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Key indicators

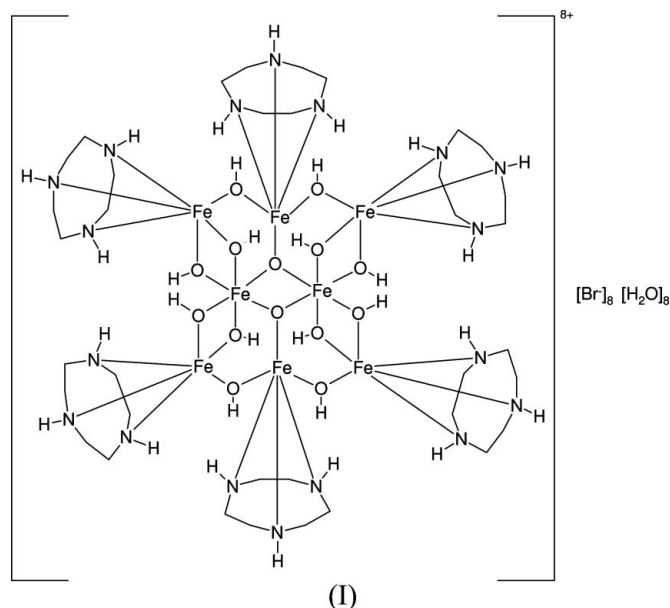
Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.012$ Å
H-atom completeness 87%
Disorder in solvent or counterion
 R factor = 0.059
 wR factor = 0.209
Data-to-parameter ratio = 28.0For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Dodeca- μ_2 -hydroxo-di- μ_3 -oxo-hexakis(1,4,7-triaza-
cyclononane- $\kappa^3\text{N},\text{N}',\text{N}''$)octairon(III) octabromide
octahydrate

The title compound, $[\text{Fe}_8(\mu_3\text{-O})_2(\mu_2\text{-OH})_{12}(\text{C}_6\text{H}_{12}\text{N}_3)_6]\text{Br}_8 \cdot 8\text{H}_2\text{O}$, which crystallizes as an octahydrate, has a different arrangement of cations, anions and water molecules from the known nonahydrate [Wieghardt, Pohl, Jibril & Huttner (1984). *Angew. Chem. Int. Ed. Engl.* **23**, 77–78.]. In the present phase, the cluster is generated by inversion symmetry and the bromide ions display substantial positional disorder.

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Comment

As in the previously reported nonahydrate structure (Wieghardt *et al.*, 1984), there is one $[(\text{C}_6\text{N}_3\text{H}_{12})_6\text{Fe}_8(\mu_3\text{-O})_2(\mu_2\text{-OH})_{12}]^{8+}$ cluster per unit cell of the title compound, (I) (Fig. 1). The two structures present a similar conformation of the cluster but a completely different distribution of bromide ions and water molecules. In (I), the cluster is generated by inversion symmetry (Table 1), whereas in the nonahydrate, the cluster possesses no symmetry.



According to charge balance, four Br^- ions per asymmetric unit should be present to compensate the positive charge of the cluster. There is substantial disorder in the positions of the anions, and the four Br^- ions appear distributed over nine sites with occupancy factors ranging from 0.181 (2) to 0.838 (2). The bromide ions in six of these sites act as acceptors for one or two hydrogen bonds with the OH groups, forming μ_2 -hydroxo bridges, and with the N atoms in the cyclic amine ligands. The sum of the occupancy factors of the six bromide sites hydrogen bonded to the cluster is 2.97, that is, each cluster is linked by hydrogen bonds to 5.94 Br atoms on

average. The bromide ions in the remaining three sites, together with the solvent water molecules, fill the space between the clusters, forming a complex network of hydrogen bonds which crosslinks the clusters and gives cohesion to the crystal structure.

