

## Crystal Structure of the *p*-Bromophenylhydrazone of Mannose

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The crystal structure of the *p*-bromophenylhydrazone of mannose has been determined by X-ray crystallographic methods and refined to an *R* index of 0.078. Three-dimensional film data were used. The compound is a true hydrazone, the mannose having an acyclic structure. The carbon atoms of the sugar form a nearly planar zigzag chain. Delocalization effects in the hydrazine part of the molecule are indicated.

Chemical evidence indicates that the phenylhydrazones of the sugars may occur in cyclic or acyclic forms.<sup>1</sup> The crystal structure of the *p*-bromophenylhydrazones of a number of sugars have been determined in this laboratory. Those of arabinose<sup>2</sup> and glucose<sup>3</sup> were found to be hydrazides with the sugar in the pyranose chair form, whereas that of ribose<sup>4</sup> is a true hydrazone, the sugar occurring in the open-chain form. In this paper the crystal structure of the mannose derivative is reported.

The formazan reaction of the phenylhydrazone of mannose would seem to indicate that it has an acyclic structure,<sup>1</sup> but recent infrared spectroscopic evidence suggests a cyclic structure for this compound in the solid state.<sup>8</sup> We have found no information on the structure of the *p*-bromophenylhydrazone of mannose in the literature.

Crystals of *p*-bromophenylhydrazone of mannose were grown from pyridine-water mixtures, and 1182 reflections recorded by taking Weissenberg photographs about the *b* axis. The structure was solved from the Patterson projection and stereochemical considerations, and refined by least squares methods to a final *R* value of 0.078. The positions of the atoms are listed in Table 1 and the corresponding bond lengths and angles in Table 2 and Fig. 1. Details of the structure analysis are given below. The estimated standard deviations are 0.02–0.03 Å in the bond lengths and 1–2° in the angles (Table 2).

It is evident from the atomic positions that the sugar has an acyclic structure and that the compound is a true hydrazone,

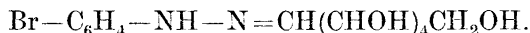


Table 1. Final atomic coordinates (in fractions of cell edges) and temperature factors. Estimated standard deviations in parentheses.

Atom	<i>x</i>	<i>y</i>	<i>z</i>	<i>B</i>
Br	0.0000	0.0000	0.0000	<sup>a</sup>
O(2)	0.2902 (21)	0.5780 (21)	0.4534 (7)	2.00 (16)
O(3)	0.9903 (22)	0.9014 (21)	0.6601 (7)	2.15 (17)
O(4)	0.9706 (21)	0.3758 (21)	0.6452 (7)	2.12 (17)
O(5)	0.5782 (22)	0.8907 (22)	0.7971 (8)	2.16 (17)
O(6)	0.5765 (25)	0.3700 (24)	0.7829 (8)	2.81 (20)
N(2)	0.8405 (30)	0.1031 (29)	0.3594 (10)	2.63 (23)
N(1)	0.8458 (25)	0.9034 (26)	0.4020 (9)	2.06 (20)
C(7)	0.2634 (35)	0.0387 (36)	0.1142 (12)	2.82 (27)
C(8)	0.4795 (38)	0.2636 (38)	0.1484 (13)	3.23 (30)
C(9)	0.6581 (52)	0.2772 (52)	0.2334 (18)	3.54 (38)
C(10)	0.6449 (31)	0.0750 (31)	0.2780 (11)	2.20 (23)
C(11)	0.4263 (36)	0.8492 (36)	0.2375 (12)	2.88 (28)
C(12)	0.2406 (37)	0.8329 (37)	0.1569 (13)	3.08 (29)
C(1)	0.0800 (28)	0.9178 (30)	0.4517 (10)	1.85 (21)
C(2)	0.0985 (27)	0.7049 (28)	0.5006 (9)	1.67 (20)
C(3)	0.2026 (25)	0.8006 (28)	0.6112 (9)	1.46 (20)
C(4)	0.2370 (28)	0.5857 (29)	0.6631 (10)	1.79 (21)
C(5)	0.3175 (32)	0.6787 (34)	0.7777 (11)	1.67 (25)
C(6)	0.3606 (29)	0.4729 (30)	0.8291 (10)	1.98 (22)
<sup>a</sup> β <sub>11</sub> = 0.04515 (75)		β <sub>22</sub> = 0.03051 (79)		
β <sub>33</sub> = 0.00286 (6)		β <sub>12</sub> = 0.03486 (103)		
β <sub>13</sub> = -0.00496 (31)		β <sub>23</sub> = 0.00443 (29)		
in exp -(β <sub>11</sub> h <sup>2</sup> + β <sub>22</sub> k <sup>2</sup> + β <sub>33</sub> l <sup>2</sup> + β <sub>12</sub> hk + β <sub>23</sub> kl + β <sub>13</sub> hl)				

