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Hemi(piperazinediium) hexaaquaaluminium(III) bis(sulfate) tetrahydrate: a new double aluminium sulfate salt

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Piperazinium aluminium sulfate decahydrate, $(C_4H_{12}N_2)_{0.5}$ $[AI(H_2O)_6](SO_4)_2.4H_2O$, exhibits a crystal structure built from isolated $[A](H_2O)_6^{3+}$, SO_4^{2-} , $C_4H_{12}N_2^{2+}$ and H_2O units connected by a complex hydrogen-bond network. The title compound shows strong similarities to many double aluminium sulfates, such as alums and Tutton's salts. However, since its structure is not derived directly from that of these compounds, it is assumed to be a new structure type.

Comment

Studies in sulfate-based chemistry have recently experienced a resurgence in interest, with the use of the SO_4^2 anion as a tetrahedral building block and amines as templates (Choudhury et al., 2001; Norquist et al., 2003). The new amine lanthanum sulfates obtained recently (Bataille $& Louër, 2002$; Xing et al., 2003) also offer a promising route to open structures. Similarly, by using trivalent aluminium, it would be expected that materials with open frameworks constructed only from tetrahedral building units could be prepared. Instead, the crystal structure of the new piperazinium aluminium sulfate decahydrate obtained here, $viz.$ $(C_4H_{12}N_2)_{0.5}$ $[AI(H_2O)_6](SO_4)_2.4H_2O$, (I), consists of isolated entities linked through a complex hydrogen-bond network. The present work also deals with the similarity of the crystal structure of (I) to those of other double aluminium sulfates, such as Tutton's salts and alums.

Alums [general formula $A^{I}M^{III}(SO_4)_2.12H_2O$] and Tutton's salts [general formula $A_2^{\text{I}}M^{\text{II}}(\text{SO}_4)_2.6\text{H}_2\text{O}$] are two large families of double salts that include the sulfate anion. The

Figure 1

A projection of the structure of (I) along the *a* axis, showing the hydrogen-bond network formed by the $[A1(H_2O)_6]^3$ (light grey) octahedra), SO_4^2 ⁻ (dark grey tetrahedra), $C_4H_{12}N_2^2$ ⁺ and H_2O entities. Hydrogen bonds formed via water molecules are represented by thick dashed lines, while hydrogen bonds involving N atoms are shown as thin dashed lines. H atoms have been omitted for clarity.

chemical formula of (I) can be written as $[(C_4H_{12}N_2)_0,$ $(SO₄)₂$ 10H₂O. Compound (I) is thus related to the alum family, except that A is a divalent cation with halved stoichiometry and the compound is decahydrated. Nevertheless, the space-group symmetry of (I) $(P2₁/n)$ is comparable to that of dimethylammonium aluminium sulfate hexahydrate $(P2₁/n;$ Galešić & Jordanovska, 1992) and Tutton's salts ($P2₁/a$; see, for example, Maslen et al., 1988), while alums crystallize with a cubic symmetry. The structure of (I) (Figs. 1 and 2) consists of Al^{3+} ions octahedrally coordinated to six water molecules, together with sulfate tetrahedra, piperazinium cations (lying about inversion centres) and non-bonded water molecules, linked by hydrogen bonds only. This bonding feature has already been observed in, for example, piperazinium sulfate monohydrate (Jayaraman et al., 2002). The crystal structure of

Figure 2

Part of the crystal structure of (I), showing the asymmetric unit and the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. Hydrogen bonds are represented by dashed lines. [Symmetry code: (i) $-x$, $-y$, $-z$.]

(I) is unlike that of other double aluminium sulfates, in which the metal atom is located at the origin of the cell, in that the Al atom in (I) lies in a general position. Selected bond lengths are given in Table 1. The $AI-OW$ bond lengths range from 1.8620 (17) to 1.8872 (17) Å [mean 1.876 (2) Å], in agreement with the value (1.876 Å) calculated using the bond-valence program VALENCE (Brown, 1996) for a sixfold oxygencoordinated Al atom. The $S-O$ distances fall in the range 1.4611 (17)-1.4809 (17) \AA [mean 1.473 (2) \AA], which is also comparable to the value (1.474 Å) calculated using the bondvalence method. This bond length agrees with that observed in many Tutton's salts, while it is generally slightly shorter in alums (Petruševski, 1994). No disorder of the amine group is observed, and the centre of the piperazinium group corresponds to the inversion centre of the unit cell. In the dimethylammonium salt (Galešić & Jordanovska, 1992), the amine group is disordered around the inversion centre, and in α -alums, the monovalent A^T cations are disordered around the special position 4b of space group Pa3 (Larson & Cromer, 1967), while in Tutton's salts, the A^T cations lie in general positions (Maslen et al., 1988). The presence of a diprotonated amine group and ten water molecules leads to a complex hydrogen-bond network, involving 22 H atoms (Table 2).

Clearly, the crystal structure of (I) has some analogies to those of other double aluminium sulfate salts. Nevertheless, these similarities are not sufficient to allow us to consider (I) as belonging to the known families of double salts, such as alums and Tutton's salts. Rather, it is likely that (I) is the first member of a new family of double aluminium sulfates. It is therefore of interest to investigate further new aluminium sulfates prepared in the presence of diprotonated amines, in order to establish whether similarities exist with (I).

Experimental

Single crystals of (I) were obtained by dissolving $Al_2(SO_4)_3.18H_2O$ (0.75 mmol) and piperazine hexahydrate (2 mmol) in $H₂O$ (7 ml) acidified with H_2SO_4 (0.5 ml). The resulting clear solution was allowed to stand for two months, until large crystals were obtained. The product was filtered off and washed with a small amount of distilled water, since the crystals appeared to be slightly soluble.

Crystal data

 $T_{\text{min}} = 0.797, T_{\text{max}} = 0.967$ 19 066 measured reflections 3911 independent reflections

Table 1

Selected interatomic distances (A) .

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

Hydrogen-bonding geometry (A, \circ) .

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

Refinement

All 26 H atoms were located from difference Fourier syntheses and their positions were refined isotropically, with $O-H$, $C-H$ and $N-\frac{1}{2}$ H distances restrained to 0.97 (2) \AA and H \cdots H distances restrained to 1.55 (3) Å. The atomic displacement parameters of the H atoms were fixed at $1.5U_{eq}$ (parent atom).

Data collection: COLLECT (Nonius, 1998); cell refinement: HKL SCALEPACK (Otwinowski & Minor, 1997); data reduction: HKL DENZO (Otwinowski & Minor, 1997) and SCALEPACK; program(s) used to solve structure: SIR97 (Altomare et al., 1999); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: DIAMOND (Brandenburg & Berndt, 1999); software used to prepare material for publication: $WinGX$ (Farrugia, 1999).

 $k = -15 \to 15$ $l = -27 \rightarrow 27$

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: DN1030). Services for accessing these data are described at the back of the journal.

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