This distribution of hydrogen bonds is very different from that observed in the previously reported nonahydrate structure. In that case, there were seven bromide ions and one water molecule acting as acceptors for hydrogen bonds between solvent water molecules and the cluster. Owing to the disorder in the bromide positions in (I), the unit-cell volume is larger in this case [$2163.0(8) \text{ \AA}^3$ versus 1959 \AA^3], in spite of containing one fewer water molecule.

Experimental

Crystals of (I) were obtained while following the literature procedure to grow crystals of the known nonahydrate (Wiegardt *et al.*, 1984).

Crystal data

$[\text{Fe}_8\text{O}_2(\text{OH})_{12}(\text{C}_6\text{H}_{12}\text{N}_3)_6]\text{Br}_8 \cdot 8\text{H}_2\text{O}$
 $M_r = 2241.56$
 Triclinic, $P\bar{1}$
 $a = 13.257(2) \text{ \AA}$
 $b = 13.564(1) \text{ \AA}$
 $c = 14.831(4) \text{ \AA}$
 $\alpha = 108.15(1)^\circ$
 $\beta = 113.34(1)^\circ$
 $\gamma = 101.43(1)^\circ$
 $V = 2163.0(8) \text{ \AA}^3$

$Z = 1$
 $D_x = 1.721 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation
 Cell parameters from 15 reflections
 $\theta = 10.8\text{--}12.6^\circ$
 $\mu = 5.06 \text{ mm}^{-1}$
 $T = 293(2) \text{ K}$
 Prism, brown
 $0.5 \times 0.37 \times 0.28 \text{ mm}$

Data collection

Enraf–Nonius CAD-4 diffractometer
 Non-profiled $\omega/2\theta$ scans
 Absorption correction: ψ scan (North *et al.*, 1968)
 $T_{\min} = 0.142$, $T_{\max} = 0.241$
 13602 measured reflections
 13085 independent reflections
 8217 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.029$
 $\theta_{\text{max}} = 30.4^\circ$
 $h = 0 \rightarrow 18$
 $k = -19 \rightarrow 18$
 $l = -21 \rightarrow 19$
 3 standard reflections every 90 reflections
 intensity decay: none

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.059$
 $wR(F^2) = 0.209$
 $S = 1.12$
 13085 reflections
 467 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.116P)^2 + 2.3967P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.048$
 $\Delta\rho_{\text{max}} = 2.53 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -1.03 \text{ e \AA}^{-3}$

Table 1

Selected bond lengths (\AA).

Fe1—O1 ⁱ	1.959 (3)	Fe4—O1	1.872 (3)
Fe1—O1	1.967 (3)	Fe4—O6	1.989 (3)
Fe1—O2	2.030 (3)	Fe4—O7	2.011 (3)
Fe1—O4	2.031 (3)	Fe4—N41	2.156 (4)
Fe1—O5	2.040 (3)	Fe4—N44	2.167 (4)
Fe1—O3	2.084 (3)	Fe4—N47	2.241 (4)
Fe3—O5	1.922 (3)	Fe2—O3	1.939 (3)
Fe3—O4	1.950 (3)	Fe2—O2	1.957 (3)
Fe3—O6	1.964 (3)	Fe2—N24	2.160 (5)
Fe3—N37	2.159 (4)	Fe2—N21	2.170 (4)
Fe3—N34	2.161 (4)	Fe2—N27	2.179 (5)
Fe3—N31	2.181 (4)		

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

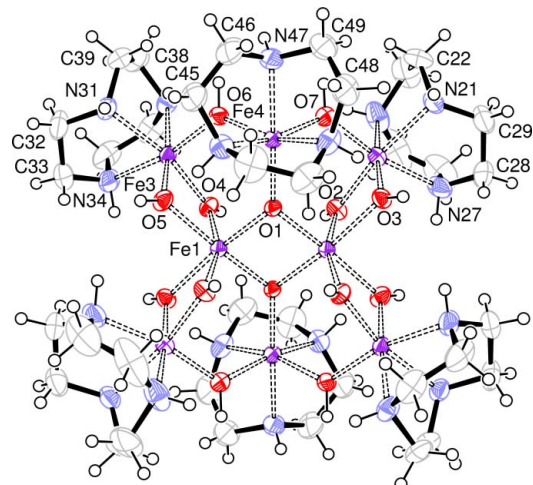


Figure 1

View of the cluster in (I), showing 50% probability displacement ellipsoids for the non-H atoms. Unlabelled atoms are generated from the labelled atoms by the symmetry operation $(1 - x, 1 - y, 1 - z)$.