Table 2. Bond lengths (Å units) and bond angles (°) with estimated standard deviations.

Bond lengths		Bond angles	
Br - C(7)	1.894 (17)	Br - C(7) - C(8)	119.8 (1.4)
C(7) - C(8)	1.372 (25)	Br - C(7) - C(12)	118.1 (1.3)
C(8) - C(9)	1.384 (29)	C(7) - C(8) - C(9)	116.0 (1.9)
C(9) - C(10)	1.380 (31)	C(8) - C(9) - C(10)	123.7 (2.2)
C(10) - C(11)	1.395 (21)	C(9) - C(10) - C(11)	117.3 (1.6)
C(11) - C(12)	1.352 (24)	C(10) - C(11) - C(12)	120.3 (1.7)
C(7) - C(12)	1.376 (27)	C(7) - C(12) - C(11)	120.5 (1.7)
C(10) - N(2)	1.373 (19)	C(12) - C(7) - C(8)	122.0 (1.7)
N(2) - N(1)	1.370 (20)	C(9) - C(10) - N(2)	119.7 (1.6)
N(1) - C(1)	1.254 (18)	C(11) - C(10) - N(2)	123.0 (1.6)
C(1) - C(2)	1.504 (22)	C(10) - N(2) - N(1)	121.4 (1.3)
C(2) - C(3)	1.513 (16)	N(2) - N(1) - C(1)	117.8 (1.3)
C(3) - C(4)	1.561 (21)	N(1) - C(1) - C(2)	120.2 (1.3)
C(4) - C(5)	1.545 (19)	C(1) - C(2) - C(3)	110.3 (1.2)
C(5) - C(6)	1.525 (24)	C(2) - C(3) - C(4)	111.5 (1.2)
C(2) - O(2)	1.422 (16)	C(3) - C(4) - C(5)	111.4 (1.2)
C(3) - O(3)	1.428 (16)	C(4) - C(5) - C(6)	112.1 (1.4)
C(4) - O(4)	1.438 (16)	C(1) - C(2) - O(2)	111.8 (1.0)
C(5) - O(5)	1.428 (19)	O(2) - C(2) - C(3)	109.7 (1.1)
C(6) - O(6)	1.433 (18)	C(2) - C(3) - O(3)	109.3 (1.0)
		O(3) - C(3) - C(4)	108.0 (1.0)
		C(3) - C(4) - O(4)	111.3 (1.0)
		O(4) - C(4) - C(5)	109.1 (1.1)
		C(4) - C(5) - O(5)	109.9 (1.1)
		O(5) - C(5) - C(6)	110.1 (1.2)
		C(5) - C(6) - O(6)	110.5 (1.1)

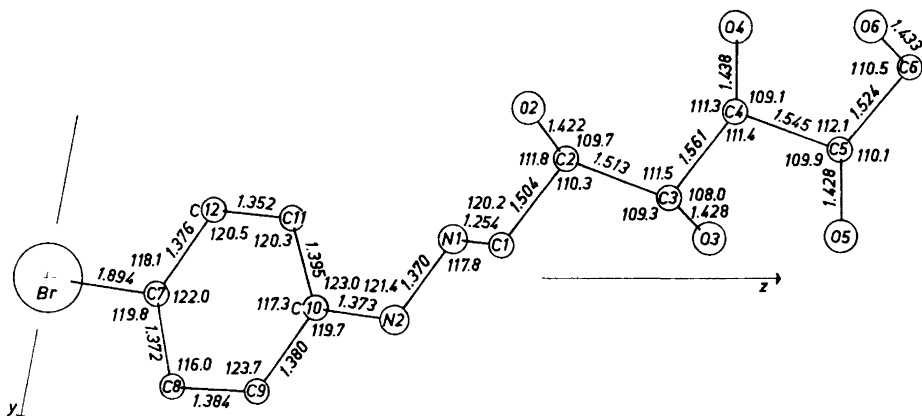


Fig. 1. The molecular structure of mannose-*p*-bromophenylhydrazone (seen along *a*). Bond lengths in Å units.

