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# 1-Deoxy-1-(4-fluorophenyl)- $\beta$ -D-ribofuranose, its hemihydrate, and 1-deoxy-1-(2,4-difluorophenyl)- $\beta$ -Dribofuranose: structural evidence for intermolecular C—H···F—C interactions

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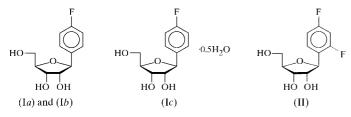
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The structures of 1-deoxy-1-(4-fluorophenyl)- $\beta$ -D-ribofuranose in two crystal forms, (Ia) and (Ib) (C<sub>11</sub>H<sub>13</sub>FO<sub>4</sub>), 1-deoxy-1-(4-fluorophenyl)- $\beta$ -D-ribofuranose hemihydrate, (Ic) (C<sub>11</sub>H<sub>13</sub>-FO<sub>4</sub>·0.5H<sub>2</sub>O) and 1-deoxy-1-(2,4-difluorophenyl)- $\beta$ -D-ribofuranose, (II) (C<sub>11</sub>H<sub>12</sub>F<sub>2</sub>O<sub>4</sub>), show two-dimensional networks of intermolecular hydrogen bonds between the hydroxyl groups. Weak intermolecular C-H···F-C and C-H··· $\pi_{arene}$  interactions complete the packing in the third dimension. The ribofuranose ring has a conformation intermediate between a C1'-*exo*,C2'-*endo* twist and a C2'-*endo* envelope for (Ia) and (Ic), a conformation intermediate between a C2'-*endo*,C3'-*exo* twist and a C2'-*endo* twist conformation for (II).

# Comment

This paper presents the crystal structures of four ribofuranosebased compounds, namely, 1-deoxy-1-(4-fluorophenyl)- $\beta$ -Dribofuranose in two crystal forms, (I*a*) and (I*b*), 1-deoxy-1-(4fluorophenyl)- $\beta$ -D-ribofuranose hemihydrate, (I*c*), and 1deoxy-1-(2,4-difluorophenyl)- $\beta$ -D-ribofuranose, (II).



The five-membered ribofuranose rings in (I*a*) and (I*c*) (Fig. 1) have rather similar conformations. The ring puckering parameters defined by Cremer & Pople (1975) are q = 0.375 Å and  $\varphi = 60.2^{\circ}$  for (I*a*), and q = 0.421 Å and  $\varphi = 63.4^{\circ}$  for (I*c*). This corresponds to a conformation intermediate between a

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Cl'*-exo*,C2'*-endo* twist and a C2'*-endo* envelope. The hydroxyl group at C2 and the phenyl group attached to C1 are both in pseudo-equatorial positions, while the hydroxyl group at C3 is in a pseudo-axial position and the methanol group attached to C4 is in a bisecting position with respect to the five-membered ring. No short intramolecular contacts are observed for this conformation.

The five-membered ribofuranose ring in (Ib) (Fig. 1) is considerably different. It has a conformation intermediate between a C2'-endo,C3'-exo twist and a C2'-endo envelope. The ring puckering parameters are q = 0.368 Å and  $\varphi = 81.6^{\circ}$ . The hydroxyl group on C2 and the phenyl group attached to C1 are in pseudo-equatorial positions, while the hydroxyl group at C3 and the methanol group attached to C4 are both in pseudo-axial positions with respect to the five-membered ring.

