



Crystal structure of rubidium peroxide ammonia disolvate

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Received 6 December 2016

Accepted 9 January 2017

Edited by T. J. Prior, University of Hull, England

Keywords: crystal structure; rubidium; peroxide; ammonia disolvate.

CCDC reference: 1526250

Supporting information: this article has supporting information at journals.iucr.org/e

The title compound, $\text{Rb}_2\text{O}_2 \cdot 2\text{NH}_3$, has been obtained as a reaction product of rubidium metal dissolved in liquid ammonia and glucuronic acid. As a result of the low-temperature crystallization, a disolvate was formed. To our knowledge, only one other solvate of an alkali metal peroxide is known: $\text{Na}_2\text{O}_2 \cdot 8\text{H}_2\text{O}$ has been reported by Grehl *et al.* [*Acta Cryst.* (1995), **C51**, 1038–1040]. We determined the peroxide bond length to be 1.530 (11) Å, which is in accordance with the length reported by Bremm & Jansen [*Z. Anorg. Allg. Chem.* (1992), **610**, 64–66]. One of the ammonia solvate molecules is disordered relative to a mirror plane, with 0.5 occupancy for the corresponding nitrogen atom.

1. Chemical context

The crystal structure of the title compound was determined in the course of investigations regarding the reactivity of carbohydrates towards alkali metals and NH_3 in solutions where liquid ammonia itself is used as solvent. The source of the peroxide anion could not be explicitly traced back but it seems to have its origin in oxygen gas from intruding atmosphere due to undetected leakage in the reaction vessel.

2. Structural commentary

The asymmetric unit contains one peroxide anion, two charge-compensating rubidium cations and two ammonia molecules (Fig. 1). Except for one nitrogen atom (N1, showing half-occupancy) and one hydrogen atom (H2B), all other atoms are located on mirror planes. The anion is surrounded by four rubidium cations located around the girth of the peroxide ion (Fig. 2). This unit forms one-dimensional infinite strands by sharing one common edge of a distorted plane of four Rb ions

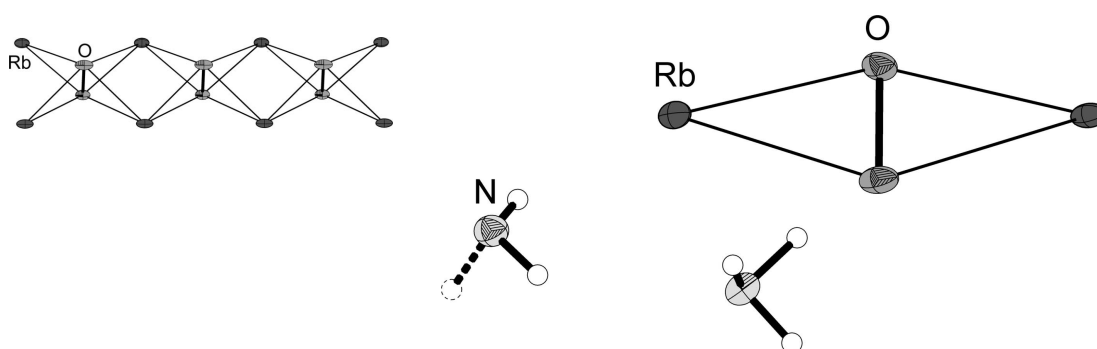


Figure 1

The asymmetric unit of the title compound, with the atom labeling and displacement ellipsoids drawn at the 50% probability level.

