

ment of the Ag atom from the centre of symmetry depending on the orientation of the neighbouring CN groups.

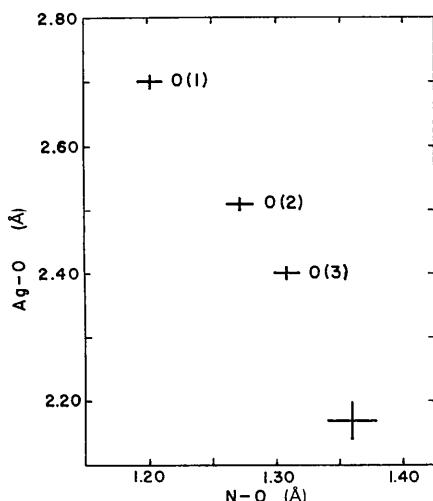


Fig. 3. N-O bond lengths *versus* Ag-O distances. The points are labelled with the numbers from Table 2. The unlabelled point corresponds to an N-O single bond and a Ag-O single bond, from CH_3ONO and $\text{Ag}_2\text{C}_2\text{O}_4$ respectively.

This work was performed in part during the tenure of a fellowship for which D.B. would like to thank the National Science Foundation. The preliminary calculations were carried out on the IBM 1620 computer of this laboratory, using programs prepared by M. Dobler, H. C. Mez, P. Strickler, and H. P. Weber. The least-squares calculations were carried out on the CDC 1604 computer at the Numerical Analysis Center of the University of Minnesota, using programs prepared at Princeton University under the direction of Professor R. Jacobson. This part of the work was performed by Mr E. O. Schlemper and Mrs Judith Konnert, and supported by a grant from the National Science Foundation. We thank them all for their help.

References

- ARNDT, U. W. & PHILLIPS, D. C. (1961). *Acta Cryst.* **14**, 807.
- BRITTON, D. & DUNITZ, J. D. (1965). *Acta Cryst.* **19**, 662.
- LINDQVIST, I. (1954). *Acta Cryst.* **7**, 635.
- PROSEN, R. J. & TRUEBLOOD, K. N. (1956). *Acta Cryst.* **9**, 741.
- SUTTON, L. E. (1958). *Interatomic Distances*. London: The Chemical Society.
- SWANSON, H. E., GILFRICH, N. T. & UGRINIC, G. M. (1955). N.B.S. Circular 539, vol. V, p. 59.
- WEST, C. D. (1935). *Z. Kristallogr.* **90**, 555.

Acta Cryst. (1965). **19**, 820

The Crystal Structure of α -Methyl D-Galactoside 6-Bromohydrin

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The crystal structure of α -methyl D-galactoside 6-bromohydrin has been determined by the heavy-atom method, at 125 °K. The bromine atom coordinates were derived from Harker sections. Three-dimensional structure factors and least-squares refinement of 1203 reflexions with anisotropic temperature factors gave a final residual $R=0.108$. All bonds, including the C(1)-O(1) bond, were found to be of normal length. The positions of the hydrogen atoms were found from the final 3-D Fourier synthesis. The hydrogen bonding system, which gives good agreement with the infrared spectroscopic data, consists of spiral linkages about the screw axes parallel to the *b* axis.

Experimental

α -Methyl D-galactoside 6-bromohydrin was prepared by Valentin (1952); unit-cell and density measurements were made by Cox, Goodwin & Wagstaff (1935). The unit-cell dimensions were re-measured at 125 °K by the extrapolation to $\theta=90^\circ$ of high order reflexions on zero layer Weissenberg photographs, calibrated with aluminum wire powder lines. The *a* and *c* axes of Cox

et al. were interchanged for convenience. The unit cell is orthorhombic with systematic absences of *h*00 for *h* odd and 0*k*0 for *k* odd. The space group is $P2_12_12$ and the cell dimensions are:

$a = 11.142 \pm 0.005$ Å	<i>cf.</i> Cox, <i>et al.</i>	11.23 Å
$b = 7.815 \pm 0.003$	at room temperature	7.81
$c = 10.612 \pm 0.010$		10.58

The value of the density observed, 1.86 g.cm^{-3} , gave a cell weight of 1035 and hence $Z=4.03$ ($\text{C}_7\text{H}_{13}\text{BrO}_5 = 257$).

