metal-organic compounds

Aquabromidobis(dimethylglyoximato)-cobalt(III)

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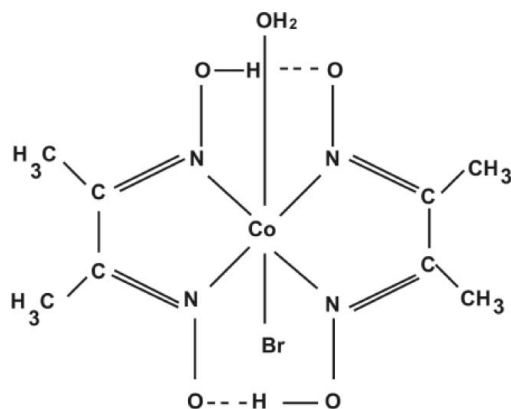Correspondence e-mail: dayalan77@gmail.com

In the title complex, $[CoBr(C_4H_7N_2O_2)_2(H_2O)]$, a crystallographic mirror plane bisects the molecule, perpendicular to the glyoxime ligands. The geometry around the cobalt(III) atom is approximately octahedral with the four glyoxime N atoms forming the square base. A bromide ion and the O atom of a water molecule occupy the remaining coordination sites. The N—Co—N bite angles are 82.18 (4) and 80.03 (16)°. The glyoximate moieties form strong intramolecular O—H···O hydrogen bonds. The coordinated water molecule forms an intermolecular O—H···O hydrogen bond with a glyoxime O atom, thereby generating supramolecular chains parallel to [010].

Related literature

For related complexes, see: Ohkubo & Fukuzumi (2005); Randall & Alberty (1970); Schrauzer (1968); Trommel et al. (2001). For similar structures, see: Bernstein et al. (1995); Mégnamisi-Bélombé et al. (1983); Meera et al. (2009); Ramesh et al. (2008). For the preparation of similar complexes, see: Vijayraghavan & Dayalan (1992). For spectroscopic studies related to the title complex, see: Folgando et al. (1986); Khan et al. (1997); Lopez et al. (1986).

Experimental

Crystal data

$[CoBr(C_4H_7N_2O_2)_2(H_2O)]$	V = 706.09 (5) Å^3
M_r = 387.09	Z = 2
Monoclinic, $P2_1/m$	Mo $K\alpha$ radiation
a = 7.5903 (3) Å	μ = 4.07 mm^{-1}
b = 8.8816 (4) Å	T = 293 K
c = 10.5343 (5) Å	0.15 × 0.10 × 0.10 mm
β = 96.137 (3)°	

Data collection

Bruker Kappa APEXII CCD diffractometer	7395 measured reflections
Absorption correction: multi-scan (SADABS; Bruker 1999)	1480 independent reflections
	1298 reflections with $I > 2\sigma(I)$
T_{min} = 0.581, T_{max} = 0.687	R_{int} = 0.028

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.034	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2)$ = 0.096	
S = 1.22	
1480 reflections	$\Delta\rho_{max}$ = 1.01 e Å^{-3}
101 parameters	$\Delta\rho_{min}$ = −0.54 e Å^{-3}
2 restraints	

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

D—H···A	D—H	H···A	D···A	D—H···A
O2—H2···O1^i	0.92 (1)	1.58 (1)	2.494 (3)	169 (4)
O3—H3···O1^{ii}	0.85 (3)	1.79 (3)	2.616 (3)	167 (4)

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

Data collection: APEX2 (Bruker, 2004); cell refinement: APEX2 and SAINT-Plus (Bruker, 2004); data reduction: SAINT-Plus and XPREP (Bruker, 2004); program(s) used to solve structure: SIR92 (Altomare et al., 1993); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 (Farrugia, 1997) and Mercury (Macrae et al., 2008); software used to prepare material for publication: PLATON (Spek, 2009).

