (12)	H2C—C3A—H3A1	91.7
O2—Gd—O1	50.45 (12)	C3A ^v —C3A—H3A2	109.2
O5—Gd—C1	159.72 (13)	C2—C3A—H3A2	109.2
O4 ⁱⁱⁱ —Gd—C1	100.86 (14)	H2C—C3A—H3A2	85.5
O3—Gd—C1	66.75 (14)	H3A1—C3A—H3A2	107.9
O7—Gd—C1	82.15 (15)	C3B ^v —C3B—C2	108 (2)
O2 ⁱⁱ —Gd—C1	89.45 (14)	C3B ^v —C3B—H3B1	110.1
O1 ⁱ —Gd—C1	85.94 (14)	C2—C3B—H3B1	110.1
O6 ^{iv} —Gd—C1	134.31 (13)	C3B ^v —C3B—H3B2	110.1
O2—Gd—C1	25.31 (13)	C2—C3B—H3B2	110.1
O1—Gd—C1	25.21 (14)	H3B1—C3B—H3B2	108.4
C1—O1—Gd ⁱ	133.0 (3)	O4—C4—O3	126.3 (5)
C1—O1—Gd	95.4 (3)	O4—C4—C4 ⁱⁱⁱ	117.1 (6)
Gd ⁱ —O1—Gd	114.80 (14)	O3—C4—C4 ⁱⁱⁱ	116.6 (6)
C1—O2—Gd ⁱⁱ	147.7 (3)	O6—C5—O5	127.0 (5)
C1—O2—Gd	95.5 (3)	O6—C5—C5 ^{iv}	116.6 (6)
Gd ⁱⁱ —O2—Gd	115.62 (14)	O5—C5—C5 ^{iv}	116.4 (6)
O5—Gd—O1—C1	-140.5 (3)	O3—Gd—O5—C5	-8.0 (5)
O4 ⁱⁱⁱ —Gd—O1—C1	145.4 (3)	O7—Gd—O5—C5	148.3 (4)
O3—Gd—O1—C1	75.1 (3)	O2 ⁱⁱ —Gd—O5—C5	74.8 (4)