The material was recrystallized from water, on a greased microscope slide, and a crystal with dimensions

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0.25 mm × 0.25 mm × 0.21 mm was selected. Full three-dimensional data were collected with the use of Cu $K\alpha$ radiation and the Weissenberg equi-inclination technique. The crystal was cooled to 125 °K by a stream of dry, cold nitrogen gas from apparatus similar to that described previously (Robertson, 1960). 1203 independent hkl reflexions were measured by the multiple film technique and correlated by hand calculation. 62 reflexions (marked with an asterisk in Table 6) were too weak to be measured. These were included in the least-squares refinement as $\frac{1}{2}F_{min}$ but without separate weighting.

Solution of the structure

The position of the bromine atom was found from the Patterson-Harker sections at $x=\frac{1}{2}$, $y=\frac{1}{2}$ and $z=0$. Examination of the three mutually perpendicular Pat-

terson projections showed that the x coordinate was general (at 5/60th a) but y was approximately $\frac{1}{4}$ and $z=0$ (or $\frac{1}{2}$). From the Harker sections it was possible to establish that the bromine atom coordinates were not quite so unfavourable as was first feared. The coordinates were specified as $x=5/60$, $y=14/60$ and $z=1/120$. A three-dimensional Fourier synthesis was computed on the Leeds University Ferranti Pegasus computer by the program written by Cruickshank, Pilling, Bujosa, Lovell & Truter (1961), with the phase angles specified by the bromine atoms. The resultant synthesis contained some spurious symmetry due to the near-special position of the bromine atom. Five definite light atom peaks were selected, two cycles of structure-factor least-squares (SFLS) refinement were carried out with the program written by Cruickshank *et al.* (1961) and the modified phase angles were used to

Table 1. Atomic coordinates and estimated standard deviations (\AA)

	x	σ	y	σ	z	σ
Br	0.8976	0.0014	1.7809	0.0015	-0.1230	0.0016
O(1)	2.855	0.007	0.535	0.008	7.511	0.009
O(2)	9.663	0.008	0.083	0.008	5.211	0.007
O(3)	5.692	0.008	1.726	0.008	4.363	0.009
O(4)	3.568	0.007	0.062	0.008	3.404	0.008
O(5)	7.409	0.007	1.627	0.009	7.671	0.009
C(1)	7.879	0.011	0.624	0.011	6.781	0.011
C(2)	9.079	0.010	1.132	0.014	6.020	0.013
C(3)	10.184	0.010	1.627	0.012	6.983	0.014
C(4)	4.033	0.010	1.223	0.012	2.722	0.013
C(5)	2.851	0.010	1.828	0.012	1.984	0.011
C(6)	2.197	0.013	0.865	0.011	1.046	0.016
C(7)	3.940	0.012	1.200	0.015	8.178	0.015

Table 2. Anisotropic temperature factors and estimated standard deviations

	U_{11}	U_{22}	U_{33}	U_{12}	U_{13}	U_{23}
Br	0.0308	0.0242	0.0363	0.0036	0.0074	-0.0169
O(1)	0.0202	0.0147	0.0315	-0.0179	-0.0017	-0.0134
O(2)	0.0238	0.0098	0.0328	0.0016	-0.0134	0.0142
O(3)	0.0202	0.0135	0.0388	0.0099	-0.0156	-0.0128
O(4)	0.0121	0.0106	0.0392	0.0139	0.0035	-0.0053
O(5)	0.0137	0.0180	0.0350	0.0111	0.0021	-0.0020
C(1)	0.0235	0.0135	0.0268	-0.0227	0.0021	0.0035
C(2)	0.0133	0.0312	0.0290	-0.0089	0.0049	0.0087
C(3)	0.0112	0.0177	0.0454	-0.0061	0.0007	0.0137
C(4)	0.0134	0.0175	0.0426	-0.0049	0.0165	-0.0035
C(5)	0.0196	0.0217	0.0169	-0.0119	0.0102	-0.0085
C(6)	0.0299	0.0090	0.0532	-0.0049	0.0117	-0.0279
C(7)	0.0159	0.0401	0.0471	-0.0372	-0.0109	-0.0100
	σ_{11}	σ_{22}	σ_{33}	σ_{12}	σ_{13}	σ_{23}
Br	0.0007	0.0007	0.0007	0.0010	0.0011	0.0010
O(1)	0.0037	0.0035	0.0042	0.0066	0.0074	0.0069
O(2)	0.0040	0.0032	0.0042	0.0060	0.0085	0.0071
O(3)	0.0034	0.0033	0.0042	0.0068	0.0086	0.0073
O(4)	0.0031	0.0031	0.0043	0.0058	0.0084	0.0066
O(5)	0.0032	0.0034	0.0040	0.0060	0.0088	0.0068
C(1)	0.0052	0.0045	0.0054	0.0085	0.0096	0.0091
C(2)	0.0040	0.0062	0.0054	0.0101	0.0117	0.0086
C(3)	0.0040	0.0048	0.0067	0.0086	0.0112	0.0090
C(4)	0.0043	0.0050	0.0067	0.0091	0.0109	0.0094
C(5)	0.0047	0.0052	0.0040	0.0093	0.0085	0.0073
C(6)	0.0059	0.0044	0.0080	0.0102	0.0108	0.0123
C(7)	0.0043	0.0073	0.0071	0.0121	0.0149	0.0110