The carbon atoms of the mannose form a nearly planar zigzag chain, the largest deviation from the best least squares plane through these atoms being only 0.04 Å. The mean bond lengths and bond angles in this part of the molecule are 1.530 Å (C—C), 1.430 Å (C—O), 111.3° (C—C—C), and 109.9° (C—C—O). The corresponding values reported for galactitol are 1.520 Å, 1.432 Å, 111.6°, and 110.0°, respectively.<sup>9</sup> The conformation of mannose is the same as in the polymorphs of mannitol.<sup>5</sup> The terminal oxygen atom O(6) is not a linear extension of the zigzag chain.

Table 3. Short intermolecular distances (Å units) and their angles (°) with adjacent bonds.

Distances		Angles	
O <sub>b</sub> (2)...N(2)	2.940 (18)	C(10)—N(2)...O <sub>b</sub> (2)	124.4 (1.1)
O(2)...N <sub>a</sub> (1)	2.934 (16)	N(1)—N(2)...O <sub>b</sub> (2)	114.2 (0.9)
O(3)...O <sub>b</sub> (4)	2.770 (17)	C <sub>a</sub> (1)—N <sub>a</sub> (1)...O(2)	117.4 (1.0)
O <sub>a</sub> (3)...O(5)	2.745 (15)	N <sub>a</sub> (2)—N <sub>a</sub> (1)...O(2)	119.2 (0.9)
O <sub>a</sub> (4)...O(6)	2.692 (15)	C <sub>b</sub> (2)—O <sub>b</sub> (2)...N(2)	98.3 (0.8)
O(5)...O <sub>b</sub> (6)	2.769 (19)	C(2)—O(2)...N <sub>a</sub> (1)	115.4 (0.8)
		N(2)...O <sub>b</sub> (2)...N <sub>ab</sub> (1)	137.2 (0.5)
		C(3)—O(3)...O <sub>b</sub> (4)	120.6 (0.8)
		C <sub>a</sub> (3)—O <sub>a</sub> (3)...O(5)	150.7 (0.9)
		C <sub>b</sub> (4)—O <sub>b</sub> (4)...O(3)	120.4 (0.8)
		C <sub>a</sub> (4)—O <sub>a</sub> (4)...O(6)	117.0 (0.8)
		C(5)—O(5)...O <sub>a</sub> (3)	116.7 (0.9)
		C(5)—O(5)...O <sub>b</sub> (6)	123.0 (1.0)
		C(6)—O(6)...O <sub>a</sub> (4)	148.9 (1.0)
		C <sub>b</sub> (6)—O <sub>b</sub> (6)...O(5)	120.4 (0.9)
		O <sub>a</sub> (3)...O <sub>ab</sub> (4)...O <sub>b</sub> (6)	92.0 (0.5)
		O <sub>ab</sub> (4)...O <sub>b</sub> (6)...O(5)	89.0 (0.5)
		O <sub>b</sub> (6)...O(5)...O <sub>a</sub> (3)	91.9 (0.5)
		O(5)...O <sub>a</sub> (3)...O <sub>ab</sub> (4)	87.9 (0.4)

A<sub>a</sub> is an atom in (*x*+1, *y*, *z*), A<sub>b</sub> in (*x*, *y*+1, *z*) and A<sub>ab</sub> in (*x*+1, *y*+1, *z*).