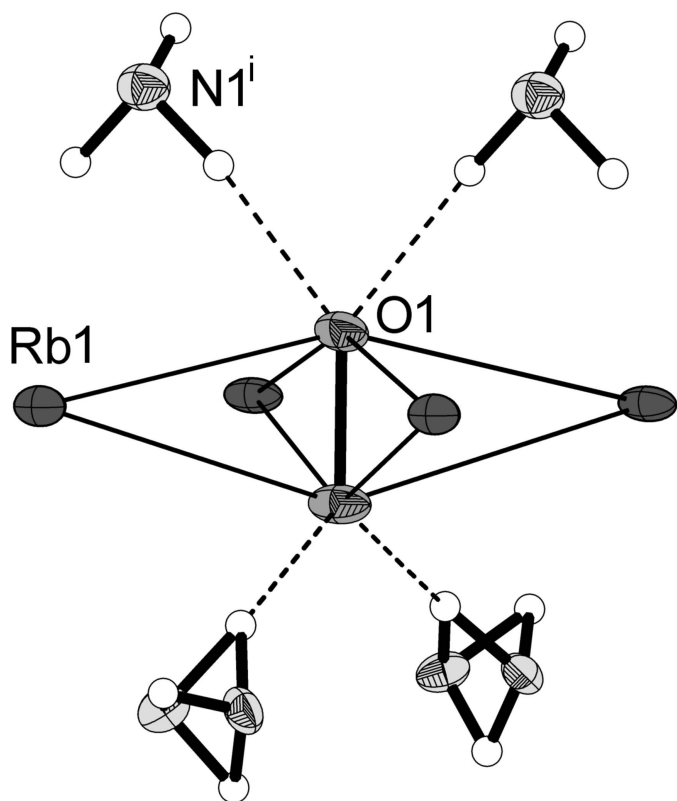


Figure 2
The environment of the peroxide anion. Displacement ellipsoids are drawn at the 50% probability level. [Symmetry code: (i) $-1 + x, y, z$.]

(Fig. 3). This structural motif can also be observed in potassium acetylide K_2C_2 (Hamberger *et al.*, 2012). The peroxide bond length was determined to be 1.530 (11) Å. The anion-cation contacts range between 2.790 (5) Å and 2.917 (6) Å. The coordination number of the cations is 8.

The O—O bond length of the peroxide anion is longer than the value found in the literature based on the work of Föppl which is approximately 1.49 Å. In Fig. 4, a comparative view of bond lengths is presented based on the work of Bremm & Jansen (1992), Föppl (1954, 1955, 1957) and Grehl *et al.* (1995).

3. Supramolecular features

Despite the low ammonia content, numerous hydrogen bonds can be observed and the NH_3 molecules bridge the peroxide

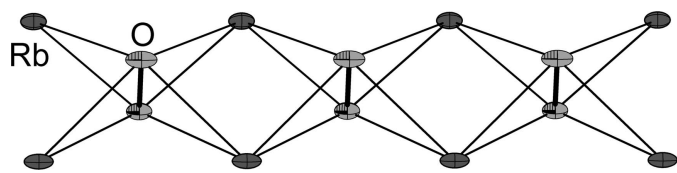


Figure 3
One-dimensional infinite strands formed by peroxide anions and rubidium cations. Displacement ellipsoids are drawn at the 50% probability level.

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N1-H1A\cdots O2$	1.05 (14)	1.98 (15)	2.941 (16)	151 (8)
$N1-H1B\cdots O2^i$	0.97 (13)	2.04 (15)	2.926 (15)	151 (8)
$N1-H1C\cdots O1^{iii}$	0.82 (14)	3.07 (16)	3.597 (16)	125 (11)
$N2-H2A\cdots N2^{iii}$	0.74 (16)	3.03 (12)	3.57 (2)	131 (6)
$N2-H2A\cdots N2^{iv}$	0.74 (16)	3.03 (12)	3.57 (2)	131 (6)
$N2-H2A\cdots N2^v$	0.74 (16)	3.03 (12)	3.57 (2)	131 (6)
$N2-H2B\cdots O1^{ii}$	1.01 (11)	1.95 (11)	2.955 (10)	173 (8)

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

anions. The peroxide anion shows five contacts to ammonia molecules, forming a three-dimensional network in the packing. The distances between donor and acceptor atoms ranges from 2.926(15) Angstrom to 3.597(16) Angstrom, which is commonly observed in ammoniates. Numerical details of the hydrogen-bonding interactions are given in Table 1.

4. Synthesis and crystallization

500 mg (2.58 mmol) D-glucuronic acid and 880 mg (10.29 mmol) rubidium were placed under an argon atmo-

