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Key indicators

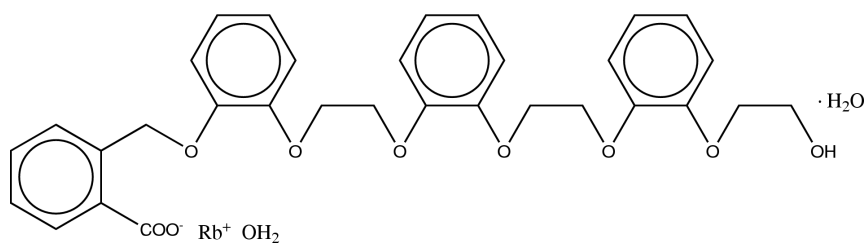
Single-crystal X-ray study
 $T = 123\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.006\text{ \AA}$
 R factor = 0.040
 wR factor = 0.115
Data-to-parameter ratio = 16.2For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.The rubidium salt of an ω -hydroxycarboxylic acid
with six ethereal O atoms

The title compound, aqua[2-({2-[2-(2-{2-[2-(2-hydroxy-ethoxy)phenoxy]ethoxy}phenoxy)ethoxy]phenoxy}methyl)-benzoato]rubidium monohydrate, $[\text{Rb}(\text{C}_{32}\text{H}_{33}\text{O}_{10})]\cdot\text{H}_2\text{O}$, shows a monomeric structure. The polyether chain encloses Rb^+ and adopts an S-like conformation. The Rb^+ centre is ninefold coordinated by one O atom of the carboxylate group, the O atom of the terminal hydroxy group, six ethereal O atoms and one water molecule. An intermolecular hydrogen bond was observed between the terminal hydroxy group and the hydrate water molecule.

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Comment

Naturally occurring acid ionophores are known to mediate active ion transport through lipophilic membranes by formation of hydrophobic complexes with ions (Taylor *et al.*, 1982, and references therein) and their structures were studied to provide information on their function (Duesler & Paul, 1983, and references therein). We have synthesized a series of relatively simple ω -hydroxycarboxylic acids as model compounds for the naturally occurring ionophores to investigate the relationship between the primary molecular structures and ion-transport properties (Kuboniwa *et al.*, 1982, 1985; Yamazaki *et al.*, 1978, 1979; Yamaguchi *et al.*, 1988, 1989).



(I)

The model compounds, (1)–(3), transport K^+ over Na^+ through the $\text{ClCH}_2\text{CH}_2\text{Cl}$ membrane using the pH gradient of outer aqueous phases. The competitive ion-transport experiments show that the amount of transported ions, as well as K^+ selectivity over Na^+ , increases in the order of (1) \rightarrow (2) \rightarrow (3) (Kuboniwa *et al.*, 1985; Yamaguchi *et al.*, 1988). The model compound, (1), having five ethereal O atoms, forms a 2:2 dimer with K^+ . The polyether chain of (1) encloses K^+ and adopts an S-like conformation. One hydroxy group, six ethereal O atoms including one from the polyether chain of the dimer, and two water molecules coordinate to K^+ and the cation has ninefold coordination. Two O atoms of the carboxylate group do not coordinate to the cation. Solvent water molecules are located between the K salt molecules

(Kasuga *et al.*, 1995). Compound (3), having eight etheral O atoms, forms 1:1 salts with K^+ , Rb^+ and Cs^+ . The conformation of the three alkali metal salts of (3) are very similar to each other. The backbone of the three salts of (3) forms a pseudocyclic ring by head-to-tail hydrogen bonding between the terminal carboxylate and hydroxy groups. The main chain of (3) is bent at O12 and O39 and wraps the cation like the seam of a tennis ball to give a complex with a lipophilic exterior. One hydroxy group, eight etheral O atoms of the main chain and one O atom of the carboxylate group participate in the coordination and K^+ is tenfold coordinated. The chloroform molecules, used as the recrystallization solvent, fill the space between the salt molecules in the crystals (Kuboniwa *et al.*, 1988; Kasuga *et al.*, 1991). Slow evaporation of a chloroform solution of the Rb salt of (2), which has six etheral O atoms and shows ion-transport properties intermediate between those of (1) and (3), afforded single crystals. This paper reports the structure of the Rb salt of (2), *i.e.* (I).