Table 4. Observed ( $F_o$ ) and calculated ( $F_c$ ) structure factors.

h	k	l	F <sub>o</sub>	F <sub>c</sub>	h	k	l	F <sub>o</sub>	F <sub>c</sub>	h	k	l	F <sub>o</sub>	F <sub>c</sub>	h	k	l	F <sub>o</sub>	F <sub>c</sub>
5	0	7	37	35	1	0	16	77	74	3	1	-4	130	129	0	1	-8	206	206
5	0	6	40	49	1	0	15	106	109	3	1	-5	114	109	0	1	-9	165	154
5	0	5	71	78	1	0	14	90	82	3	1	-6	105	96	0	1	-10	201	209
5	0	4	89	94	1	0	13	102	101	3	1	-7	93	94	0	1	-11	168	171
5	0	3	88	59	1	0	12	135	132	3	1	-8	104	94	0	1	-12	108	104
5	0	2	54	55	1	0	11	184	168	3	1	-9	89	73	0	1	-13	104	94
5	0	1	72	75	1	0	10	113	106	3	1	-10	103	93	0	1	-14	84	73
5	0	0	110	129	1	0	9	158	147	3	1	-11	55	53	0	1	-15	127	129
5	0	-1	94	100	1	0	8	279	265	3	1	-12	61	53	0	1	-16	84	93
5	0	-2	43	38	1	0	7	107	107	3	1	-13	29	27	0	1	-17	60	74
5	0	-3	85	82	1	0	6	473	482	3	1	-14	74	78	-1	1	16	32	49
5	0	-4	65	63	1	0	5	91	87	3	1	-15	28	23	-1	1	15	94	90
5	0	-5	50	51	1	0	4	274	272	2	1	14	72	75	-1	1	14	47	47
5	0	-6	38	31	1	0	3	491	486	2	1	13	79	74	-1	1	13	102	94
5	0	-7	35	37	1	0	2	402	396	2	1	12	95	91	-1	1	12	128	120
4	0	11	52	71	1	0	1	117	144	2	1	11	143	139	-1	1	11	157	150
4	0	10	58	58	1	0	0	303	319	2	1	10	117	127	-1	1	10	176	159
4	0	9	87	88	1	0	-1	86	88	2	1	9	129	126	-1	1	9	166	154
4	0	8	74	71	1	0	-2	621	722	2	1	8	91	72	-1	1	8	128	113
4	0	7	75	77	1	0	-3	532	594	2	1	7	89	80	-1	1	7	261	296
4	0	6	131	142	1	0	-4	125	125	2	1	6	351	351	-1	1	6	203	194
4	0	5	95	95	1	0	-5	254	254	2	1	5	131	128	-1	1	5	149	148
4	0	4	66	66	1	0	-6	192	188	2	1	4	182	172	-1	1	4	107	110
4	0	3	212	223	1	0	-7	226	233	2	1	3	162	168	-1	1	3	352	375
4	0	2	131	123	1	0	-8	167	157	2	1	2	190	197	-1	1	2	538	668
4	0	1	98	104	1	0	-9	195	189	2	1	1	219	209	-1	1	1	384	448
4	0	0	71	80	1	0	-10	148	134	2	1	0	213	214	-1	1	0	112	124
4	0	-1	45	45	1	0	-11	169	166	2	1	-1	88	96	-1	1	-1	231	251
4	0	-2	73	86	1	0	-12	174	168	2	1	-2	301	298	-1	1	-2	177	187
4	0	-3	221	237	1	0	-13	61	54	2	1	-3	214	203	-1	1	-3	259	318
4	0	-4	108	105	1	0	-14	109	106	2	1	-4	152	163	-1	1	-4	201	203
4	0	-5	78	78	1	0	-15	56	60	2	1	-5	161	150	-1	1	-5	342	350
4	0	-6	89	85	1	0	-16	51	81	2	1	-6	121	123	-1	1	-6	274	269
4	0	-7	80	83	1	0	-17	58	73	2	1	-7	201	198	-1	1	-7	253	242
4	0	-8	128	125	1	0	-18	101	106	2	1	-8	101	106	-1	1	-8	145	134
4	0	-9	27	32	0	0	17	57	62	2	1	-9	144	135	-1	1	-9	201	198
4	0	-10	50	42	0	0	16	36	33	2	1	-10	74	68	-1	1	-10	168	152
4	0	-11	68	60	0	0	15	71	70	2	1	-11	113	115	-1	1	-11	173	165
4	0	-12	71	68	0	0	14	120	124	2	1	-12	115	107	-1	1	-12	146	139
3	0	13	79	80	0	0	13	119	112	2	1	-13	97	88	-1	1	-13	111	109
3	0	12	85	90	0	0	12	118	111	2	1	-14	86	71	-1	1	-14	80	79