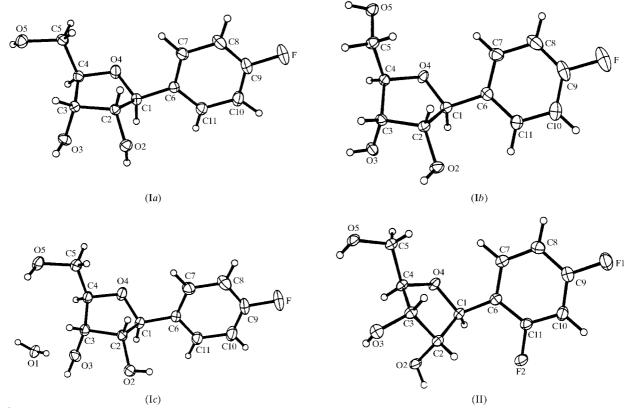
The five-membered ribofuranose ring in (II) (Fig. 1) is different again. It approximately has an unsymmetrical C2'exo,C3'-endo twist conformation. The ring puckering parameters are q = 0.390 Å and  $\varphi = 274.8^{\circ}$ . The hydroxyl group at C2 is in an axial position, the hydroxyl group at C3 is in an equatorial position, the methanol group attached to C4 is in a pseudo-equatorial position and the phenyl group attached to C1 is in a pseudo-axial position with respect to the fivemembered ring.

Molecules (Ib) and (II) show a short intramolecular contact distance of 2.42 (2) for (Ib) and 2.44 (2) Å for (II) between O4 and H7. This distance is equal to the van der Waals contact distance of 2.4 Å between O and H. Each molecule of (Ia) and (Ib) is connected by intermolecular hydrogen bonding (Tables 1 and 2, Figs. 2 and 3) to six neighbouring molecules, leading to a two-dimensional network in the *ab* direction.

For (Ic), a rather similar two-dimensional network is found in the bc direction. In this case, the water molecule is included in the network. Each molecule of (Ic) is connected by hydrogen bonding (Table 3, Fig. 4) to three neighbouring molecules of (Ic) and to two different water molecules.

Each molecule of (II) is connected by intermolecular hydrogen bonding (Table 4, Fig. 5) to four different neighbouring molecules, leading to a two-dimensional network in the *ab* direction. Two-dimensional networks of hydrogen bonds have also been observed in the crystal structures of 1deoxy-1-phenyl- $\beta$ -D-ribofuranose (Matulic-Adamic *et al.*, 1996), 1-deoxy-1-(2-fluorophenyl)- $\beta$ -D-ribofuranose (Bats *et al.*, 1999*a*) and 1-deoxy-1-(3-fluorophenyl)- $\beta$ -D-ribofuranose (Bats *et al.*, 1999*b*).

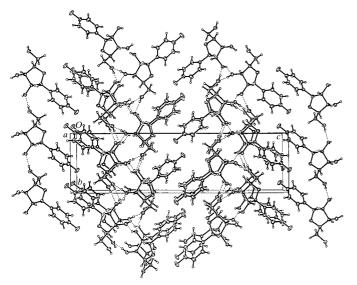
The packing in the direction of the long dimension of the unit cell consists entirely of intermolecular interactions between the fluorophenyl groups. Rather short intermolecular  $C-H\cdots F-C$  contacts are observed in (Ia), (Ib) and (Ic). The intermolecular H10 $\cdots$ F distance is 2.30 (2) in (Ia), 2.42 (2) in (Ib) and 2.38 (2) Å in (Ic). These distances are comparable to the van der Waals contact distance between F and H [2.35 Å according to Pauling (1960) or 2.47 Å according to Bondi (1964)]. Each 4-fluorophenyl group is involved in two of these contacts, resulting in zigzag chains of molecules (Fig. 6). In addition to these  $C-H\cdots F-C$  interactions, the crystal

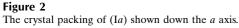


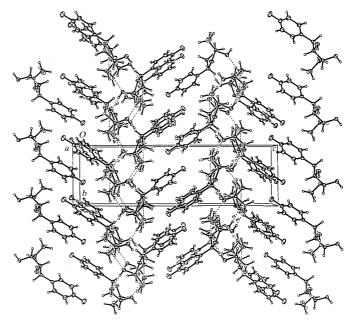
#### Figure 1

The molecules of (Ia), (Ib), (Ic) and (II) with 50% probability displacement ellipsoids. H atoms are shown as spheres of arbitrary radii.