References

Altomare, A., Cascarano, G., Giacovazzo, C. & Guagliardi, A. (1993). J. Appl. Cryst. **26**, 343–350.
Bernstein, J., Davis, R. E., Shimoni, L. & Chang, N.-L. (1995). Angew. Chem. Int. Ed. Engl. **34**, 1555–1573.
Bruker (1999). SADABS. Bruker AXS Inc., Madison, Wisconsin, USA.
Bruker (2004). APEX2 and SAINT-Plus. Bruker AXS Inc., Madison, Wisconsin, USA.
Farrugia, L. J. (1997). J. Appl. Cryst. **30**, 565.
Folgando, J. V., Coronado, E. & Bltran, D. (1986). J. Chem. Soc. Dalton Trans. 1, pp. 1061–1064.

Khan, T. A., Shahjahan, & Zaidi, S. A. A. (1997). *Indian J. Chem. Sect. A*, **36**, 153–156.

Lopez, C., Alvarez, S., Solans, X. & Font-Altaba, M. (1986). *Inorg. Chem.* **25**, 2962–2969.

Macrae, C. F., Bruno, I. J., Chisholm, J. A., Edgington, P. R., McCabe, P., Pidcock, E., Rodriguez-Monge, L., Taylor, R., van de Streek, J. & Wood, P. A. (2008). *J. Appl. Cryst.* **41**, 466–470.

Meera, P., Revathi, C. & Dayalan, A. (2009). *Acta Cryst.* E**65**, m140–m141.

Mégnamisi-Bélombé, M., Endres, H. & Rossato, E. (1983). *Acta Cryst.* C**39**, 705–707.

Ohkubo, K. & Fukuzumi, S. (2005). *J. Phys. Chem.* **109**, 1105–1113.

Ramesh, P., SubbiahPandi, A., Jothi, P., Revathi, C. & Dayalan, A. (2008). *Acta Cryst.* E**64**, m300–m301.

Randall, W. C. & Alberty, R. A. (1970). *Biochemistry*, **9**, 1886–1892.

Schrauzer, G. N. (1968). *Acc. Chem. Res.* **1**, 97–103.

Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.

Spek, A. L. (2009). *Acta Cryst.* D**65**, 148–155.

Trommel, J. S., Warncke, K. & Marzilli, L. G. (2001). *J. Am. Chem. Soc.* **123**, 3358–3366.

Vijayraghavan, V. R. & Dayalan, A. (1992). *J. Indian Chem. Soc.* **69**, 383–384.

supporting information

Aquabromidobis(dimethylglyoximato)cobalt(III)

Parthasarathy Meera, Madhavan Amutha Selvi, Pachaimuthu Jothi and Arunachalam Dayalan

S1. Comment

A number of cobalt complexes have been proposed as model systems for vitamin-B_{12}(Trommel *et al.*, 2001; Ohkubo & Fukuzumi, 2005). The most commonly mentioned model system is bis(dimethylglyoximato)cobalt(III) complexes on which Schrauzer has carried out a great amount of research.The common feature of the different models is that each possesses a strong equatorial ligand field (Schrauzer,1968). A variety of cobalt(III) complexes have been discovered possessing stable axial cobalt-carbon bonds. Simple alkyl cobaloximes, are thermally stable upto about 200°C and are therefore among the most stable organo metallic compounds known. Halide ions can coordinate to cobalt(III) as other common anionic ligands. Cobalt(III) complexes, being low spin, are conveniently studied in aqueous medium (Randall & Alberty, 1970). We report here the synthesis and X-ray crystal structure of the title compound.