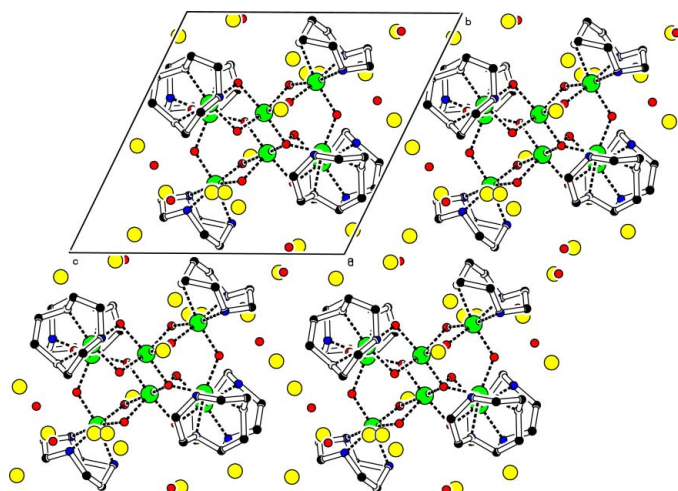


Figure 2

Molecular packing in (I) viewed along the a axis. H atoms have been omitted for clarity. Colour key: Fe green, C black, N blue, O red, Br yellow. Fe—N and Fe—O bonds are shown as dashed lines.

Table 2

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

$D\text{—}H\cdots A$	$D\text{—}H$	$H\cdots A$	$D\cdots A$	$D\text{—}H\cdots A$
O5—H5 \cdots Br1	0.93	2.24	3.142 (3)	165
O6—H6 \cdots Br2	0.93	2.52	3.271 (3)	139
O7—H7 \cdots Br2	0.93	2.52	3.306 (3)	143
O3—H3 \cdots Br3 ⁱ	0.93	2.32	3.242 (3)	171
O2—H2 \cdots Br6	0.93	2.31	3.116 (10)	145
O4—H4 \cdots Br6 ⁱ	0.93	2.29	3.123 (10)	149
O2—H2 \cdots Br8	0.93	2.55	3.274 (5)	135
O4—H4 \cdots Br8 ⁱ	0.93	2.55	3.298 (5)	138
N24—H24 \cdots Br7	0.91	2.64	3.519 (6)	164
N24—H24 \cdots Br8	0.91	2.96	3.591 (8)	128
N27—H27 \cdots Br1	0.91	2.72	3.562 (5)	155
N31—H31 \cdots Br1	0.91	2.94	3.643 (4)	136
N34—H34 \cdots Br3 ⁱ	0.91	2.54	3.403 (5)	159
N37—H37 \cdots Br7 ⁱ	0.91	2.79	3.578 (5)	146
N37—H37 \cdots Br2	0.91	3.00	3.731 (5)	139
N41—H41 \cdots Br3	0.91	3.08	3.853 (5)	144
N47—H47 \cdots Br2	0.91	2.89	3.689 (5)	148

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

H atoms in the cluster were placed in calculated positions (C–H = 0.97 Å, N–H = 0.91 Å and O–H = 0.93 Å) and refined as riding with the constraint $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{carrier})$ applied. The H atoms of the solvent water molecules could not be located in difference Fourier maps. The highest peak and deepest hole at the end of the refinement were located close to atoms Br3 and O2w, respectively.

Data collection: *CAD-4 EXPRESS* (Enraf–Nonius, 1994); cell refinement: *CAD-4 EXPRESS*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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