3	0	11	115	115	0	0	11	154	142	2	1	-15	86	76	-1	1	-15	109	106
3	0	10	51	38	0	0	10	105	106	2	1	-16	88	71	-1	1	-16	108	108
3	0	9	161	162	0	0	9	272	263	2	1	-17	111	101	-1	1	-17	60	84
3	0	8	147	149	0	0	8	300	275	1	1	15	74	77	-1	1	-17	60	84
3	0	7	106	102	0	0	7	131	111	1	1	14	81	77	-2	1	16	25	33
3	0	6	96	87	0	0	6	423	426	1	1	13	115	102	-2	1	15	76	73
3	0	5	115	110	0	0	5	457	451	1	1	12	113	107	-2	1	14	72	64
3	0	4	207	200	0	0	4	228	226	1	1	11	141	128	-2	1	13	89	81
3	0	3	211	211	0	0	3	252	231	1	1	10	58	54	-2	1	12	113	102
3	0	2	162	160	0	0	2	77	66	1	1	9	199	200	-2	1	11	117	101
3	0	1	96	97	0	0	1	90	64	1	1	8	225	205	-2	1	10	99	89
3	0	0	206	217	0	0	0	20	26	1	1	7	135	123	-2	1	9	239	229
3	0	-1	211	224	5	1	3	35	24	1	1	6	219	206	-2	1	8	206	85
3	0	-2	74	67	5	1	2	74	87	1	1	5	265	273	-2	1	8	138	136
3	0	-3	88	86	5	1	1	34	21	1	1	4	200	192	-2	1	6	81	69
3	0	-4	159	173	5	1	0	77	73	1	1	3	295	295	-2	1	5	356	379
3	0	-5	176	182	5	1	-1	74	34	1	1	2	111	117	-2	1	4	371	430
3	0	-6	224	216	5	1	-2	40	5	1	1	1	153	163	-2	1	3	159	159
3	0	-7	111	102	5	1	-3	80	73	1	1	0	516	598	-2	1	2	81	76
3	0	-8	146	139	5	1	-4	35	50	1	1	-1	180	192	-2	1	1	193	186
3	0	-9	64	72	5	1	-5	27	43	1	1	-2	302	328	-2	1	0	246	260
3	0	-10	117	109	4	1	9	64	65	1	1	-3	124	144	-2	1	-1	204	230
3	0	-11	54	60	4	1	8	48	50	1	1	-4	246	269	-2	1	-2	219	232
3	0	-12	32	31	4	1	7	70	75	1	1	-5	223	233	-2	1	-3	250	293
3	0	-13	47	48	4	1	6	91	76	1	1	-6	219	219	-2	1	-4	310	312
3	0	-14	43	42	4	1	5	79	73	1	1	-7	186	189	-2	1	-5	150	138
2	0	15	71	71	4	1	4	95	92	1	1	-8	120	136	-2	1	-6	164	162
2	0	14	98	104	4	1	3	35	20	-1	1	-9	191	191	-2	1	-7	214	210
2	0	13	79	71	4	1	2	105	92	1	1	-10	122	125	-2	1	-8	175	162
2	0	12	116	115	4	1	1	105	96	1	1	-11	94	93	-2	1	-9	234	220
2	0	11	153	145	4	1	0	89	82	1	1	-12	140	131	-2	1	-10	105	97
2	0	10	141	140	4	1	-1	99	103	1	1	-13	190	194	-2	1	-11	131	117
2	0	9	147	143	4	1	-2	80	69	1	1	-14	92	80	-2	1	-12	122	118
2	0	8	151	139	4	1	-3	59	49	1	1	-15	85	46	-2	1	-13	160	168
2	0	7	96	91	4	1	-4	106	83	1	1	-16	25	34	-2	1	-14	72	70
2	0	6	292	289	4	1	-5	54	35	1	1	-17	54	50	-2	1	-15	107	114
2	0	5	196	193	4	1	-6	103	93	0	1	16	54	61	-2	1	-16	38	48
2	0	4	212	194	4	1	-7	53	36	0	1	15	83	86	-3	1	15	37	47
2	0	3	287	283	4	1	-8	44	53	0	1	14	93	99	-3	1	14	68	64
2	0	2	132	129	4	1	-9	49	43	0	1	13	59	83	-3	1	13	59	48
2	0	1	266	285	4	1	-10	76	68	0	1	12	113	101	-3	1	12	92	85
2	0	0	200	222	4	1	-11	34	29	0	1	11	109	104	-3	1	11	86	69
2	0	-1	208	216	3	1	12	84	83	0	1	10	175	160	-3	1	10	59	55
2	0	-2	190	211	3	1	11	68	62	0	1	9	153	144	-3	1	9	139	131
2	0	-3	199	203	3	1	10	83	78	0	1	8	241	214	-3	1	8	114	104
2	0	-4	286	296	3	1	9												