The geometry around the cobalt(III) is approximately octahedral with the four glyoximate N atoms forming the square base;whereas, the coordinated bromide (Br1) and oxygen (O3) and the coordinated oxygen of water form the apex. The bite angles of the glyoximates with cobalt are N(1)#1-Co(1)—N(1) 82.18 (14)° and N(2)#1-Co(1)—N(2) 80.03 (03)°, respectively. Further N(1)#1-Co(1)—N(2)#1 178.29 (10)° confirms the distorted octahedral geometry of the molecule. The bond lengths Co(1)—N(1)#1, 1.883 (2) Å,Co(1)—N(2)#1, 1.911 (2) Å agree well with the previously reported structures (Meera *et al.*, 2009, Ramesh *et al.*, 2008) and the axial Co–Br distance d(Co1–Br1) = 2.3563 (6)Å agrees well with the reported structure of *trans*-aquabromobis[ethanedial dioximato(1-)-N,*N*']cobalt(III)(Mégnamisi-Bélombé *et al.*, 1983). The glyoximate moieties are further bound by strong intraomecular O—H···O hydrogen bonds showing an $S(6)$ ring motif (Bernstein *et al.*, 1995). Thecoordinated water forms an intermolecular hydrogen bond O3—H3···O1ii[symmetry code (ii): -x + 1, y - 1/2, -z + 1] with the glyoximate oxygen atoms which links the inversion related title compound thus forming a ring motif of $R_2^2(10)$. Fused rings of $R_2^2(10)$ generates a supramolecular one dimensional chain extending parallel to [010] direction. The structure is further stabilized through van der waals interaction.

S2. Experimental

Cobalt(II) bromide hexahydrate was thoroughly grinded and exposed to microwave for 30 s.The dehydrated cobalt(II) bromide was mixed with dimethylglyoxime in 1:2 molar ratio in acetone medium and allowed to stir for an hour (Vijayraghavan & Dayalan, 1992). The dibromo complex obtained was filtered dried and then it was refluxed with water for two hours. The resulting brown mass was filtered washed with ether and dried over desiccator. The elemental analysis data, obtained by analytical methods agree well with the theoretical data expected for the formula of the complex, $C_8H_{16}N_4O_5BrCo$ proposed *viz*.,[Co(dmgH)$_2$(H$_2$O)Br]: Anal,% (cald,%): C, 25.12(24.8); H,4.82(4.13); N,14.50(14.47). The C=N stretching vibration of oxime in its complex was observed at 1580 cm^{-1} and the intra molecular hydrogen bonded OH around 3100 cm^{-1}. A moderate peak around 1070 cm^{-1} may be assigned to the C=N—O stretching of the oxime. The peak around 510 cm^{-1} could be attributed to cobalt(III)-nitrogen stretching (Khan *et al.*, 1997; Folgando *et al.*, 1986). The ^{1}H NMR spectra of the complex in DMSO-d$_6$ shows a sharp intense singlet at 2.3 p.p.m. corresponding to

methyl protons of the oxime.The oxime –OH resonates at 13.08 p.p.m..A singlet around 8.5 ppm represents the –OH of the aquo ligand (Lopez *et al.*, 1986).

S3. Refinement

The H– atoms bound to C– atoms were constrained to riding atoms with d(C—H) = 0.96Å and U_{iso}(H) = 1.5U_{equ}(C).The positions of the hydrogen atoms,bound to the glyoximate and water O atoms, were identified from difference in the electron density map and restrained to a distance of d(O2—H2) = 0.92 (1)Å and d(O3—H3) = 0.85 (1) Å. A difference elctron density peak of 1.008 e A^{-3} was observed after the final refinement. Since the observed peak position is meaningless it is ignored.
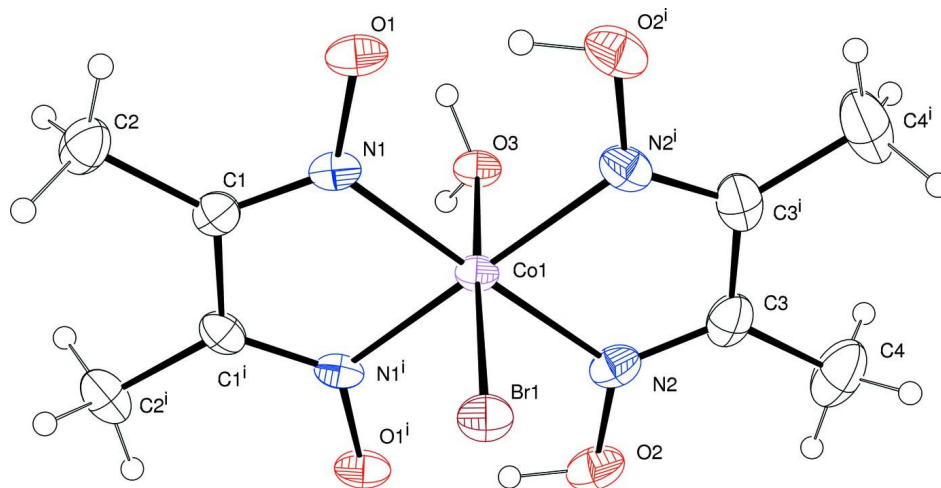
Figure 1