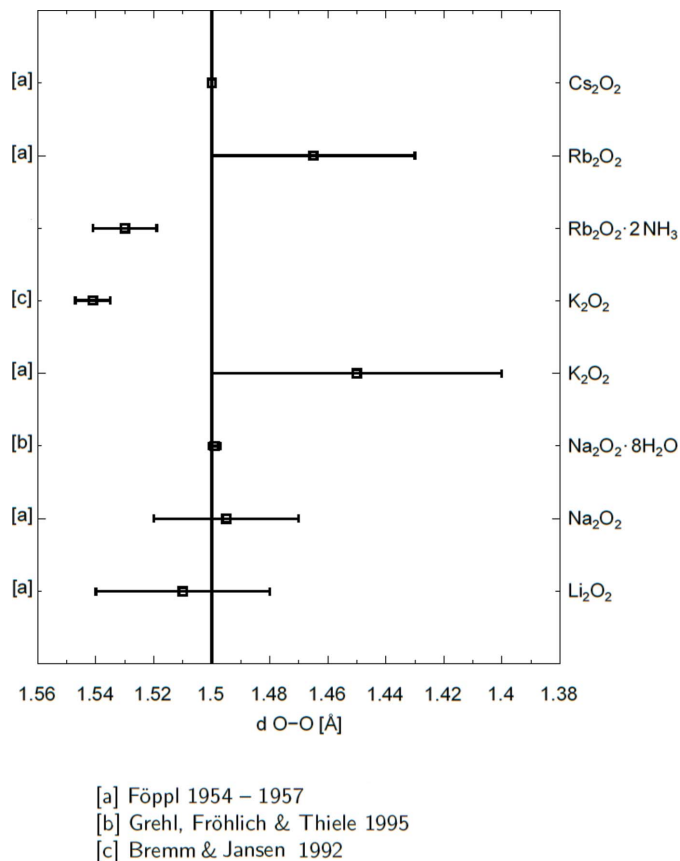


Figure 4
Comparison of peroxide bond lengths in different compounds. The vertical line shows the peroxide bond length commonly used in the literature. Each data point is shown with its standard uncertainties.

Table 2
Experimental details.

Crystal data	
Chemical formula	Rb ₂ O ₂ ·2NH ₃
<i>M_r</i>	237.01
Crystal system, space group	Orthorhombic, <i>Pnma</i>
Temperature (K)	123
<i>a</i> , <i>b</i> , <i>c</i> (Å)	7.3957 (7), 4.0932 (6), 18.1873 (17)
<i>V</i> (Å ³)	550.57 (11)
<i>Z</i>	4
Radiation type	Mo <i>K</i> α
<i>μ</i> (mm ⁻¹)	17.66
Crystal size (mm)	0.24 × 0.09 × 0.08
Data collection	
Diffractometer	Agilent SuperNova Dual Source diffractometer with an Eos detector
Absorption correction	Analytical [<i>CrysAlis PRO</i> (Agilent, 2012), based on expressions derived by Clark & Reid (1995)]
<i>T_{min}</i> , <i>T_{max}</i>	0.064, 0.354
No. of measured, independent and observed [<i>I</i> > 2σ(<i>I</i>)] reflections	2921, 641, 570
<i>R_{int}</i>	0.057
(sin θ/λ) _{max} (Å ⁻¹)	0.625
Refinement	
<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.050, 0.118, 1.35
No. of reflections	641
No. of parameters	51
H-atom treatment	Only H-atom coordinates refined
Δρ _{max} , Δρ _{min} (e Å ⁻³)	1.12, -1.79

Computer programs: *CrysAlis PRO* (Agilent, 2012), *olex2.solve* (Bourhis *et al.*, 2015), *SHELXL2014* (Sheldrick, 2015), *DIAMOND* (Brandenburg & Putz, 2012) and *OLEX2* (Dolomanov *et al.*, 2009).

sphere in a reaction vessel and 25 ml of dry liquid ammonia were condensed. The mixture was stored at 237 K for five days.

The flask was then stored at 161 K for several months. After that period, clear needle-shaped colorless crystals of the title compound could be found at the wall of the flask.

5. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. The nitrogen atom N1 is disordered with 0.5 as the site occupation factor. All hydrogen atoms could be located in difference map and their positions were refined freely with a common *U*_{iso}(H) parameter. The isotropic displacement parameters were fixed to 0.025.

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supporting information

Acta Cryst. (2017). E73, 200-202 [https://doi.org/10.1107/S2056989017000354]

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Computing details

Data collection: *CrysAlis PRO* (Agilent, 2012); cell refinement: *CrysAlis PRO* (Agilent, 2012); data reduction: *CrysAlis PRO* (Agilent, 2012); program(s) used to solve structure: *olex2.solve* (Bourhis *et al.*, 2015); program(s) used to refine structure: *SHELXL2014* (Sheldrick, 2015); molecular graphics: *DIAMOND* (Brandenburg & Putz, 2012); software used to prepare material for publication: *OLEX2* (Dolomanov *et al.*, 2009).