compute a second three-dimensional Fourier synthesis. In this all the atoms of the structure, with the exception of the carbon atom of the methyl group, could be distinguished. Seven cycles of SFLS were carried out with isotropic temperature factors for all the atoms except the methyl group carbon atom. This atom was then found by computing a difference Fourier synthesis for the sections where it was thought the methyl group should occur.

This atom was then introduced and four cycles of SFLS refinement were carried out with anisotropic temperature factors. Form factors published by Berghuis, Haanappel, Potters, Loopstra, MacGillavry & Veenendaal were used for carbon and oxygen atoms and by Thomas & Umeda (1957) for bromine. The weighting scheme used was $w = 1/(2F_{\min} + F + 2F^2/F_{\max})$. The final cycle gave a residual, $R = 0.108$ and the observed and calculated structure factors are tabulated in Table 6.

A final three-dimensional Fourier synthesis was carried out and examined for variations of electron density. Some small positive regions were noted, of peak heights $0.7-1.2 \text{ e.}\text{\AA}^{-3}$. Some of these were in positions in which would be plausible for hydrogen atoms to occur. As confirmation, the standard deviation of the electron density was calculated by the formulae given by Cruickshank (1949) and by Cruickshank & Rollett (1953):

$$\sigma(\rho) = \frac{1}{V} \{ \sum (\Delta F)^2 \}^{1/2}$$

using a program written by one of us (B.S.) and found to be

$$\sigma(\rho) = 0.37 \text{ e.}\text{\AA}^{-3},$$

i.e. a peak height of $0.8 \text{ e.}\text{\AA}^{-3}$ would probably be significant. Thirteen possible hydrogen atoms were found and their positions were accurate enough to define the hydrogen bonding system. The coordinates of the atoms of one molecule are given in Table 1, together with the estimated standard deviations. The anisotropic temperature factors and their estimated standard deviations are given in Table 2, and the positions of the hydrogen atoms in Table 3.

Attempts were made to refine the data further by including the hydrogen atoms isotropically ($R = 0.105$)

Table 3. Atomic coordinates of hydrogen atoms (unrefined) (\AA)

	x	y	z
H(1)	4.04	0.80	6.19
H(2)	8.73	1.91	5.40
H(3)	10.46	0.78	7.57
H(4)	4.74	1.17	2.03
H(5)	8.73	1.43	9.37
H(6)	1.76	1.40	1.68
H(7)	1.80	0.00	8.91
H(8)	4.36	1.04	9.29
H(9)	4.55	1.04	7.16
H(10)	8.91	1.82	1.77
H(11)	6.59	1.04	4.51
H(12)	4.77	0.00	3.86
H(13)	9.42	0.13	6.37

and by correcting the data (40 reflexions) for the presence of secondary extinction ($R = 0.096$). These measures had little effect on the atomic coordinates and the reduction of the e.s.d.'s was comparable to the reduction of the residual R . The stereochemical arrangement is shown in Fig. 1.

The bond lengths, bond angles and related e.s.d.'s were calculated by a program written by Dr Mary R. Truter and the results are shown in Figs. 2 and 3 and Tables 4 and 5.