packing of (I*a*), (I*b*) and (I*c*) also shows intermolecular C– H··· $\pi_{arene}$  interactions. The H8···*Cg* distance is 2.82 in (I*a*), 2.84 in (I*b*) and 2.82 Å in (I*c*) (*Cg* is the centroid of the phenyl ring). Intermolecular C–H·· $\pi_{arene}$  interactions play a significant role in crystal structures (Brady *et al.*, 1998; Gallagher *et al.*, 1998). Koch & Egert (1995) have shown these interactions to be of an electrostatic nature. The crystal structure of (II) shows pairs of molecules related by a twofold axis, with intermolecular distances of 2.57 (2) Å between F1 and H10(-x, y, -z) (Fig. 7). These distances are longer than the van der Waals contact distance. No other intermolecular interactions between the phenyl groups are

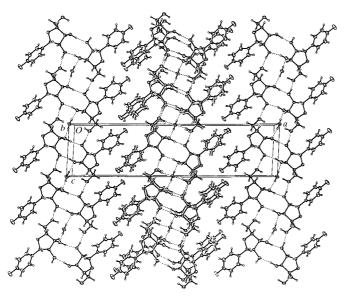






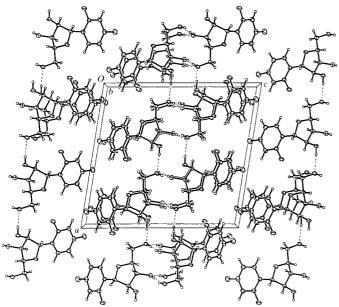
**Figure 3** The crystal packing of (I*b*) shown down the *a* axis.

found in this case, which can explain the crystal packing in the *c* direction. Intermolecular  $F \cdots H$  contacts have also been reported for other crystal structures (Karipides *et al.*, 1988; Bruno *et al.*, 1997; Antolić *et al.*, 1996; Vij *et al.*, 1996; Shimoni *et al.*, 1994; Weiss *et al.*, 1997). A search of the Spring 1999 release of the Cambridge Structural Database (Allen & Kennard, 1993) for structures containing fluorophenyl groups, with H-atom coordinates included and no disorder reported, revealed 1144 crystal structures. Among these structures, 1596 different  $C-H \cdots F-C$  contacts were found with  $H \cdots F$  distances shorter than 2.6 Å (652 shorter than 2.5, 182 shorter than 2.4 and 37 shorter than 2.3 Å). About one third of these interactions were  $C_{phenyl}-H \cdots F-C_{phenyl}$  contacts. The



### Figure 4

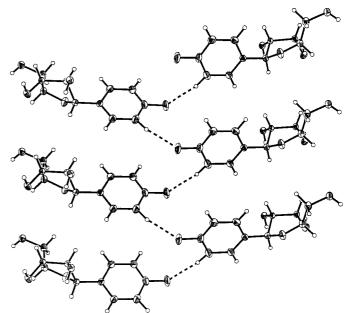
The crystal packing of (Ic) shown down the b axis.



**Figure 5** The crystal packing of (II) shown down the *b* axis.

number of dimeric structures as found for (II) is rather small. Only 35 dimers were found with both  $C-H\cdots F-C$  contacts shorter than 2.6 Å. Larsen & Marthi (1994) and Kubota & Ohba (1992) have shown the F atom in fluorophenyl groups to bear a small negative charge of about  $-0.1 \text{ e } \text{Å}^{-3}$ . Thus, the C-F bond is a likely candidate for electrostatic interactions with electropositive C-H bonds.

Howard et al. (1996) calculated the C-H···F-C interaction energy of a complex between CH<sub>3</sub>F and H<sub>4</sub>C to be approximately  $-0.2 \text{ kcal mol}^{-1}$  (1 cal = 4.184 J), which is an order of magnitude smaller than the corresponding value for a complex between CH<sub>3</sub>F and H<sub>2</sub>O. They termed the C- $H \cdot \cdot \cdot F - C$  interaction as a van der Waals interaction rather than a hydrogen bond. Thalladi et al. (1998) showed the packing in fluorobenzenes containing no other elements than C, H and F to be dominated by  $C-H \cdots F-C$  interactions. They also showed the  $C-H \cdot \cdot F$  interactions to have similar characteristics to those of weak C-H···O and C-H···N hydrogen bonds. These interactions are rather weak and thus are easily overridden by stronger intermolecular interactions. In the crystal structures reported here, however, stronger interactions are missing in one direction. Thus, weaker  $C-H\cdots F-C$  and  $C-H\cdots \pi$  interactions dominate the crystal packing in this direction.



