

The Rb analogue of grimselite, $\text{Rb}_6\text{Na}_2[(\text{UO}_2)(\text{CO}_3)_3]_2(\text{H}_2\text{O})$

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The crystal structure of the Rb analogue of grimselite, rubidium sodium uranyl tricarbonate hydrate, $\text{Rb}_6\text{Na}_2[(\text{UO}_2)(\text{CO}_3)_3]_2(\text{H}_2\text{O})$, consists of a uranyl hexagonal bipyramid that shares three non-adjacent equatorial edges with carbonate triangles, resulting in a uranyl tricarbonate cluster of composition $[(\text{UO}_2)(\text{CO}_3)_3]$. These uranyl tricarbonate clusters form layers perpendicular to $[001]$ and are interconnected by NaO_8 polyhedra. The title compound is isostructural with grimselite, with a reduced occupancy of the H_2O site (25% versus 50% in grimselite).

Comment

The compound $\text{Rb}_6\text{Na}_2[(\text{UO}_2)(\text{CO}_3)_3]_2(\text{H}_2\text{O})$ was obtained by evaporation of a solution containing $\text{UO}_2(\text{NO}_3)_2(\text{H}_2\text{O})_6$, RbNO_3 and Na_2CO_3 at ambient temperature and pressure. $\text{Rb}_6\text{Na}_2[(\text{UO}_2)(\text{CO}_3)_3]_2(\text{H}_2\text{O})$ is structurally analogous to the mineral grimselite, $\text{K}_3\text{Na}[(\text{UO}_2)(\text{CO}_3)_3](\text{H}_2\text{O})$ (Li & Burns, 2001). Each contains the uranyl tricarbonate cluster, $[(\text{UO}_2)(\text{CO}_3)_3]$, and crystallizes in space group $P\bar{6}2c$. However, these two compounds differ in their hydration state.

There is one symmetrically distinct U^{VI} atom in the structure of $\text{Rb}_6\text{Na}_2[(\text{UO}_2)(\text{CO}_3)_3]_2(\text{H}_2\text{O})$. This atom is part of a linear uranyl ion, $(\text{UO}_2)^{2+}$, which is coordinated by six equatorial O atoms, forming a uranyl hexagonal bipyramid. Three non-adjacent equatorial edges of the uranyl hexagonal bipyramid share edges with CO_3 groups, giving the uranyl tricarbonate cluster, $[(\text{UO}_2)(\text{CO}_3)_3]$ (Fig. 1). These clusters share edges with NaO_8 hexagonal bipyramids, to form heteropolyhedral sheets parallel to (001) (Fig. 2). These sheets are stacked along $[001]$, with voids in the resulting framework containing Rb cations and H_2O molecules. The H_2O sites are 25% occupied in the structure of $\text{Rb}_6\text{Na}_2[(\text{UO}_2)(\text{CO}_3)_3]_2(\text{H}_2\text{O})$, whereas they are 50% occupied in the structure of grimselite. This difference in H_2O content may be attributable to the larger size of the Rb^+ cation.

Bond-valence sums were calculated for each ion using the parameters of Brese & O'Keeffe (1991), giving the following values: 6.18 for U, 4.15 for C, 1.29 for Rb and 1.15 for Na

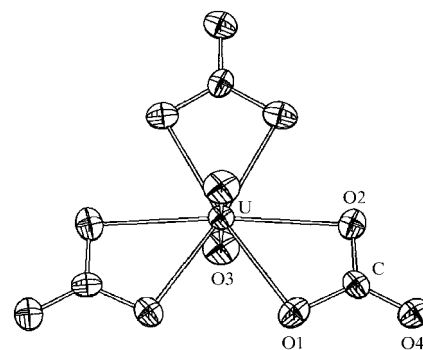


Figure 1

A view of the $[(\text{UO}_2)(\text{CO}_3)_3]$ clusters in $\text{Rb}_6\text{Na}_2[(\text{UO}_2)(\text{CO}_3)_3]_2(\text{H}_2\text{O})$, shown with 50% probability displacement ellipsoids.

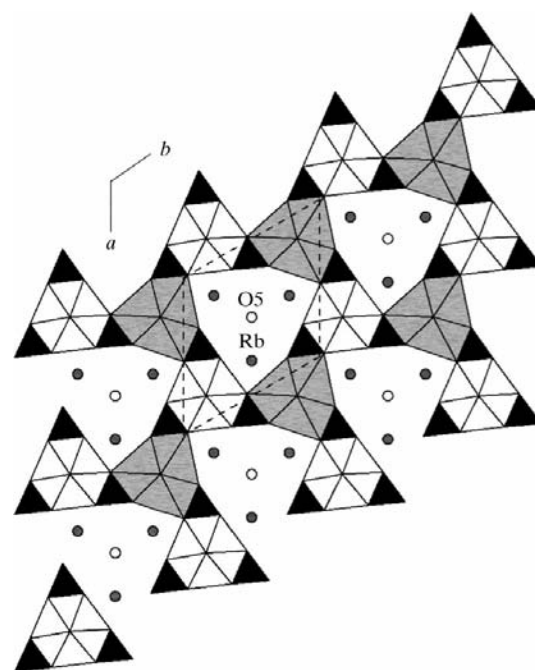


Figure 2

The heteropolyhedral layer of NaO_8 hexagonal bipyramids (grey) and $[(\text{UO}_2)(\text{CO}_3)_3]$ clusters (UO_8 shown in white, CO_3 shown in black), projected along $[001]$ (unit cell shown as dashed lines).