Displacement ellipsoid plot of the title compound drawn at 30% probability level.The equivalent symbol i represents the mirror symmetry (*x*, 1/2; -*y*,*z*) at one fourth of b axes.
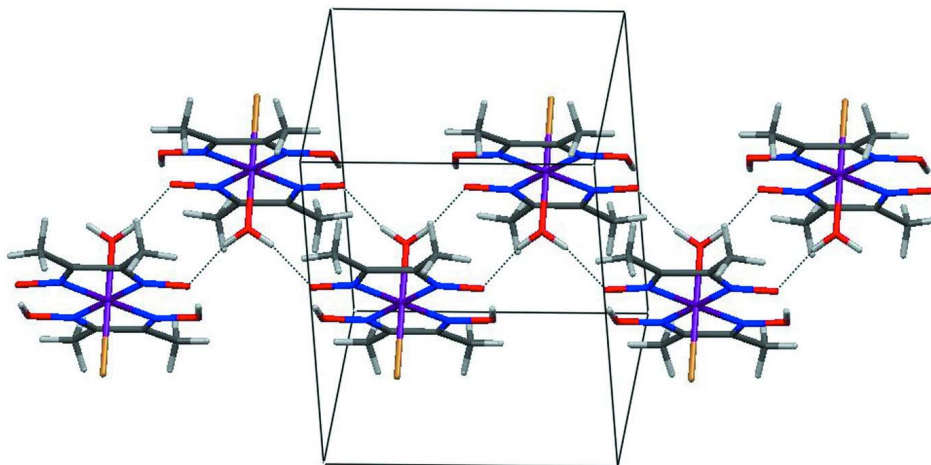
Figure 2

Part of the crystal structure of the title compound showing the formation of one dimensional chain through O3—H3⋯O1^{ii} hydrogen bond extending along [010] direction [Symmetry codes: (i) *x*, -*y* + 1/2, *z*; (ii) -*x* + 1, *y* - 1/2, -*z* + 1].

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Crystal data

$[CoBr(C_4H_7N_2O_2)_2(H_2O)]$	$F(000) = 388$
$M_r = 387.09$	$D_x = 1.821$ Mg m^{-3}
Monoclinic, $P2_1/m$	Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Hall symbol: -P 2yb	Cell parameters from 3745 reflections
$a = 7.5903$ (3) Å	$\theta = 2.7–30.7°$
$b = 8.8816$ (4) Å	$\mu = 4.07$ mm^{-1}
$c = 10.5343$ (5) Å	$T = 293$ K
$\beta = 96.137$ (3)°	Block, brown
$V = 706.09$ (5) Å^3	$0.15 \times 0.10 \times 0.10$ mm
$Z = 2$	

Data collection

Bruker Kappa APEXII CCD diffractometer	7395 measured reflections
Radiation source: fine-focus sealed tube	1480 independent reflections
Graphite monochromator	1298 reflections with $I > 2\sigma(I)$
ω and φ scans	$R_{int} = 0.028$
Absorption correction: multi-scan (*SADABS*; Bruker 1999)	$\theta_{max} = 26.0°$, $\theta_{min} = 2.7°$
$T_{min} = 0.581$, $T_{max} = 0.687$	$h = -9 \rightarrow 9$
	$k = -10 \rightarrow 9$
	$l = -12 \rightarrow 12$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.034$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.096$	$w = 1/[\sigma^2(F_o^2) + (0.0564P)^2 + 0.1583P]$ where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.22$	
1480 reflections	$(\Delta/\sigma)_{max} = 0.001$
101 parameters	$\Delta\rho_{max} = 1.01$ e Å^{-3}
2 restraints	$\Delta\rho_{min} = -0.54$ e Å^{-3}
Primary atom site location: structure-invariant direct methods	