Rubidium peroxide ammonia disolvate

Crystal data

Rb₂O₂·2NH₃

$M_r = 237.01$

Orthorhombic, *Pnma*

$a = 7.3957$ (7) Å

$b = 4.0932$ (6) Å

$c = 18.1873$ (17) Å

$V = 550.57$ (11) Å³

$Z = 4$

$F(000) = 440$

$D_x = 2.859$ Mg m⁻³

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

Cell parameters from 1711 reflections

$\theta = 3.4$ – 28.3°

$\mu = 17.66$ mm⁻¹

$T = 123$ K

Block, clear colourless

$0.24 \times 0.09 \times 0.08$ mm

Data collection

Agilent SuperNova Dual Source
diffractometer with an Eos detector

Mirror monochromator

Detector resolution: 7.9851 pixels mm⁻¹

phi and ω scans

Absorption correction: analytical

[*CrysAlis PRO* (Agilent, 2012), based on
expressions derived by Clark & Reid (1995)]

$T_{\min} = 0.064$, $T_{\max} = 0.354$

2921 measured reflections

641 independent reflections

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

$R_{\text{int}} = 0.057$

$\theta_{\max} = 26.4^\circ$, $\theta_{\min} = 3.6^\circ$

$h = -9 \rightarrow 9$

$k = -5 \rightarrow 4$

$l = -22 \rightarrow 22$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.050$

$wR(F^2) = 0.118$

$S = 1.35$

641 reflections

51 parameters

0 restraints

Hydrogen site location: difference Fourier map

Only H-atom coordinates refined

$w = 1/[\sigma^2(F_o^2) + (0.0361P)^2 + 7.6682P]$

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

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

$\Delta\rho_{\max} = 1.12$ e Å⁻³

$\Delta\rho_{\min} = -1.79$ e 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.

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)
Rb1	0.11196 (14)	0.2500	0.42075 (5)	0.0164 (4)	
Rb2	-0.25614 (15)	1.2500	0.28803 (5)	0.0182 (4)	
O1	-0.1266 (11)	0.7500	0.3825 (4)	0.019 (2)	
O2	0.0133 (11)	0.7500	0.3206 (4)	0.022 (2)	
N2	0.6843 (16)	0.2500	0.4710 (6)	0.020 (3)	
N1	0.4060 (18)	0.650 (4)	0.3327 (7)	0.017 (4)	0.5
H1A	0.28 (2)	0.7500	0.340 (8)	0.025*	
H1B	0.42 (2)	0.7500	0.285 (8)	0.025*	
H1C	0.46 (2)	0.7500	0.364 (8)	0.025*	
H2A	0.58 (2)	0.2500	0.472 (8)	0.025*	
H2B	0.739 (13)	0.43 (3)	0.441 (5)	0.025*	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Rb1	0.0107 (5)	0.0268 (8)	0.0116 (5)	0.000	-0.0022 (4)	0.000
Rb2	0.0088 (5)	0.0345 (8)	0.0113 (5)	0.000	0.0003 (4)	0.000
O1	0.012 (4)	0.027 (6)	0.020 (4)	0.000	0.003 (3)	0.000
O2	0.011 (4)	0.040 (7)	0.014 (4)	0.000	0.000 (3)	0.000
N2	0.017 (5)	0.025 (7)	0.018 (5)	0.000	0.000 (4)	0.000
N1	0.016 (6)	0.019 (12)	0.016 (6)	-0.007 (6)	-0.002 (5)	0.000 (6)

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

Rb1—Rb2 ⁱ	3.6383 (15)	Rb2—N1 ^x	3.507 (15)
Rb1—O1	2.790 (5)	Rb2—N1 ^{xiii}	2.988 (14)
Rb1—O1 ⁱⁱ	3.579 (8)	Rb2—N1 ^{ix}	2.988 (14)
Rb1—O1 ⁱ	2.790 (6)	O1—Rb1 ⁱⁱ	3.579 (8)
Rb1—O2	2.836 (5)	O1—Rb1 ^{viii}	2.790 (5)
Rb1—O2 ⁱ	2.836 (5)	O1—Rb2 ⁱ	2.839 (6)
Rb1—N2 ⁱⁱⁱ	3.292 (12)	O1—O2	1.530 (11)
Rb1—N2 ^{iv}	3.215 (8)	O2—Rb1 ^{viii}	2.836 (5)
Rb1—N2 ^v	3.215 (8)	O2—Rb2 ^{xiv}	3.316 (6)
Rb1—N2 ^{vi}	3.215 (8)	O2—Rb2 ^{xv}	3.316 (6)
Rb1—N1 ^{vii}	3.157 (14)	O2—Rb2 ⁱ	2.917 (6)
Rb1—N1	3.157 (14)	N2—Rb1 ^{iv}	3.215 (8)
Rb2—O1	2.839 (6)	N2—Rb1 ^{xvi}	3.292 (12)
Rb2—O1 ^{viii}	2.839 (6)	N2—Rb1 ^{xvii}	3.215 (8)
Rb2—O2 ^{viii}	2.917 (6)	N2—Rb2 ^{xviii}	3.356 (11)

