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manuela@pollux.fis.uc.pt**Key indicators**Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
Disorder in main residue
 R factor = 0.028
 wR factor = 0.076
Data-to-parameter ratio = 14.2For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.**Tris(L-argininium) hexafluorochromate(III) trihydrate**

The title compound, $(\text{C}_6\text{H}_{15}\text{N}_4\text{O}_2)_3 \cdot \text{CrF}_6 \cdot 3\text{H}_2\text{O}$, is isostructural with the vanadium analogue. One of the arginium cations is partially disordered over two different conformations. A complex network of $\text{N}-\text{H} \cdots \text{O}$, $\text{N}-\text{H} \cdots \text{F}$, $\text{O}-\text{H} \cdots \text{O}$ and $\text{O}-\text{H} \cdots \text{F}$ hydrogen bonds supports the structure.

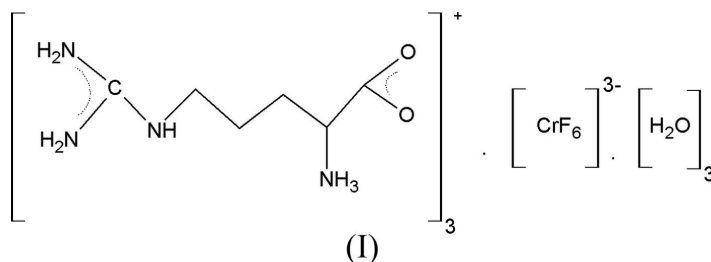
Received 12 October 2005

Accepted 27 October 2005

Online 31 October 2005

Comment

The title compound, (I) (Fig. 1), is isostructural with tris-(L-argininium) hexafluorovanadate(III) trihydrate, (II) (Ramos Silva, Matos Beja, Paixão & Alte da Veiga, 2000). The bond lengths and angles for (I) (Table 1) and the molecular conformations are all very similar to those of the equivalent species in (II). The arginium species in (I) are positively charged, with the guanidyl and amino groups protonated and the carboxyl groups deprotonated. One of the arginium cations is partially disordered, adopting two alternative conformations (Fig. 1). In the two ordered arginium cations (containing atoms C1 and C7), $\text{C}\gamma$ (*i.e.* atoms C3 and C9, respectively) is in a staggered *trans* position with respect to the carboxyl C atom, while in the disordered C13-containing molecule, $\text{C}\gamma$ (atom C15) is in a *gauche* position with respect to both the amino and carboxyl C atoms (Table 1).



There is an extended and complex three-dimensional network of hydrogen bonds in the crystal structure of (I), three of which are bifurcated, involving 15 donors and 31 H atoms (Table 2). Every F atom accepts at least two hydrogen bonds (Fig. 2), as also seen in a related compound (Ramos Silva, Matos Beja, Costa *et al.*, 2000). Isomorphous compounds of hexafluorovanadate(III) and hexafluorochromate(III) have already been found, for instance trihydrazinium hexafluorochromate(III) and trihydrazinium hexafluorovanadate(III) (Kojić-Prodić *et al.*, 1972).

Magnetic susceptibility measurements on (I) were performed down to 5 K in an applied field of 50 Oe using a powdered sample. The variation of the induced magnetic moment with temperature shows paramagnetic behaviour down to the lowest temperature with no sign of magnetic ordering.

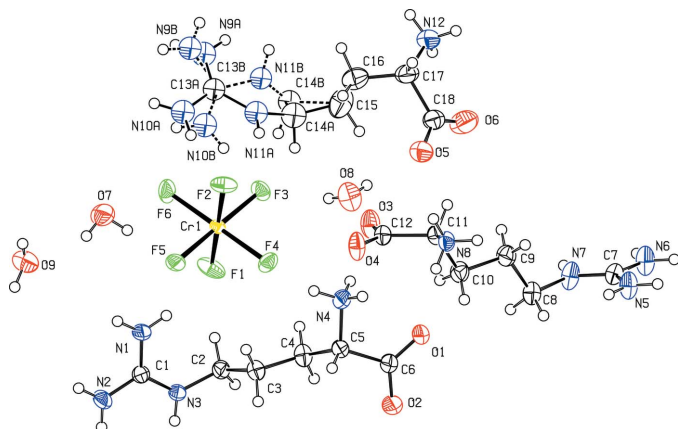


Figure 1
A view of (I), showing 50% probability displacement ellipsoids. Both disorder components are shown.

Experimental

Hydrofluoric acid (Merck, 40%) was reacted with powdered chromium (99.7%) in a plastic flask. The resulting solution was then added to an aqueous solution of L-arginine (Aldrich, 98%) in a 1:3 Cr-arginine molar proportion. Several recrystallizations were carried out using water and ethanol as solvent over a period of several weeks, eventually yielding green prisms of (I).