#### Figure 6

Zigzag chains of molecules connected by  $C-H\cdots F$  interactions in (I*a*). Similar chains occur in (I*b*) and (I*c*).

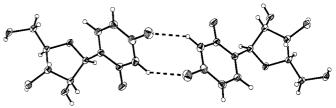


Figure 7 Dimers of molecules of (II) connected by  $C-H\cdots F$  interactions.

# **Experimental**

The title compounds were prepared according to Krohn et al. (1992). Thus, 1-bromo-4-fluorobenzene [for (I)] or 1-bromo-2,4-difluorobenzene [for (II)] was lithiated and added to 2',3',5'-tri-O-benzyl-Dribono-1,4-lactone (Barker & Fletcher, 1961; Timpe et al., 1975). The resulting intermediate lactol was directly dehydroxylated to yield a sterically uniform  $\beta$ -C-riboside. Debenzylation of the hydroxyl groups resulted in the unprotected C-nucleoside. Compound (I) was recrystallized from methanol, resulting in two different crystalline modifications, (Ia) and (Ib), while recrystallization from water yielded the hemihydrate (Ic). Compound (II) was also recrystallized from water.

### Compound (Ia)

#### Crystal data

 $C_{11}H_{13}FO_4$  $M_r = 228.21$ Orthorhombic,  $P2_12_12_1$ a = 5.4741 (9) Åb = 7.3018 (7) Å c = 26.791(3) Å V = 1070.9 (2) Å<sup>3</sup> Z = 4 $D_x = 1.416 \text{ Mg m}^{-3}$ 

#### Data collection

Siemens SMART diffractometer  $\omega$  scans 18 913 measured reflections 3431 independent reflections 3273 reflections with  $I > 2\sigma(I)$  $R_{\rm int}=0.084$  $\theta_{\rm max} = 31.3^{\circ}$ 

#### Refinement

Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.037$  $wR(F^2) = 0.111$ S = 1.3933431 reflections 197 parameters

Table 1 Hydrogen-bonding geometry (Å, °) for (Ia).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$\begin{array}{c} O2 - HO2 \cdots O3^{i} \\ O3 - HO3 \cdots O5^{ii} \\ O5 - HO5 \cdots O2^{iii} \end{array}$	0.85 (2) 0.88 (2) 0.80 (2)	1.96 (2) 1.91 (2) 1.89 (2)	2.729 (1) 2.774 (1) 2.682 (1)	151 (2) 169 (2) 172 (2)

Symmetry codes: (i)  $1 - x, \frac{1}{2} + y, \frac{3}{2} - z$ ; (ii)  $2 - x, \frac{1}{2} + y, \frac{3}{2} - z$ ; (iii) x, y - 1, z.

# Compound (Ib)

Crystal data

C11H13FO4  $M_r = 228.21$ Orthorhombic,  $P2_12_12_1$ a = 6.7109 (14) Åb = 6.7653 (7) Åc = 22.846 (3) Å V = 1037.2 (3) Å<sup>3</sup> Z = 4 $D_x = 1.461 \text{ Mg m}^{-3}$ 

Mo  $K\alpha$  radiation Cell parameters from 161 reflections  $\theta = 3-23^{\circ}$  $\mu = 0.122 \text{ mm}^{-1}$ T = 135 (2) KRod, colourless  $0.66 \times 0.45 \times 0.28 \text{ mm}$ 