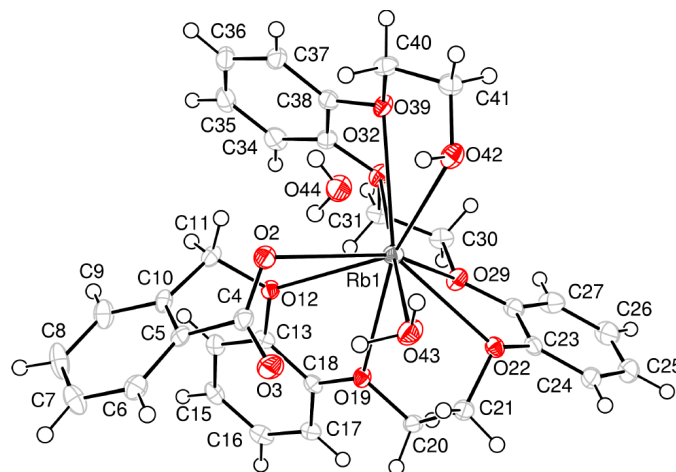
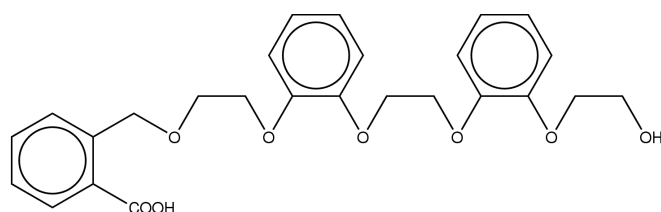
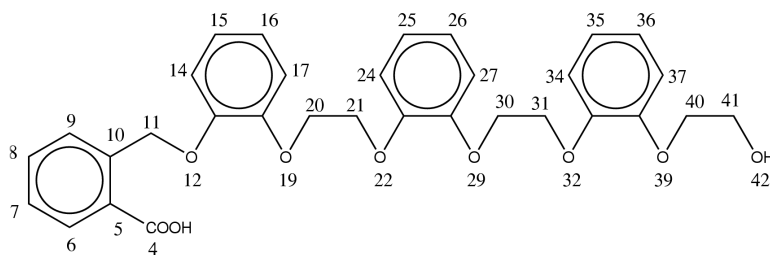


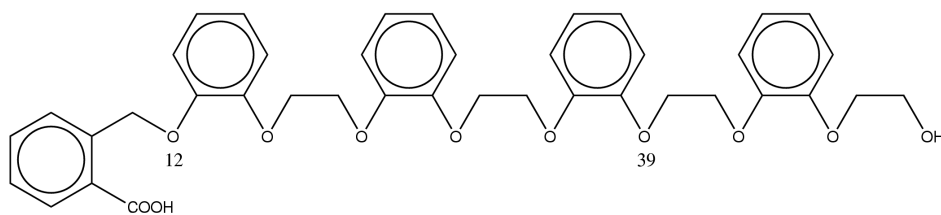
Figure 1
An ORTEP-3 drawing (Farrugia, 1997) of (I) with 50% probability displacement ellipsoids.



(1)



(2)



(3)

The molecular structure of (I) (Fig. 1) reveals that the salt forms a 1:1 monomeric structure. The Rb—O distances [2.877 (3)–3.147 (3) Å] indicate that one O atom of the carboxylate group, six etheral O atoms, the terminal hydroxy group and a water molecule coordinate to Rb^+ . As a result, Rb^+ is ninefold coordinated. Water coordination to the cation is observed in the structure of the K salt of (1), but not in the structure of the K salt of (3). In (I), another water molecule (O44) lies near the coordinating water molecule (O43), with an O43...O44 distance of 3.151 (4) Å. The O42...O44 distance of 2.724 (4) Å shows that intermolecular hydrogen bonding is formed between the terminal hydroxy group and the solvent water molecule. The head-to-tail hydrogen bonding seen in the structure of the K salt of (3) is not observed in (I). Generally, $Csp^3-Csp^3-O-Csp^2$ (quaternary) torsion angles are near 180° , Csp^2-Csp^2 (quaternary)-O- Csp^3 are 0° and O- Csp^3-Csp^3-O are *gauche*. In (I), this rule also holds except for four torsion angles, C10—C11—O12—C13 [-65.0 (4) $^\circ$], C14—C13—O12—C11 [-22.4 (5) $^\circ$], C20—C21—O22—C23 [70.4 (4) $^\circ$] and C24—C23—O22—C21 [69.8 (4) $^\circ$]. Consequently, the backbone bend of (2) in (I) occurs in a shorter period (O12 and O22) compared with that in the alkali metal salts of (3) (O12 and O39).