Table 4. Continued.

h	k	l	F <sub>o</sub>	F <sub>c</sub>	h	k	l	F <sub>o</sub>	F <sub>c</sub>	h	k	l	F <sub>o</sub>	F <sub>c</sub>	h	k	l	F <sub>o</sub>	F <sub>c</sub>
-3	1	-4	187	185	2	2	4	139	141	-1	2	-11	103	83	-5	2	-9	59	81
-3	1	-7	136	136	2	2	3	133	143	-1	2	-12	155	144	-6	2	3	52	61
-3	1	-8	104	93	2	2	2	112	114	-1	2	-13	163	158	-8	2	2	40	50
-3	1	-9	155	134	2	2	1	68	70	-1	2	-14	96	96	-6	2	-1	39	52
-3	1	-10	191	190	2	2	0	258	243	-1	2	-15	152	157	4	3	4	20	13
-3	1	-11	106	101	2	2	-1	300	209	-1	2	-17	72	97	4	3	3	30	33
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-3	3	5	137	127	-2	4	-4	81	91	-1	4	-16	59	80	-5	4	-6	70	81
-3	3	4	95	99	-2	4</													

The atoms N(1), N(2), C(1), and C(2) are coplanar (to within 0.002 Å), as is to be expected for a true hydrazone structure. This plane forms an angle of 21.3° with the benzene ring plane and of 124.0° with the plane of the sugar carbon atoms. The bond N(1)—C(1) is *trans* to the bisecting line of the angle O(2)C(2)C(3). Essentially the same relative orientation of the sugar and hydrazone part of the molecule with respect to the bond C(1)—C(2) was found also in the ribose derivative.<sup>4</sup> This conformational feature may well be preferred for all hydrazones as it implies *cis* relation between N(1) and the hydrogen atom at C(2). The N(1)—C(1) bond length of 1.25 Å is close to a normal double bond, whereas the bonds N(1)—N(2) (1.37 Å) and N(2)—C(10) (1.37 Å) are significantly shorter than normal single bonds. The angles at N(1) and N(2) are close to 120°, and it is possible that both these atoms are nearly  $sp^2$  hybridized. At least some of the shortening may be explained on this basis, but additional delocalization effects probably also occur.

The azimuthal angle at the N—N bond is 157°, not far from the value of 165° found in the ribose derivative.

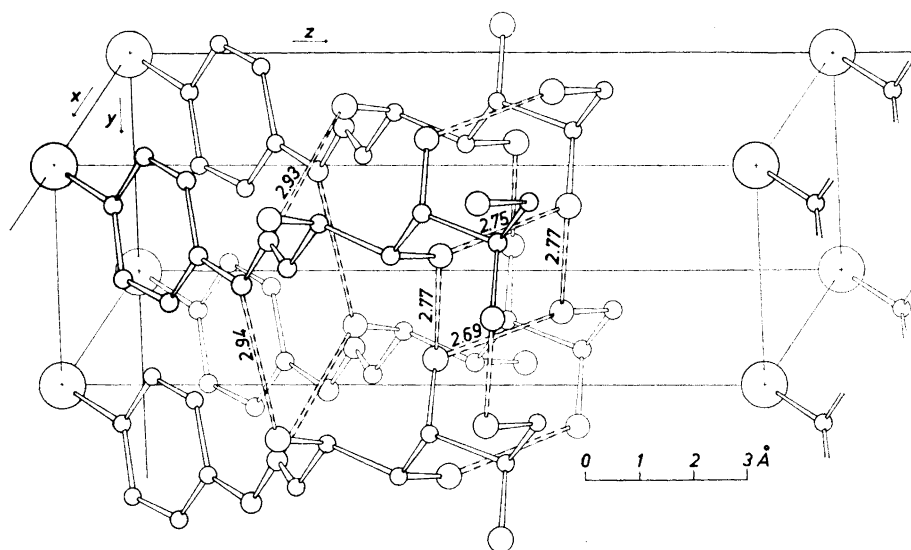


Fig. 2. Arrangement of the molecules in the crystal. Four neighbouring molecules are projected into the plane (111). Broken lines indicate hydrogen bonds.