Rb2—O2 ^{ix}	3.316 (6)	N1—Rb1 ^{viii}	3.652 (15)
Rb2—O2	2.917 (6)	N1—Rb2 ^{xiv}	2.988 (14)
Rb2—O2 ^x	3.316 (6)	N1—Rb2 ^{xv}	3.507 (15)
Rb2—N2 ^{xi}	3.356 (11)	N1—Rb2 ^{xvi}	3.598 (14)
Rb2—N1 ^{xi}	3.095 (15)	N1—Rb2 ^{xviii}	3.095 (15)
Rb2—N1 ^{xii}	3.095 (15)	N1—N1 ^{xix}	0.82 (3)
O1 ⁱⁱ —Rb1—Rb2 ⁱ	133.30 (13)	O2—Rb2—N1 ^x	53.2 (3)
O1 ⁱ —Rb1—Rb2 ⁱ	50.32 (13)	O2—Rb2—N1 ^{xi}	150.4 (3)
O1—Rb1—Rb2 ⁱ	50.32 (13)	O2—Rb2—N1 ^{xii}	97.3 (3)
O1 ⁱ —Rb1—O1 ⁱⁱ	105.56 (17)	O2 ^{ix} —Rb2—N1 ^x	103.6 (3)
O1—Rb1—O1 ⁱ	94.4 (2)	O2 ^{viii} —Rb2—N1 ^{ix}	59.4 (3)
O1—Rb1—O1 ⁱⁱ	105.56 (17)	O2 ^{viii} —Rb2—N1 ^x	112.7 (3)
O1 ⁱ —Rb1—O2	101.92 (19)	O2 ^{viii} —Rb2—N1 ^{xi}	97.3 (3)
O1 ⁱ —Rb1—O2 ⁱ	31.5 (2)	O2 ^x —Rb2—N1 ^x	51.0 (3)
O1—Rb1—O2	31.5 (2)	O2—Rb2—N1 ^{xiii}	59.4 (3)
O1—Rb1—O2 ⁱ	101.92 (19)	O2 ^{viii} —Rb2—N1 ^{xiii}	105.0 (3)
O1 ⁱ —Rb1—N2 ^v	156.3 (3)	O2—Rb2—N1 ^{ix}	105.0 (3)
O1 ⁱ —Rb1—N2 ⁱⁱⁱ	57.42 (17)	N2 ^{xi} —Rb2—N1 ^x	131.7 (3)
O1 ⁱ —Rb1—N2 ^{iv}	156.3 (3)	N1 ^{xi} —Rb2—O2 ^x	93.8 (3)
O1—Rb1—N2 ^v	89.0 (2)	N1 ^{xi} —Rb2—O2 ^{ix}	54.2 (3)
O1—Rb1—N2 ^{iv}	89.0 (2)	N1 ^{ix} —Rb2—O2 ^{ix}	55.3 (3)
O1—Rb1—N2 ^{vi}	156.3 (3)	N1 ^{xii} —Rb2—O2 ^{ix}	93.8 (3)
O1—Rb1—N2 ⁱⁱⁱ	57.42 (17)	N1 ^{xiii} —Rb2—O2 ^x	55.3 (3)
O1 ⁱ —Rb1—N2 ^{vi}	88.98 (19)	N1 ^{xii} —Rb2—O2 ^x	54.2 (3)
O1—Rb1—N1	85.9 (3)	N1 ^{xiii} —Rb2—O2 ^{ix}	96.1 (3)
O1 ⁱ —Rb1—N1 ^{vii}	85.9 (3)	N1 ^{ix} —Rb2—O2 ^x	96.1 (3)
O1 ⁱ —Rb1—N1	133.5 (3)	N1 ^{xii} —Rb2—N2 ^{xi}	68.5 (3)
O1—Rb1—N1 ^{vii}	133.5 (3)	N1 ^{xiii} —Rb2—N2 ^{xi}	141.3 (3)
O2 ⁱ —Rb1—Rb2 ⁱ	51.77 (13)	N1 ^{xi} —Rb2—N2 ^{xi}	68.5 (3)
O2—Rb1—Rb2 ⁱ	51.77 (13)	N1 ^{ix} —Rb2—N2 ^{xi}	141.3 (3)
O2—Rb1—O1 ⁱⁱ	130.55 (13)	N1 ^{xii} —Rb2—N1 ^{xi}	63.8 (6)
O2 ⁱ —Rb1—O1 ⁱⁱ	130.55 (13)	N1 ^{ix} —Rb2—N1 ^{xi}	103.1 (5)
O2—Rb1—O2 ⁱ	92.4 (2)	N1 ^{ix} —Rb2—N1 ^{xii}	143.7 (3)
O2—Rb1—N2 ^v	93.14 (17)	N1 ^{xiii} —Rb2—N1 ^x	11.3 (4)
O2 ⁱ —Rb1—N2 ^{iv}	166.8 (2)	N1 ^{xiii} —Rb2—N1 ^{ix}	66.4 (6)
O2 ⁱ —Rb1—N2 ^{vi}	93.14 (17)	N1 ^{ix} —Rb2—N1 ^x	77.7 (3)
O2—Rb1—N2 ⁱⁱⁱ	86.0 (2)	N1 ^{xiii} —Rb2—N1 ^{xii}	103.1 (5)
O2—Rb1—N2 ^{iv}	93.14 (17)	N1 ^{xii} —Rb2—N1 ^x	94.0 (2)
O2—Rb1—N2 ^{vi}	166.8 (2)	N1 ^{xiii} —Rb2—N1 ^{xi}	143.7 (3)
O2 ⁱ —Rb1—N2 ^v	166.8 (2)	N1 ^{xi} —Rb2—N1 ^x	144.2 (4)
O2 ⁱ —Rb1—N2 ⁱⁱⁱ	86.0 (2)	Rb1 ^{viii} —O1—Rb1 ⁱⁱ	74.44 (17)
O2—Rb1—N1	58.5 (3)	Rb1—O1—Rb1 ^{viii}	94.4 (2)
O2—Rb1—N1 ^{vii}	103.0 (3)	Rb1—O1—Rb1 ⁱⁱ	74.44 (17)
O2 ⁱ —Rb1—N1	103.0 (3)	Rb1—O1—Rb2 ⁱ	80.53 (7)
O2 ⁱ —Rb1—N1 ^{vii}	58.5 (3)	Rb1 ^{viii} —O1—Rb2	80.53 (7)
N2 ⁱⁱⁱ —Rb1—Rb2 ⁱ	57.66 (19)	Rb1 ^{viii} —O1—Rb2 ⁱ	153.3 (3)
N2 ^{vi} —Rb1—Rb2 ⁱ	139.21 (14)	Rb1—O1—Rb2	153.3 (3)