In conclusion, the three-dimensional structure of the Rb salt of (2) is intermediate between that of the K salt of (1) and that of the alkali metal salts of (3).

Experimental

Model compound (2) was synthesized using successive Williamson's methods (Kuboniwa *et al.*, 1985). An ion-transport experiment using (2) through a ClCH₂CH₂Cl liquid membrane was carried out in a U-tube apparatus as described previously (Yamaguchi *et al.*, 1988), however, transported ions were analyzed by atomic absorption spectrometry instead of flame analysis. The analysis showed that 22% of Na⁺ and 73% of K⁺ were transported by (2) with errors of about 5% in 5 d. A solution of RbOH in MeOH (0.54 ml, 3.74 mM) was added to the solution of (2) (11.3 mg) in 20 ml MeOH and the mixture was stirred overnight. After removal of the solvent, the solid was dried *in vacuo*. Compound (2) and its Rb salt were characterized by EA, IR, ¹H and ¹³C NMR. A chloroform solution of the Rb salt of (2) was slowly evaporated to give single crystals of (1). Crystals of the K salt of (2) were also obtained from chloroform solution, however, the quality of the reflection data was not good.

Crystal data

[Rb(C ₃₂ H ₃₃ O ₁₀)]·H ₂ O	Z = 2
M _r = 681.07	D _x = 1.50 Mg m ⁻³
Triclinic, P1̄	Mo Kα radiation
a = 10.0948 (10) Å	Cell parameters from 1053 reflections
b = 11.2408 (10) Å	θ = 2.3–13.7°
c = 14.055 (3) Å	μ = 1.70 mm ⁻¹
α = 85.524 (11)°	T = 123 (2) K
β = 71.679 (11)°	Platelet, colorless
γ = 89.852 (14)°	0.30 × 0.10 × 0.05 mm
V = 1509.0 (4) Å ³	

Data collection

Rigaku R-Axis-RAPID Imaging Plate diffractometer	6684 independent reflections
ω scans	4520 reflections with I > 2σ(I)
Absorption correction: multi-scan (ABSCOR; Higashi, 1995)	R _{int} = 0.054
T _{min} = 0.616, T _{max} = 0.919	θ _{max} = 27.3°
12 683 measured reflections	h = -13 → 13
	k = -14 → 14
	l = -18 → 18

Refinement

Refinement on F ²	H atoms treated by a mixture of independent and constrained refinement
R[F ² > 2σ(F ²)] = 0.040	w = 1/[σ ² (F _o ²) + (0.0549P) ²]
wR(F ²) = 0.115	where P = (F _o ² + 2F _c ²)/3
S = 0.94	(Δ/σ) _{max} < 0.001
6684 reflections	Δρ _{max} = 0.53 e Å ⁻³
413 parameters	Δρ _{min} = -0.52 e Å ⁻³

Table 1

Selected geometric parameters (Å, °).

O2—Rb1	3.020 (3)	O32—Rb1	3.012 (3)
O12—Rb1	2.958 (2)	O39—Rb1	3.037 (2)
O19—Rb1	2.945 (2)	O42—Rb1	2.994 (3)
O22—Rb1	2.877 (3)	O43—Rb1	3.023 (3)
O29—Rb1	3.147 (3)		
C10—C11—O12—C13	-65.0 (4)	C24—C23—O22—C21	69.8 (4)
C21—C20—O19—C18	178.4 (3)	C27—C28—O29—C30	5.6 (5)
C20—C21—O22—C23	70.4 (4)	C34—C33—O32—C31	0.7 (5)
C31—C30—O29—C28	-178.8 (3)	C37—C38—O39—C40	-2.9 (5)
C30—C31—O32—C33	-170.6 (3)	O19—C20—C21—O22	65.6 (4)
C41—C40—O39—C38	-175.0 (3)	O29—C30—C31—O32	-64.8 (4)
C14—C13—O12—C11	-22.4 (5)	O39—C40—C41—O42	62.6 (4)
C17—C18—O19—C20	-1.5 (5)		

Table 2

Hydrogen-bonding geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
O42—H42...O44	0.807 (19)	1.93 (2)	2.724 (4)	169 (4)

H atoms attached to C atoms were included in idealized positions and refined using a riding model (C—H = 0.93 Å). H atoms attached to O atoms were found in the difference Fourier maps and their positional parameters were refined with restraints on the O—H distances. The O—H distances were 0.803 (18)–1.00 (4) Å.

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Molecular Structure Corporation and Rigaku, 2001); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); 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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