Structure of the molecule

The molecule has the expected chair form with the configuration $1a2e3e4a$, as found for β -arabinose by Fur-

Table 4. Bond lengths and estimated standard deviations

Bond	$l(\text{\AA})$	e.s.d. (\AA)
Br-C(6)	1.974	0.014
C(1)-C(2)	1.509	0.016
C(2)-C(3)	1.551	0.017
C(3)-C(4)	1.502	0.017
C(4)-C(5)	1.524	0.015
C(5)-C(6)	1.492	0.018
O(1)-C(1)	1.430	0.014
O(1)-C(7)	1.436	0.016
O(2)-C(2)	1.448	0.015
O(3)-C(3)	1.414	0.014
O(4)-C(4)	1.425	0.014
O(5)-C(5)	1.461	0.013
O(5)-C(1)	1.421	0.014

Table 5. Bond angles and estimated standard deviations

Angle ($^\circ$)	e.s.d.
Br-C(6)-C(5)	111°02'
O(5)-C(1)-C(2)	109 56
O(5)-C(1)-O(1)	110 16
O(1)-C(1)-C(2)	107 39
C(1)-C(2)-C(3)	111 18
C(1)-C(2)-O(2)	111 02
O(2)-C(2)-C(3)	107 09
C(2)-C(3)-C(4)	109 05
C(2)-C(3)-O(3)	110 19
O(3)-C(3)-C(4)	109 49
C(3)-C(4)-C(5)	108 28
C(3)-C(4)-O(4)	113 55
O(4)-C(4)-C(5)	107 26
C(4)-C(5)-C(6)	112 46
C(4)-C(5)-O(5)	110 03
C(6)-C(5)-O(5)	108 13
C(5)-O(5)-C(1)	113 40
C(1)-O(1)-C(7)	113 20

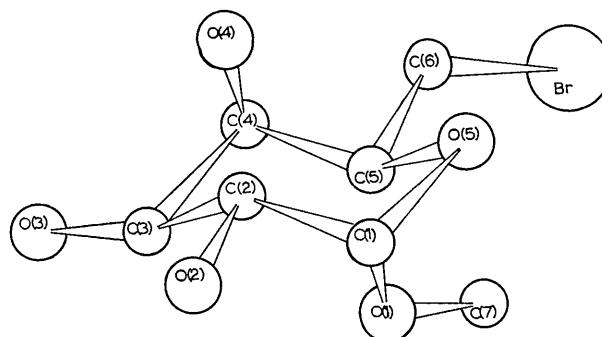


Fig. 1. Stereochemical arrangement.

Table 6. Observed and calculated structure factors

In each group headed by the values of h and k , the columns give l , $10|F_0|$, $10F_c$, $10A_c$, $10B_c$.