Crystal data

$(C_6H_{15}N_4O_2)_3[CrF_6] \cdot 3H_2O$
 $M_r = 745.71$
 Monoclinic, $P2_1$
 $a = 10.8194$ (15) Å
 $b = 9.0434$ (15) Å
 $c = 16.998$ (2) Å
 $\beta = 96.890$ (12)°
 $V = 1651.2$ (4) Å³
 $Z = 2$
 $D_x = 1.500$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 25 reflections
 $\theta = 9.8$ – 15.7 °
 $\mu = 0.44$ mm⁻¹
 $T = 293$ (2) K
 Prism, green
 $0.34 \times 0.22 \times 0.15$ mm

Data collection

Enraf–Nonius CAD-4 diffractometer
 Profile-fitted $\omega/2\theta$ scans
 Absorption correction: ψ scan North *et al.* (1968)
 $T_{min} = 0.884$, $T_{max} = 0.984$
 12219 measured reflections
 6118 independent reflections
 5397 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.028$
 $\theta_{max} = 25.5$ °
 $h = -13 \rightarrow 13$
 $k = -10 \rightarrow 10$
 $l = -20 \rightarrow 20$
 3 standard reflections
 frequency: 180 min
 intensity decay: 7%

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.028$
 $wR(F^2) = 0.076$
 $S = 1.05$
 6118 reflections
 432 parameters
 H atoms treated by a mixture of independent and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0451P)^2 + 0.3634P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} < 0.001$
 $\Delta\rho_{max} = 0.28$ e Å⁻³
 $\Delta\rho_{min} = -0.29$ e Å⁻³
 Absolute structure: Flack (1983), with 2845 Friedel pairs
 Flack parameter: 0.010 (14)

Table 1

Selected geometric parameters (Å, °).

Cr1—F2	1.8855 (14)	Cr1—F3	1.9010 (13)
Cr1—F1	1.8857 (14)	Cr1—F5	1.9020 (13)
Cr1—F6	1.8902 (13)	Cr1—F4	1.9096 (13)
C15—C16—C17—N12	72.1 (3)	C3—C4—C5—C6	144.5 (2)
C15—C16—C17—C18	−52.0 (4)	C9—C10—C11—C12	142.9 (2)

Table 2

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N12—H12A ⁱ ···O5 ⁱ	0.89	2.20	3.034 (3)	157
N12—H12B ⁱⁱ ···O3 ⁱⁱ	0.89	2.23	2.948 (3)	137
N12—H12C ⁱⁱⁱ ···O7 ⁱⁱⁱ	0.89	1.92	2.756 (3)	156
N10A—H10B ^{iv} ···F6	0.86	1.89	2.620 (5)	141
N10A—H10B ^v ···F3	0.86	2.60	3.302 (5)	140
N10B—H10D ^{vi} ···F6	0.86	2.32	2.748 (6)	111
N10B—H10C ^{vii} ···F3	0.86	2.03	2.803 (6)	150
N10A—H10A ^{viii} ···O6 ^{viii}	0.86	1.89	2.740 (5)	169
N11B—H11B ^{ix} ···F2 ^{ix}	0.86	1.91	2.744 (5)	163
N11A—H11A ^x ···F3	0.86	1.98	2.823 (4)	165
N9A—H9E ^{xi} ···O8 ^{xi}	0.86	2.43	3.102 (5)	136
N9B—H9H ^{xii} ···O8 ^{xii}	0.86	2.41	2.855 (6)	113
N9B—H9G ^{xiii} ···O6 ^{xiii}	0.86	2.29	3.117 (6)	160
N9A—H9F ^{xiv} ···F2 ^{xiv}	0.86	2.13	2.793 (5)	133
N9B—H9H ^{xv} ···F2 ^{xv}	0.86	2.33	3.006 (6)	136
N9A—H9F ^{xvi} ···F3 ^{xvi}	0.86	2.48	3.072 (5)	127
N9B—H9H ^{xvii} ···F3 ^{xvii}	0.86	2.64	3.384 (6)	146
N1—H1A ^{xviii} ···O1 ^{xviii}	0.86	2.23	3.080 (2)	171
N1—H1B ^{xix} ···F1	0.86	1.89	2.732 (2)	167
N2—H21 ^{xx} ···O2 ^{xx}	0.86	1.96	2.818 (3)	176
N2—H22 ^{xxi} ···F4 ^{xxi}	0.86	2.00	2.820 (2)	158
N3—H3 ^{xxii} ···F5 ^{xxii}	0.86	1.88	2.726 (2)	168
N4—H4A ^{xxiii} ···O4	0.89	2.03	2.901 (3)	168
N4—H4B ^{xxiv} ···O9 ^{xxiv}	0.89	2.09	2.946 (3)	161
N4—H4C ^{xxv} ···F4	0.89	1.84	2.696 (2)	161
N5—H5B ^{xxvi} ···O4 ^{xxvi}	0.86	2.07	2.929 (3)	173
N5—H5A ^{xxvii} ···F1 ^{xxvii}	0.86	1.92	2.747 (2)	162
N6—H6A ^{xxviii} ···F4 ^{xxviii}	0.86	2.42	3.028 (3)	128
N6—H6A ^{xxix} ···F2 ^{xxix}	0.86	2.63	3.256 (3)	131
N6—H6B ^{xxx} ···O3 ^{xxx}	0.86	1.99	2.846 (3)	171
N7—H7 ^{xxxi} ···F2 ^{xxxi}	0.86	1.89	2.723 (2)	162
N8—H8A ^{xxxii} ···O5	0.89	2.03	2.860 (3)	154
N8—H8B ^{xxxiii} ···F6 ^{xxxiii}	0.89	1.75	2.640 (2)	176
N8—H8C ^{xxxiv} ···O1	0.89	2.00	2.884 (2)	171
O7—H71 ^{xxxv} ···O9	0.79 (4)	2.17 (4)	2.875 (3)	148 (4)
O7—H72 ^{xxxvi} ···F5	0.88 (4)	1.81 (4)	2.687 (2)	175 (4)
O8—H81 ^{xxxvii} ···O5	0.78 (4)	2.05 (4)	2.804 (3)	165 (5)
O8—H82 ^{xxxviii} ···F3	0.79 (4)	2.06 (4)	2.787 (3)	154 (4)
O9—H91 ^{xxxix} ···O2 ^{xxxix}	0.89 (4)	1.85 (4)	2.718 (3)	166 (4)
O9—H92 ^{xl} ···O3 ^{xl}	0.80 (4)	2.23 (4)	2.996 (3)	160 (4)