The arrangement of the molecules in the crystal is shown in Fig. 2. The hydrogen atoms were not located in the present analysis, but distances and directions strongly suggest that all oxygen and nitrogen atoms in the structure are involved in hydrogen bonds. There are six intermolecular O...O and O...N distances shorter than 3.0 Å, corresponding to six different hydrogen bonds, one for each hydrogen atom attached to oxygen or nitrogen atoms. The atom

O(2) is linked to nitrogen atoms in two different neighbouring molecules by bonds N(2)—H...O(2) (2.94 Å) and O(2)—H...N(1) (2.93 Å). The remaining four oxygen atoms are all connected to two different neighbouring molecules by O—H...O bond of lengths 2.69—2.77 Å. These bonds form a closed roughly planar square ring, the angle between the two hydrogen bonds being close to 90° at each oxygen atom (range 88°—92°).

As there is only one molecule per unit cell, all molecules have the same orientation. They are stacked in layers parallel to the *xy* plane, with their "axes" roughly perpendicular to this plane, which is also the predominant crystal face. There are no hydrogen bonds between these layers.

#### EXPERIMENTAL. STRUCTURE ANALYSIS

The compound was prepared by mixing aqueous solutions of D-mannose and *p*-bromophenylhydrazine hydrochloride at room temperature. It is almost insoluble in most of the usual solvents, except boiling pyridine. Crystals were grown by slow cooling of a saturated pyridine-water solution. Flat needles, elongated along *a*, and with (001) as the predominant face, were obtained.

X-Ray diagrams showed that the crystals were triclinic. The unit cell dimensions were derived from 15 well resolved low order lines recorded on a Guinier camera calibrated against KCl. The following values were found:  $a=4.750$  (0.006) Å,  $b=5.685$  (0.007) Å,  $c=13.577$  (0.021) Å,  $\alpha=100.43^\circ$  (0.11),  $\beta=92.13^\circ$  (0.09),  $\gamma=107.09^\circ$  (0.12). The uncertainties indicated are estimated standard deviations. As the compound is optically active, the space group is *P*1. There is only one molecule in the unit cell. The density was found to be 1.70 g/cm<sup>3</sup>. The calculated value is 1.69 g/cm<sup>3</sup>.

A crystal of dimensions 0.26 × 0.09 × 0.04 mm was used to record layers *h*0*l*—*h*4*l* by Weissenberg integrating techniques, employing CuK $\alpha$  radiation ( $\lambda=1.542$  Å). The intensities were measured on a Hilger & Watts photometer. The weakest ones were estimated visually. In all 1182 non-equivalent reflections were recorded with measurable intensities. They were corrected for absorption effects. The non-observed reflections were not included in the analysis.

The structure was solved by the heavy atom procedure, deriving first the (*h*0*l*) projection. A program based on the "minimum residual method"<sup>16</sup> proved useful in locating the atoms. The structure was refined by block diagonal least squares calculations. The weighting scheme used was  $w=10.0$  for  $F \leq 6.0$  and of the form  $w=17.1 F^{-0.30}$  for  $F > 6.0$ . The scale factors for the different layers were adjusted by means of the calculated values of the structure factors. Anisotropic temperature factor was used only for the bromine atom, the vibrations of the light atoms being treated as isotropic. The temperature factors are given in Table 1. The hydrogen atoms could not be unequivocally located from difference maps. The positions of 13 of them were postulated on the basis of stereochemical considerations, but not refined. A temperature factor  $B=2.5$  Å<sup>2</sup> was applied. The contribution from the remaining four hydrogen atoms [those attached to O(3), O(4), O(5), and O(6)] was neglected. The final value of the *R* index is 0.078. In Table 4 observed and calculated structure factors are compared.

The atomic form factors of Hanson *et al.*<sup>7</sup> were used. The programs applied were written, or revised, for CDC 3300 by T. Dahl, F. Gram, P. Groth, B. Klewe and Chr. Rømming.



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