N2 ^{iv} —Rb1—Rb2 ⁱ	139.21 (14)	Rb2—O1—Rb1 ⁱⁱ	127.97 (15)
N2 ^v —Rb1—Rb2 ⁱ	139.21 (14)	Rb2 ⁱ —O1—Rb1 ⁱⁱ	127.97 (15)
N2 ^{iv} —Rb1—O1 ⁱⁱ	51.21 (19)	Rb2 ⁱ —O1—Rb2	92.3 (2)
N2 ^v —Rb1—O1 ⁱⁱ	51.21 (19)	O2—O1—Rb1 ⁱⁱ	135.7 (5)
N2 ⁱⁱⁱ —Rb1—O1 ⁱⁱ	75.6 (2)	O2—O1—Rb1	75.9 (3)
N2 ^{vi} —Rb1—O1 ⁱⁱ	51.21 (19)	O2—O1—Rb1 ^{viii}	75.9 (3)
N2 ^v —Rb1—N2 ⁱⁱⁱ	106.3 (2)	O2—O1—Rb2	77.4 (3)
N2 ^{vi} —Rb1—N2 ⁱⁱⁱ	106.3 (2)	O2—O1—Rb2 ⁱ	77.4 (3)
N2 ^{iv} —Rb1—N2 ⁱⁱⁱ	106.3 (2)	Rb1—O2—Rb1 ^{viii}	92.4 (2)
N2 ^v —Rb1—N2 ^{vi}	79.1 (2)	Rb1 ^{viii} —O2—Rb2	78.45 (8)
N2 ^v —Rb1—N2 ^{iv}	0.0 (5)	Rb1—O2—Rb2	144.3 (3)
N2 ^{iv} —Rb1—N2 ^{vi}	79.1 (2)	Rb1 ^{viii} —O2—Rb2 ^{xiv}	134.1 (3)
N1 ^{vii} —Rb1—Rb2 ⁱ	100.3 (2)	Rb1—O2—Rb2 ^{xiv}	78.75 (10)
N1—Rb1—Rb2 ⁱ	100.3 (2)	Rb1—O2—Rb2 ⁱ	78.45 (8)
N1—Rb1—O1 ⁱⁱ	119.1 (3)	Rb1—O2—Rb2 ^{xv}	134.1 (3)
N1 ^{vii} —Rb1—O1 ⁱⁱ	119.1 (3)	Rb1 ^{viii} —O2—Rb2 ⁱ	144.3 (3)
N1—Rb1—N2 ^v	70.0 (3)	Rb1 ^{viii} —O2—Rb2 ^{xv}	78.75 (10)
N1 ^{vii} —Rb1—N2 ^{vi}	70.0 (3)	Rb2 ⁱ —O2—Rb2	89.1 (2)
N1—Rb1—N2 ^{iv}	70.0 (3)	Rb2 ^{xiv} —O2—Rb2 ^{xv}	76.22 (17)
N1 ^{vii} —Rb1—N2 ^{iv}	108.5 (3)	Rb2—O2—Rb2 ^{xv}	78.31 (10)
N1—Rb1—N2 ^{vi}	108.5 (3)	Rb2 ⁱ —O2—Rb2 ^{xiv}	78.32 (10)
N1—Rb1—N2 ⁱⁱⁱ	143.4 (3)	Rb2—O2—Rb2 ^{xiv}	131.6 (3)
N1 ^{vii} —Rb1—N2 ⁱⁱⁱ	143.4 (3)	Rb2 ⁱ —O2—Rb2 ^{xv}	131.6 (3)
N1 ^{vii} —Rb1—N2 ^v	108.5 (3)	O1—O2—Rb1	72.6 (3)
N1—Rb1—N1 ^{vii}	62.4 (6)	O1—O2—Rb1 ^{viii}	72.6 (3)
O1—Rb2—O1 ^{viii}	92.3 (2)	O1—O2—Rb2	71.8 (3)
O1 ^{viii} —Rb2—O2 ^{ix}	95.11 (15)	O1—O2—Rb2 ^{xv}	141.86 (9)
O1—Rb2—O2 ^{viii}	98.78 (18)	O1—O2—Rb2 ^{xiv}	141.86 (9)
O1—Rb2—O2 ^x	95.11 (15)	O1—O2—Rb2 ⁱ	71.8 (3)
O1 ^{viii} —Rb2—O2 ^x	168.22 (18)	Rb1 ^{iv} —N2—Rb1 ^{xvi}	73.7 (2)
O1—Rb2—O2 ^{ix}	168.22 (18)	Rb1 ^{iv} —N2—Rb1 ^{xvii}	79.1 (2)
O1 ^{viii} —Rb2—O2 ^{viii}	30.8 (2)	Rb1 ^{xvii} —N2—Rb1 ^{xvi}	73.7 (2)
O1 ^{viii} —Rb2—O2	98.77 (18)	Rb1 ^{xvi} —N2—Rb2 ^{xviii}	66.4 (2)
O1—Rb2—O2	30.8 (2)	Rb1 ^{xvii} —N2—Rb2 ^{xviii}	123.1 (3)
O1—Rb2—N2 ^{xi}	56.23 (17)	Rb1 ^{iv} —N2—Rb2 ^{xviii}	123.1 (3)
O1 ^{viii} —Rb2—N2 ^{xi}	56.23 (17)	Rb1—N1—Rb1 ^{viii}	73.5 (3)
O1 ^{viii} —Rb2—N1 ^{ix}	85.1 (3)	Rb1—N1—Rb2 ^{xvi}	161.2 (5)
O1—Rb2—N1 ^{ix}	134.7 (3)	Rb1—N1—Rb2 ^{xv}	116.4 (4)
O1—Rb2—N1 ^x	76.0 (3)	Rb2 ^{xiv} —N1—Rb1	79.2 (4)
O1—Rb2—N1 ^{xi}	119.6 (3)	Rb2 ^{xvi} —N1—Rb1 ^{viii}	93.0 (4)
O1—Rb2—N1 ^{xii}	74.5 (3)	Rb2 ^{xv} —N1—Rb1 ^{viii}	66.4 (3)
O1 ^{viii} —Rb2—N1 ^{xiii}	134.7 (3)	Rb2 ^{xviii} —N1—Rb1	114.5 (5)
O1 ^{viii} —Rb2—N1 ^{xii}	119.6 (3)	Rb2 ^{xiv} —N1—Rb1 ^{viii}	116.9 (4)
O1 ^{viii} —Rb2—N1 ^{xi}	74.5 (3)	Rb2 ^{xviii} —N1—Rb1 ^{viii}	162.1 (5)
O1 ^{viii} —Rb2—N1 ^x	140.3 (3)	Rb2 ^{xviii} —N1—Rb2 ^{xvi}	75.0 (3)
O1—Rb2—N1 ^{xiii}	85.1 (3)	Rb2 ^{xviii} —N1—Rb2 ^{xv}	118.8 (4)
O2 ^x —Rb2—O2 ^{ix}	76.22 (17)	Rb2 ^{xv} —N1—Rb2 ^{xvi}	67.5 (3)
O2—Rb2—O2 ^{ix}	154.83 (12)	Rb2 ^{xiv} —N1—Rb2 ^{xv}	77.7 (3)