1	234	311	311	81	66	-66	9	212	214	-212	-69	6	118	130	-96	44	136	157	157	-54	115	
352	485	485	485	105	105	-105	105	105	105	-105	105	9	193	183	-153	105	111	121	121	58	77	
956	1146	1146	1146	105	105	-105	105	105	105	-105	105	105	157	189	-115	157	111	124	124	58	77	
303	488	488	488	105	105	-105	105	105	105	-105	105	105	149	169	-133	89	111	124	124	58	77	
578	631	631	631	105	105	-105	105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77	
381	567	567	567	105	105	-105	105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77	
307	292	292	292	105	105	-105	105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77	
358	373	373	373	292	308	-308	409	375	375	-507	471	507	129	128	125	125	367	391	391	527	177	
11	256	240	240	105	105	-105	105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77	
12	262	219	219	105	105	-105	105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77	
13	41	105	105	105	105	-105	105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77	
1	186	170	170	170	292	308	-308	292	308	-308	292	308	266	278	278	278	266	291	291	246	246	177
151	250	250	250	105	105	-105	105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77	
28	133	133	133	105	105	-105	105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77	
112	88	105	105	105	105	-105	105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77	
44	105	105	105	105	105	-105	105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77	
164	156	156	156	156	156	-156	156	156	156	-156	156	156	156	156	-156	95	111	124	124	58	77	
78	56	56	56	56	56	-56	56	56	56	-56	56	56	56	56	-56	95	111	124	124	58	77	
59	7	7	7	7	7	-7	7	7	7	-7	7	7	7	7	-7	95	111	124	124	58	77	
*11	43	30	30	30	30	-30	30	30	30	-30	30	30	30	30	-30	95	111	124	124	58	77	
12	110	97	97	97	97	-97	97	97	97	-97	97	97	97	97	-97	95	111	124	124	58	77	
13	122	102	102	102	102	-102	102	102	102	-102	102	102	102	102	-102	95	111	124	124	58	77	
2	1153	1277	1277	1277	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77	
1104	1251	1251	1251	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
631	725	725	725	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
734	853	853	853	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
653	708	708	708	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
930	1189	1189	1189	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
235	250	250	250	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
535	640	640	640	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
333	334	334	334	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
294	303	303	303	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
202	211	211	211	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
11	311	277	277	277	105	105	-105	105	105	-105	105	105	105	-105	95	111	124	124	58	77		
12	197	188	188	188	105	105	-105	105	105	-105	105	105	105	-105	95	111	124	124	58	77		
3	368	308	308	308	105	105	-105	105	105	-105	105	105	105	-105	95	111	124	124	58	77		
273	245	245	245	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
15	15	15	15	15	15	-15	15	15	-15	15	15	15	15	-15	95	111	124	124	58	77		
208	226	226	226	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
101	110	110	110	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
48	32	32	32	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
294	302	302	302	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
28	7	7	7	7	7	-7	7	7	-7	7	7	7	105	105	-105	95	111	124	124	58	77	
11	164	132	132	132	105	105	-105	105	105	-105	105	105	105	-105	95	111	124	124	58	77		
12	164	132	132	132	105	105	-105	105	105	-105	105	105	105	-105	95	111	124	124	58	77		
13	53	54	54	54	54	-54	54	54	-54	54	54	54	54	-54	95	111	124	124	58	77		
4	972	1023	1023	1023	105	105	-105	105	105	-105	105	105	105	-105	95	111	124	124	58	77		
340	373	373	373	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
640	703	703	703	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
622	659	659	659	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
404	615	615	615	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
657	714	714	714	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
314	321	321	321	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
239	226	226	226	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
159	135	135	135	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
305	314	314	314	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
11	99	98	98	98	105	105	-105	105	105	-105	105	105	105	-105	95	111	124	124	58	77		
12	161	157	157	157	105	105	-105	105	105	-105	105	105	105	-105	95	111	124	124	58	77		
5	367	334	334	334	105	105	-105	105	105	-105	105	105	105	-105	95	111	124	124	58	77		
521	307	307	307	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
351	321	321	321	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
449	412	412	412	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
142	109	109	109	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
269	228	228	228	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
262	247	247	247	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
135	151	151	151	105	105	-105	105	105	-105	105	105	105	105	-105	95	111	124	124	58	77		
123	10																					

Table 6 (*cont.*)

Table 6 (*cont.*)

berg & Hordvik (1957). All the bond lengths are normal with an average carbon–carbon length of 1.516 Å and an average carbon–oxygen length of 1.434 Å. The average estimated standard deviations are 0.015 Å and 0.013 Å respectively. The largest deviation from the mean, for the C–C bonds is the C(2)–C(3) distance of 1.551 Å, a difference of 0.035 Å, about 2.3 standard deviations and, for the C–O bonds, the O(5)–C(5) distance of 1.461 Å, a difference of 0.027 Å, *i.e.* about 2.1 standard deviations.

Previous structure determinations of monosaccharides have been carried out on α -glucose (McDonald & Beevers, 1952), α -rhamnose (McGeachin & Beevers, 1957), β -arabinose (Furberg & Hordvik, 1957) and β -glucose (Ferrier, 1963). In all these cases the C(1)-

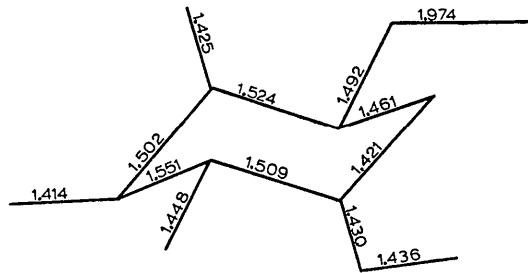


Fig. 2. Bond lengths (Å).