#### Data collection

Siemens SMART diffractometer	$R_{\rm int} = 0.058$
$\omega$ scans	$\theta_{\rm max} = 31.9^{\circ}$
Absorption correction: numerical	$h = -9 \rightarrow 9$
(SHELXTL; Sheldrick, 1996)	$k = -10 \rightarrow 10$
$T_{\min} = 0.932, T_{\max} = 0.968$	$l = -34 \rightarrow 31$
19 895 measured reflections	275 standard reflections
3512 independent reflections	frequency: 540 min
2944 reflections with $I > 2\sigma(I)$	intensity decay: none
Refinement	
Refinement on $F^2$	$w = 1/[\sigma^2(F_0^2) + (0.05P)^2]$ where
$R[F^2 > 2\sigma(F^2)] = 0.039$	$w = 1/[\sigma^2(F_o^2) + (0.05P)^2] \text{ where} P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.090$	$\Delta \rho_{\rm max} = 0.30 \text{ e} \text{ \AA}^{-3}$
S = 1.114	$\Delta \rho_{\rm A} = -0.20  {\rm e}  {\rm \AA}^{-3}$

 $wR(F^2) = 0.090$ S = 1.1143512 reflections 198 parameters All H-atom parameters refined

### Table 2

Hydrogen-bonding geometry (Å,  $^{\circ}$ ) for (Ib).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdots A$
$\begin{array}{c} O2 - HO2 \cdots O5^{i} \\ O3 - HO3 \cdots O2^{ii} \\ O5 - HO5 \cdots O3^{iii} \end{array}$	0.83 (2)	1.91 (2)	2.706 (1)	160 (1)
	0.79 (2)	1.94 (2)	2.721 (1)	170 (1)
	0.88 (2)	1.91 (2)	2.782 (1)	174 (1)

 $\Delta \rho_{\rm min} = -0.20 \ {\rm e} \ {\rm \AA}^{-3}$ 

(Sheldrick, 1997)

 $h = -40 \rightarrow 37$ 

207 standard reflections

frequency: 540 min

intensity decay: none

 $k = -7 \rightarrow 6$  $l = -9 \rightarrow 10$ 

Extinction correction: SHELXL97

Extinction coefficient: 0.010 (3)

Symmetry codes: (i) x - 1, y, z; (ii)  $-x, y - \frac{1}{2}, \frac{3}{2} - z$ ; (iii)  $1 - x, y - \frac{1}{2}, \frac{3}{2} - z$ .

### Compound (Ic)

Crystal data

$D_x = 1.378 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation
Cell parameters from 185
reflections
$\theta = 3-23^{\circ}$
$\mu = 0.116 \text{ mm}^{-1}$
T = 154 (2) K
Block, colourless
$0.45\times0.44\times0.40$ mm

Data collection

Siemens SMART diffractometer  $\omega$  scans 5566 measured reflections 2827 independent reflections 2746 reflections with  $I > 2\sigma(I)$  $R_{\rm int} = 0.041$  $\theta_{\rm max} = 29.0^{\circ}$ 

## Refinement

 $w = 1/[\sigma^2(F_o^2) + (0.05P)^2] \text{ where } P = (F_o^2 + 2F_c^2)/3$ Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.034$   $wR(F^2) = 0.100$  $\Delta \rho_{\rm max} = 0.35 \ {\rm e} \ {\rm \AA}^{-3}$  $\Delta \rho_{\rm min} = -0.17 \ {\rm e} \ {\rm \AA}^{-3}$ S = 1.4862827 reflections Extinction correction: SHELXL97 207 parameters (Sheldrick, 1997) All H-atom parameters refined Extinction coefficient: 0.006 (2)

# Table 3

Hydrogen-bonding geometry (Å, °) for (Ic).

$D-\mathrm{H}\cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$\begin{array}{c} O1 - HO1 \cdots O3^{i} \\ O2 - HO2 \cdots O5^{ii} \\ O3 - HO3 \cdots O2^{iii} \\ O5 - HO5 \cdots O1^{iv} \end{array}$	0.84 (2)	1.86 (2)	2.701 (1)	176 (2)
	0.88 (2)	1.79 (2)	2.661 (1)	167 (2)
	0.92 (2)	1.77 (2)	2.669 (1)	168 (2)
	0.93 (2)	1.85 (2)	2.777 (1)	172 (2)

Symmetry codes: (i) 1 - x, y, -z; (ii) x, y, 1 + z; (iii) 1 - x, y, 1 - z; (iv) x, y - 1, z.