O2 ^{viii} —Rb2—O2 ^{ix}	92.24 (11)	Rb2 ^{xiv} —N1—Rb2 ^{xvi}	119.1 (4)
O2 ^{viii} —Rb2—O2	89.1 (2)	Rb2 ^{xiv} —N1—Rb2 ^{xviii}	80.9 (4)
O2 ^{viii} —Rb2—O2 ^x	154.83 (12)	N1 ^{xix} —N1—Rb1 ^{viii}	47.7 (2)
O2—Rb2—O2 ^x	92.24 (11)	N1 ^{xix} —N1—Rb1	121.2 (3)
O2 ^{viii} —Rb2—N2 ^{xi}	83.6 (2)	N1 ^{xix} —N1—Rb2 ^{xv}	45.5 (2)
O2 ^{ix} —Rb2—N2 ^{xi}	121.53 (19)	N1 ^{xix} —N1—Rb2 ^{xvi}	46.9 (3)
O2 ^x —Rb2—N2 ^{xi}	121.53 (19)	N1 ^{xix} —N1—Rb2 ^{xiv}	123.2 (3)
O2—Rb2—N2 ^{xi}	83.6 (2)	N1 ^{xix} —N1—Rb2 ^{xviii}	121.9 (3)
O2 ^{viii} —Rb2—N1 ^{xii}	150.4 (3)		