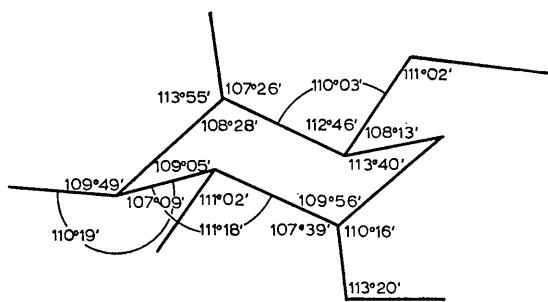


Fig. 3. Bond angles.

O(1) bond length has been found to be much shorter than other, similar bonds in the same molecule, e.g.

	Average C–O (Å)	C(1)–O(1) (Å)	e.s.d. (Å)
α -D-Glucose	1.40	1.32	—
α -Rhamnose	1.435	1.376	—
β -Arabinose	1.434	1.382	0.009
β -D-Glucose	1.444	1.404	0.010
<i>cf.</i> α -Methyl galactoside			
6-bromohydrin	1.434	1.430	0.014

The amount of shortening in each of the four cases is equivalent to about 10% double bond character (Pauling, 1960) and may be related to the differences of reactivity noted for this hydroxyl group. In the case of α -methyl D-galactoside 6-bromohydrin, in which the O(1) carries a methyl group and not a hydrogen atom, the bond length C(1)-O(1) is not shorter than the average but is almost exactly equal to the average. This would suggest that the short bond in the α position is an effect which is eliminated completely when the hydrogen atom is substituted.

Crystal structure and hydrogen bonding

The crystals have cleavage planes perpendicular to the z axis but are resistant to cleavage in any other direction. Examination of the crystal structure shows that the bromine atoms are arranged in the form of layers with the galactose molecules filling the space between the layers. No hydrogen bonds can pass through the layer of bromine atoms which thus becomes the cleavage plane, but the molecules are bound together by the hydrogen bonding system in the other directions. Details of the crystal structure are shown in Figs. 4 and 5.

Before the structure was solved the infrared spectrum was examined in order to gain some information about

the hydrogen bonds to be expected. The spectrum is shown in Fig. 6. This shows no peak at a wavelength of 2.8μ which would correspond to a free hydroxyl group. The peaks which do occur were thought to represent a double-weight peak at 3.03μ and a single-weight peak at 2.92μ . These may be related to the length of the $O-H \cdots O$ distance to be expected following Nakamoto, Margoshes & Rundle (1955). The values calculated by this process were:

$$\begin{aligned} 2 \text{ O}-\text{H}\cdots\text{O} \text{ distances of } 2.78 \text{ \AA} \\ 1 \text{ O}-\text{H}\cdots\text{O} \text{ distances of } 2.85 \end{aligned}$$

The values obtained in the final structure were:

	<i>l</i>	σ
O(4)-H---O(2)	2.733 Å	0.010 Å
O(3)-H---O(4)	2.778	0.011
O(2)-H---O(3)	2.824	0.011

and this agreement is quite good.

The hydrogen bonds form spiral arrangements around the screw axes of the unit cell at $x=\frac{1}{4}$ or $\frac{3}{4}$ and $z=\frac{1}{2}$, and extend through the crystal parallel to the *b* axis. Examination of the hydrogen atom positions showed that the directions of the bonds are as shown in the diagrams, *i.e.* the spiral around the $\frac{1}{4}, y, \frac{1}{2}$ screw axis is directed in a positive direction with respect to the *b* axis while the spiral around the $\frac{3}{4}, y, \frac{1}{2}$ screw axis has the opposite sense. The absolute sense of the direction of these spirals follows from the choice of the correct absolute configuration for the D-galactose molecule.

The dimensions of the unit cell in the *x* and *y* directions appear to be mainly dependent upon the size and packing of the bromine atoms. This produces some gaps or holes in the structure parallel to the *b* axis at $\frac{1}{2}, y, \frac{2}{5}$ and $0, y, \frac{3}{5}$. These channels are quite empty as they are not quite wide enough to contain water molecules and the material is thus anhydrous.

We wish to thank those named in the text for the use of their programs, the Director and staff of the University of Leeds Computing Laboratory for the calculations carried out on the Ferranti Pegasus computer and Sir Gordon Cox, K.B.E., F.R.S., for his initiation and encouragement of this work.