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)

	x	y	z	U_{iso}*/U_{eq}
C1	0.7937 (4)	0.3331 (3)	0.5395 (3)	0.0330 (6)
C2	0.8935 (4)	0.4238 (4)	0.6415 (3)	0.0487 (8)
H2A	0.8818	0.5288	0.6207	0.073*

H2B	1.0164	0.3960	0.6486	0.073*
H2C	0.8469	0.4052	0.7213	0.073*
C3	0.3600 (4)	0.1669 (4)	0.1163 (3)	0.0448 (8)
C4	0.2476 (6)	0.0761 (5)	0.0221 (4)	0.0726 (13)
H4A	0.2737	−0.0288	0.0358	0.109*
H4B	0.2709	0.1038	−0.0626	0.109*
H4C	0.1250	0.0942	0.0318	0.109*
N1	0.6996 (3)	0.3893 (2)	0.4417 (2)	0.0302 (5)
N2	0.4668 (3)	0.1117 (3)	0.2069 (2)	0.0374 (6)
O1	0.6810 (3)	0.5377 (2)	0.4230 (2)	0.0395 (5)
O2	0.4819 (3)	−0.0402 (3)	0.2186 (2)	0.0501 (6)
O3	0.3771 (4)	0.2500	0.4145 (3)	0.0314 (6)
Co1	0.58731 (6)	0.2500	0.32505 (5)	0.02601 (18)
Br1	0.83644 (6)	0.2500	0.20942 (4)	0.04048 (18)
H2	0.557 (4)	−0.051 (4)	0.293 (2)	0.059 (12)*
H3	0.366 (5)	0.173 (3)	0.460 (3)	0.050 (10)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0234 (13)	0.0376 (16)	0.0391 (15)	−0.0028 (12)	0.0091 (11)	−0.0058 (12)
C2	0.0354 (17)	0.060 (2)	0.0503 (18)	−0.0080 (15)	0.0039 (14)	−0.0164 (17)
C3	0.0321 (15)	0.068 (2)	0.0356 (16)	−0.0038 (15)	0.0097 (13)	−0.0091 (15)
C4	0.054 (2)	0.106 (4)	0.056 (2)	−0.018 (2)	0.0022 (19)	−0.030 (2)
N1	0.0284 (12)	0.0230 (12)	0.0415 (13)	−0.0032 (9)	0.0139 (10)	−0.0029 (10)
N2	0.0327 (13)	0.0388 (15)	0.0430 (14)	−0.0053 (11)	0.0140 (11)	−0.0082 (11)
O1	0.0446 (12)	0.0229 (10)	0.0538 (13)	−0.0022 (9)	0.0180 (10)	−0.0031 (9)
O2	0.0540 (15)	0.0376 (13)	0.0604 (15)	−0.0075 (11)	0.0142 (12)	−0.0163 (11)
O3	0.0300 (14)	0.0259 (15)	0.0401 (16)	0.000	0.0130 (12)	0.000
Co1	0.0251 (3)	0.0227 (3)	0.0312 (3)	0.000	0.0076 (2)	0.000
Br1	0.0364 (3)	0.0432 (3)	0.0441 (3)	0.000	0.01498 (19)	0.000

Geometric parameters (Å, º)

C1—N1	1.289 (4)	C4—H4C	0.9600
C1—C1^i	1.476 (6)	N1—O1	1.338 (3)
C1—C2	1.485 (4)	N1—Co1	1.883 (2)
C2—H2A	0.9600	N2—O2	1.358 (3)
C2—H2B	0.9600	N2—Co1	1.911 (2)
C2—H2C	0.9600	O2—H2	0.921 (10)
C3—N2	1.283 (4)	O3—Co1	1.938 (3)
C3—C3^i	1.475 (7)	O3—H3	0.85 (3)
C3—C4	1.477 (4)	Co1—N1^i	1.883 (2)
C4—H4A	0.9600	Co1—N2^i	1.911 (2)
C4—H4B	0.9600	Co1—Br1	2.3563 (6)
N1—C1—C1^i	112.79 (16)	C3—N2—O2	119.2 (3)
N1—C1—C2	124.4 (3)	C3—N2—Co1	117.3 (2)