Symmetry codes: (i) $x, y-1, z$; (ii) $-x, -y+1, -z+1$; (iii) $x-1, y, z$; (iv) $-x+1, -y+1, -z+1$; (v) $-x+1, y+1/2, -z+1$; (vi) $-x+1, y-1/2, -z+1$; (vii) $x, -y+1/2, z$; (viii) $x, y+1, z$; (ix) $x-1/2, y+1, -z+1/2$; (x) $x-1/2, y, -z+1/2$; (xi) $x-1, y+1, z$; (xii) $x-1, -y+3/2, z$; (xiii) $x-1/2, -y+3/2, -z+1/2$; (xiv) $x+1/2, y-1, -z+1/2$; (xv) $x+1/2, y, -z+1/2$; (xvi) $x+1, y, z$; (xvii) $-x+1, -y, -z+1$; (xviii) $x+1, y-1, z$; (xix) $x, -y+3/2, z$.

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1A \cdots O2	1.05 (14)	1.98 (15)	2.941 (16)	151 (8)
N1—H1B \cdots O2 ^{xx}	0.97 (13)	2.04 (15)	2.926 (15)	151 (8)
N1—H1C \cdots O1 ^{xvi}	0.82 (14)	3.07 (16)	3.597 (16)	125 (11)
N2—H2A \cdots N2 ^{iv}	0.74 (16)	3.03 (12)	3.57 (2)	131 (6)
N2—H2A \cdots N2 ^{vi}	0.74 (16)	3.03 (12)	3.57 (2)	131 (6)
N2—H2A \cdots N2 ^v	0.74 (16)	3.03 (12)	3.57 (2)	131 (6)
N2—H2B \cdots O1 ^{xvi}	1.01 (11)	1.95 (11)	2.955 (10)	173 (8)

Symmetry codes: (iv) $-x+1, -y+1, -z+1$; (v) $-x+1, y+1/2, -z+1$; (vi) $-x+1, y-1/2, -z+1$; (xvi) $x+1, y, z$; (xx) $x+1/2, -y+3/2, -z+1/2$.