(Burns *et al.*, 1997). The bond-valence sum for Rb is based on its coordination environment when the O5 site is occupied, and the partial occupancy of this site may, therefore, represent a way to limit the overbonding of the Rb site. For atoms O1, O2, O3 and O4, the bond-valence sums are within the normal range (1.88–2.25). The bond-valence sum for atom O5 (0.32) is consistent with a water molecule.

Experimental

Transparent yellow crystals of $\text{Rb}_6\text{Na}_2[(\text{UO}_2)(\text{CO}_3)_3]_2(\text{H}_2\text{O})$, up to 1 mm in maximum dimension, were synthesized from a solution containing Na_2CO_3 (0.026 g), RbNO_3 (0.111 g) and $\text{UO}_2(\text{NO}_3)_2(\text{H}_2\text{O})_6$ (0.123 g) in ultra-pure water (25 ml). Following evaporation

in an open beaker for 28 d at 293 K under ambient pressure, crystals were recovered from the beaker walls.

Crystal data

Rb₆Na₂[(UO₂)(CO₃)₃]₂(H₂O)
M_r = 1476.94
 Hexagonal, *P*6̄2*c*
a = 9.4316 (7) Å
c = 8.3595 (8) Å
V = 643.99 (9) Å³
Z = 1
D_x = 3.808 Mg m⁻³

Mo *K*α radiation
 Cell parameters from 1623 reflections
 $\theta = 3.3\text{--}33.8^\circ$
 $\mu = 23.95\text{ mm}^{-1}$
T = 297 (2) K
 Anhydrous, yellow
 0.15 × 0.12 × 0.12 mm

Data collection

Bruker APEX CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: empirical via ψ scan (*SHELXTL-NT*; Bruker, 1998)
T_{min} = 0.044, *T_{max}* = 0.056
 12 787 measured reflections

958 independent reflections
 787 reflections with *I* > 2σ(*I*)
R_{int} = 0.073
 $\theta_{\text{max}} = 34.6^\circ$
h = -14 → 14
k = -14 → 14
l = -13 → 13

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.034
wR(*F*²) = 0.085
S = 0.97
 958 reflections
 40 parameters
 $w = 1/[\sigma^2(F_o^2) + (0.052P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 (Δ/σ)_{max} < 0.001

$\Delta\rho_{\text{max}} = 2.04\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -1.69\text{ e \AA}^{-3}$
 Extinction correction: *SHELXTL-NT* (Bruker, 1998)
 Extinction coefficient: 0.044 (8)
 Absolute structure: Flack (1983), with 407 Friedel pairs
 Flack parameter = -0.02 (4)

The positions of the U, Rb, Na and C atoms, and of most of the O atoms, were determined by direct methods. The remaining O atoms were located by subsequent difference Fourier syntheses. The locations of the highest peak and deepest hole in the difference Fourier map are 0.75 Å from U and 0.51 Å from Rb, respectively. H-atom positions were not determined in the refinement. However, atom O2 is a possible hydrogen-bond acceptor, as it is the nearest O atom to O5 [O2...O5 = 3.122 (9) Å].

Data collection: *SMART-NT* (Bruker, 2001); cell refinement: *SAINT-NT* (Bruker, 2000); data reduction: *SAINT-NT*; program(s)

Table 1

Selected bond lengths (Å).

U—O3 ⁱ	1.779 (8)	Na—O4 ^{ix}	2.368 (9)
U—O2 ⁱⁱ	2.418 (8)	Na—O3 ^{viii}	2.401 (8)
U—O1	2.433 (6)	Na—O1 ^{viii}	3.018 (6)
Rb—O2 ⁱⁱⁱ	2.822 (6)	C—O4	1.237 (14)
Rb—O1 ^{iv}	2.897 (9)	C—O2	1.281 (14)
Rb—O4 ^v	2.992 (7)	C—O1	1.295 (17)
Rb—O5 ^{vi}	3.086 (19)		

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

used to solve structure: *SHELXTL-NT* (Bruker, 1998); program(s) used to refine structure: *SHELXTL-NT*; molecular graphics: *ATOMS* (Dowty, 2000); software used to prepare material for publication: *SHELXTL-NT* and *WinGX* (Farrugia, 1999).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: BC1032). Services for accessing these data are described at the back of the journal.

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