C1ⁱ—C1—C2	122.85 (19)	O2—N2—Co1	123.3 (2)
C1—C2—H2A	109.5	N2—O2—H2	103 (2)
C1—C2—H2B	109.5	Co1—O3—H3	114 (3)
H2A—C2—H2B	109.5	N1—Co1—N1ⁱ	82.18 (14)
C1—C2—H2C	109.5	N1—Co1—N2ⁱ	98.88 (11)
H2A—C2—H2C	109.5	N1ⁱ—Co1—N2ⁱ	178.29 (10)
H2B—C2—H2C	109.5	N1—Co1—N2	178.29 (10)
N2—C3—C3ⁱ	112.51 (19)	N1ⁱ—Co1—N2	98.88 (11)
N2—C3—C4	124.4 (4)	N2ⁱ—Co1—N2	80.03 (16)
C3ⁱ—C3—C4	123.1 (2)	N1—Co1—O3	91.24 (9)
C3—C4—H4A	109.5	N1ⁱ—Co1—O3	91.24 (9)
C3—C4—H4B	109.5	N2ⁱ—Co1—O3	87.40 (10)
H4A—C4—H4B	109.5	N2—Co1—O3	87.40 (10)
C3—C4—H4C	109.5	N1—Co1—Br1	90.29 (7)
H4A—C4—H4C	109.5	N1ⁱ—Co1—Br1	90.29 (7)
H4B—C4—H4C	109.5	N2ⁱ—Co1—Br1	91.04 (7)
C1—N1—O1	122.7 (2)	N2—Co1—Br1	91.04 (7)
C1—N1—Co1	116.1 (2)	O3—Co1—Br1	177.97 (9)
O1—N1—Co1	121.18 (18)		
C1ⁱ—C1—N1—O1	179.81 (18)	C1—N1—Co1—O3	91.1 (2)
C2—C1—N1—O1	−0.7 (4)	O1—N1—Co1—O3	−88.7 (2)
C1ⁱ—C1—N1—Co1	0.02 (19)	C1—N1—Co1—Br1	−90.28 (19)
C2—C1—N1—Co1	179.5 (2)	O1—N1—Co1—Br1	89.92 (19)
C3ⁱ—C3—N2—O2	−179.62 (19)	C3—N2—Co1—N1ⁱ	174.1 (2)
C4—C3—N2—O2	0.8 (5)	O2—N2—Co1—N1ⁱ	−2.4 (2)
C3ⁱ—C3—N2—Co1	3.8 (2)	C3—N2—Co1—N2ⁱ	−4.6 (3)
C4—C3—N2—Co1	−175.8 (3)	O2—N2—Co1—N2ⁱ	178.98 (17)
C1—N1—Co1—N1ⁱ	0.0 (2)	C3—N2—Co1—O3	83.2 (2)
O1—N1—Co1—N1ⁱ	−179.82 (14)	O2—N2—Co1—O3	−93.2 (2)
C1—N1—Co1—N2ⁱ	178.6 (2)	C3—N2—Co1—Br1	−95.5 (2)
O1—N1—Co1—N2ⁱ	−1.2 (2)	O2—N2—Co1—Br1	88.1 (2)

Symmetry code: (i) *x*, −*y*+1/2, *z*.

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
O2—H2···O1ⁱ	0.92 (1)	1.58 (1)	2.494 (3)	169 (4)
O3—H3···O1ⁱⁱ	0.85 (3)	1.79 (3)	2.616 (3)	167 (4)

Symmetry codes: (i) *x*, −*y*+1/2, *z*; (ii) −*x*+1, *y*−1/2, −*z*+1.