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

The amino acid H atoms were placed in calculated positions and refined as riding on their parent atoms, with C—H = 0.97 Å and N—H = 0.86–0.89 Å, and with $U_{iso}(H) = 1.2$ or $1.5U_{eq}(C,N)$. Water H atoms were located in a Fourier difference map and their positions were freely refined, with $U_{iso}(H) = 1.5U_{eq}(O)$. Atoms N11, C14, N9 and N10 and their attached H atoms were found to be disordered over two positions in the ratio 0.547 (4):0.453 (4) and were refined with isotropic displacement factors. Examination of the crystal structure with PLATON (Spek, 2003) showed that there are no solvent-accessible voids in the crystal structure.

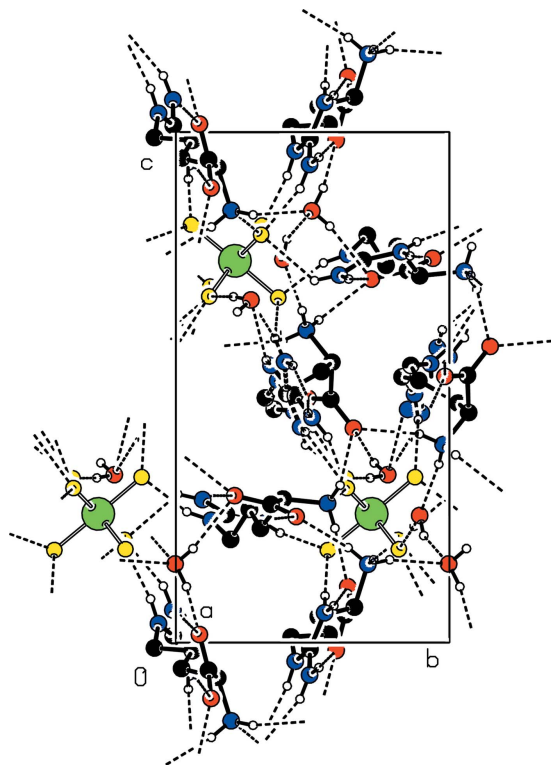


Figure 2

The packing of (I), projected along the *a* axis. Hydrogen bonds are shown as dashed lines. H atoms not involved in hydrogen bonding have been omitted for clarity.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *HELENA* (Spek, 1997); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

This work was supported by Fundação para a Ciência e a Tecnologia (FCT) under project POCI/FIS/57876/2004. The authors are indebted to Professor Margarida Godinho (FCUL) for using the SQUID magnetometer, and to H. Natal da Luz (MSc) for using X-ray fluorescence to confirm the presence of vanadium and chromium in the isostructural samples.

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