References

- BERGHUIS, J., HAANAPPEL, IJ. M., POTTERS, M., LOOPSTRA, B. O., MACGILLAVRY, C. H. & VEENENDAAL, A. L. (1955). *Acta Cryst.* **8**, 478.
- COX, E. G., GOODWIN, J. H. & WAGSTAFF, A. I. (1935). *J. Chem. Soc.* p. 978.
- CRUICKSHANK, D. W. J. (1949). *Acta Cryst.* **2**, 65.
- CRUICKSHANK, D. W. J. & ROLLETT, J. S. (1953). *Acta Cryst.* **6**, 705.
- CRUICKSHANK, D. W. J., PILLING, D. E., BUJOSA, A., LOVELL, F. M. & TRUTER, M. R. (1961). Symposium on Computer Methods. London: Pergamon Press.
- FERRIER, W. G. (1963). *Acta Cryst.* **16**, 1023.
- FURBERG, S. & HORDVIK, A. (1957). *Acta Chem. Scand.* **11**, 1594.

- MCDONALD, T. R. R. & BEEVERS, C. A. (1952). *Acta Cryst.* **5**, 654.
- MCGEACHIN, H. McD. & BEEVERS, C. A. (1957). *Acta Cryst.* **10**, 227.
- NAKAMOTO, K., MARGOSHES, M. & RUNDLE, R. E. (1955). *J. Amer. Chem. Soc.* **77**, 6480.
- PAULING, L. (1960). *The Nature of the Chemical Bond*. 3rd Edn. London: Oxford Univ. Press.
- ROBERTSON, J. H. (1960). *J. Sci. Instrum.* **37**, 41.
- THOMAS, L. H. & UMEDA, M. (1957). *J. Chem. Phys.* **26**, 293.
- VALENTIN, F. (1952). *Coll. Trav. Chim. Tschecoslovaquie*, **4**, 364.

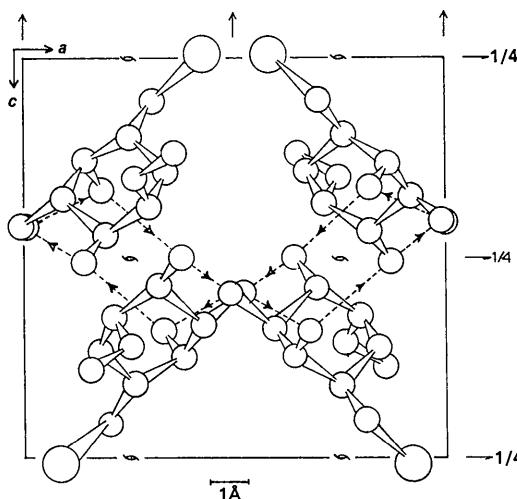


Fig. 4. Diagram of the unit cell projected down the *b* axis. The channels through the structure at $\frac{1}{4}, y, \frac{2}{5}$ and $0, y, \frac{3}{5}$ are prominent. Hydrogen bonds are shown as dashed lines.

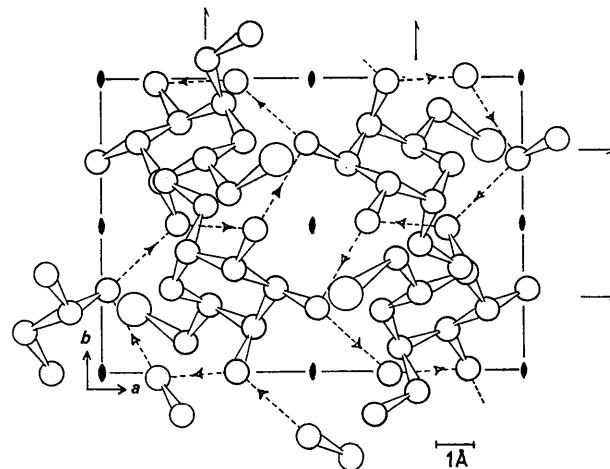


Fig. 5. Diagram of the unit cell projected down the *c* axis.

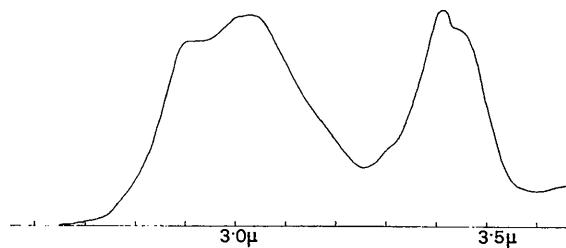


Fig. 6. Infrared spectrum.