O7—Gd—O1—C1	-70.5 (3)	O1 ⁱ —Gd—O5—C5	-136.2 (4)
O2 ⁱⁱ —Gd—O1—C1	2.4 (3)	O6 ^{iv} —Gd—O5—C5	1.7 (4)
O1 ⁱ —Gd—O1—C1	-143.0 (4)	O2—Gd—O5—C5	117.5 (4)
O6 ^{iv} —Gd—O1—C1	97.7 (3)	O1—Gd—O5—C5	-138.6 (4)
O2—Gd—O1—C1	-3.1 (3)	C1—Gd—O5—C5	170.0 (4)
O5—Gd—O1—Gd ⁱ	2.5 (3)	Gd ⁱ —O1—C1—O2	-126.2 (4)
O4 ⁱⁱⁱ —Gd—O1—Gd ⁱ	-71.57 (16)	Gd—O1—C1—O2	5.5 (5)
O3—Gd—O1—Gd ⁱ	-141.88 (18)	Gd ⁱ —O1—C1—C2	55.3 (6)
O7—Gd—O1—Gd ⁱ	72.44 (17)	Gd—O1—C1—C2	-173.1 (4)
O2 ⁱⁱ —Gd—O1—Gd ⁱ	145.34 (15)	Gd ⁱ —O1—C1—Gd	-131.6 (4)
O1 ⁱ —Gd—O1—Gd ⁱ	-0.001 (2)	Gd ⁱⁱ —O2—C1—O1	-170.9 (4)
O6 ^{iv} —Gd—O1—Gd ⁱ	-119.29 (18)	Gd—O2—C1—O1	-5.5 (5)
O2—Gd—O1—Gd ⁱ	139.9 (2)	Gd ⁱⁱ —O2—C1—C2	7.6 (9)
C1—Gd—O1—Gd ⁱ	143.0 (4)	Gd—O2—C1—C2	173.0 (4)
O5—Gd—O2—C1	139.9 (3)	Gd ⁱⁱ —O2—C1—Gd	-165.4 (6)
O4 ⁱⁱⁱ —Gd—O2—C1	-34.5 (3)	O5—Gd—C1—O1	86.4 (5)
O3—Gd—O2—C1	-79.2 (3)	O4 ⁱⁱⁱ —Gd—C1—O1	-34.7 (3)
O7—Gd—O2—C1	108.2 (3)	O3—Gd—C1—O1	-95.2 (3)
O2 ⁱⁱ —Gd—O2—C1	-171.4 (4)	O7—Gd—C1—O1	107.9 (3)
O1 ⁱ —Gd—O2—C1	40.5 (3)	O2 ⁱⁱ —Gd—C1—O1	-177.9 (3)
O6 ^{iv} —Gd—O2—C1	-122.4 (3)	O1 ⁱ —Gd—C1—O1	33.2 (3)
O1—Gd—O2—C1	3.1 (3)	O6 ^{iv} —Gd—C1—O1	-108.5 (3)
O5—Gd—O2—Gd ⁱⁱ	-48.6 (3)	O2—Gd—C1—O1	174.4 (5)
O4 ⁱⁱⁱ —Gd—O2—Gd ⁱⁱ	136.96 (15)	O5—Gd—C1—O2	-88.0 (5)
O3—Gd—O2—Gd ⁱⁱ	92.21 (16)	O4 ⁱⁱⁱ —Gd—C1—O2	150.9 (3)
O7—Gd—O2—Gd ⁱⁱ	-80.42 (17)	O3—Gd—C1—O2	90.4 (3)
O2 ⁱⁱ —Gd—O2—Gd ⁱⁱ	0.0	O7—Gd—C1—O2	-66.5 (3)
O1 ⁱ —Gd—O2—Gd ⁱⁱ	-148.11 (15)	O2 ⁱⁱ —Gd—C1—O2	7.7 (3)
O6 ^{iv} —Gd—O2—Gd ⁱⁱ	49.0 (2)	O1 ⁱ —Gd—C1—O2	-141.2 (3)
O1—Gd—O2—Gd ⁱⁱ	174.5 (2)	O6 ^{iv} —Gd—C1—O2	77.1 (3)
C1—Gd—O2—Gd ⁱⁱ	171.4 (4)	O1—Gd—C1—O2	-174.4 (5)
O5—Gd—O3—C4	-72.7 (4)	O1—C1—C2—C3A	66.3 (7)
O4 ⁱⁱⁱ —Gd—O3—C4	-5.0 (4)	Gd ⁱⁱⁱ —O4—C4—O3	-175.4 (4)
O7—Gd—O3—C4	146.7 (3)	Gd ⁱⁱⁱ —O4—C4—C4 ⁱⁱⁱ	5.3 (7)
O2 ⁱⁱ —Gd—O3—C4	-159.5 (4)	Gd—O3—C4—O4	-175.0 (4)
O1 ⁱ —Gd—O3—C4	39.7 (4)	Gd—O3—C4—C4 ⁱⁱⁱ	4.3 (7)
O6 ^{iv} —Gd—O3—C4	-82.1 (4)	Gd ^{iv} —O6—C5—O5	178.4 (4)
O2—Gd—O3—C4	135.3 (4)	Gd ^{iv} —O6—C5—C5 ^{iv}	-0.3 (7)
O1—Gd—O3—C4	81.5 (4)	Gd—O5—C5—O6	179.1 (4)
C1—Gd—O3—C4	108.1 (4)	Gd—O5—C5—C5 ^{iv}	-2.1 (8)
O4 ⁱⁱⁱ —Gd—O5—C5	-67.3 (4)		

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

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O7—H7A \cdots O3 ⁱⁱ	0.84 (6)	2.00 (5)	2.799 (4)	160

$O7-H7B\cdots O6^{vi}$	0.83 (7)	2.07 (7)	2.888 (5)	168
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Symmetry codes: (ii) $-x+1, -y+1, -z+1$; (vi) $-x, -y+1, -z$.