$\theta = 3-23^{\circ}$
$\mu = 0.118 \text{ mm}^{-1}$
T = 134 (2)  K
Plate, colourless
$1.30 \times 0.64 \times 0.15 \text{ mm}$

Mo  $K\alpha$  radiation

reflections

 $h = -7 \rightarrow 8$  $k = -10 \rightarrow 10$ 

Cell parameters from 279

$l = -38 \rightarrow 35$
369 standard reflections
frequency: 540 min
intensity decay: none

All H-atom parameters refined  $w = 1/[\sigma^2(F_o^2) + (0.06P)^2]$  where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\rm max} = 0.002$  $\Delta \rho_{\rm max} = 0.36 \ {\rm e} \ {\rm \mathring{A}}^{-3}$  $\Delta \rho_{\rm min} = -0.20 \ {\rm e} \ {\rm \AA}^{-3}$ 

# Compound (II)

#### Crystal data

 $C_{11}H_{12}F_2O_4$  $M_r = 246.21$ Monoclinic, C2 a = 14.684 (2) Åb = 4.6609 (6) Å c = 15.772 (2) Å  $\beta = 99.476 (11)^{\circ}$  $V = 1064.7 (2) \text{ Å}^3$ Z = 4Data collection Siemens SMART diffractometer  $\omega$  scans 9578 measured reflections 3048 independent reflections 2794 reflections with  $I > 2\sigma(I)$  $R_{\rm int} = 0.068$  $\theta_{\rm max} = 30.6^\circ$ 

#### Refinement

Refinement on $F^2$	All H-atom parameters refined
$R[F^2 > 2\sigma(F^2)] = 0.046$	$w = 1/[\sigma^2(F_o^2) + (0.06P)^2]$ where
$wR(F^2) = 0.122$	$P = (F_o^2 + 2F_c^2)/3$
S = 1.400	$\Delta \rho_{\rm max} = 0.37 \ {\rm e} \ {\rm \AA}^{-3}$
3048 reflections	$\Delta \rho_{\rm min} = -0.25 \text{ e} \text{ Å}^{-3}$
202 parameters	

 $D_x = 1.536 \text{ Mg m}^{-3}$ 

Cell parameters from 217

Thick plate, colourless

 $0.75 \times 0.70 \times 0.20 \text{ mm}$ 

183 standard reflections

frequency: 540 min

intensity decay: none

Mo  $K\alpha$  radiation

reflections

T = 134 (2) K

 $h = -20 \rightarrow 21$ 

 $k = -6 \rightarrow 6$ 

 $l = -22 \rightarrow 21$ 

 $\theta = 3-23^{\circ}$  $\mu = 0.138 \text{ mm}^{-1}$ 

For all compounds, data collection: *SMART* (Siemens, 1995); cell refinement: *SMART*; data reduction: *SAINT* (Siemens, 1995); program(s) used to solve structure: *SHELXTL* (Sheldrick, 1996); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997);

#### Table 4

Hydrogen-bonding geometry (Å, °) for (II).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdots A$
$\begin{array}{c} O2 - HO2 \cdots O5^{i} \\ O3 - HO3 \cdots O5^{ii} \\ O5 - HO5 \cdots O3^{ii} \end{array}$	0.82 (2)	1.97 (2)	2.780 (1)	171 (2)
	0.76 (2)	2.07 (2)	2.827 (1)	174 (2)
	0.81 (2)	1.92 (2)	2.724 (1)	171 (2)

Symmetry codes: (i)  $\frac{1}{2} + x$ ,  $y - \frac{1}{2}$ , z; (ii)  $-\frac{1}{2} - x$ ,  $y - \frac{1}{2}$ , 1 - z.

molecular graphics: *XP* in *SHELXTL*; software used to prepare material for publication: *CIF* in *SHELXL*97.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: KA1343). Services for accessing these data are described at the back of the journal.

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