

# Crystal and Molecular Structure of 3-Hydroxybiuret Potassium Salt (2:1), $\text{KH}(\text{C}_2\text{H}_4\text{N}_3\text{O}_3)_2$

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The structure of the 3-hydroxybiuret potassium salt (2:1) ( $\text{C}_2\text{H}_4\text{KN}_3\text{O}_3$ ,  $\text{C}_2\text{H}_5\text{N}_3\text{O}_3$ ) has been determined, using three-dimensional diffractometer-collected X-ray data. The salt crystallizes in the space group  $P2_1/c$  with 4 formula units  $\text{C}_4\text{H}_9\text{KN}_6\text{O}_6$  in the unit cell with the dimensions  $a = 3.868(2)$ ,  $b = 20.38(1)$ ,  $c = 12.400(7)$  Å,  $\beta = 90.38(7)^\circ$ . The structure was solved by the heavy-atom method and refined by full-matrix least-squares calculations to a conventional  $R$  of 0.032.

The crystals are built up of  $\text{K}^+$  ions and  $\text{H}(\text{C}_2\text{H}_4\text{N}_3\text{O}_3)_2^-$  ions. The two 3-hydroxybiuret residues are joined by a very short hydrogen bond (2.442(3) Å) between the hydroxyl oxygen atoms. In addition the anions are connected by several  $\text{NH}\cdots\text{O}$  hydrogen bonds. Each potassium ion is coordinated to eight oxygen atoms. The  $\text{K}\cdots\text{O}$  distances lie in the range 2.663–3.026 Å.

The structure determination of 3-hydroxybiuret,  $\text{H}_2\text{N}-(\text{C}=\text{O})-\text{N}(\text{OH})-(\text{C}=\text{O})-\text{NH}_2$ , was undertaken as part of an X-ray study of organic hydroxylamine derivatives, which are inhibitors of DNA synthesis in several cell systems. 3-Hydroxybiuret (3-HB) and 1-hydroxybiuret (1-HB) were by Gale *et al.*<sup>1</sup> shown to inhibit DNA synthesis in an *in vitro* Ehrlich ascites tumor test system. The 3-HB used in these experiments was prepared by the method described by Exner,<sup>2</sup> but following the same method the present author obtained a compound with a different melting point (138–139 °C decomp.; Exner: 133 °C decomp.). The elementary analysis suggested the compound to be a 3-HB potassium salt (2:1), and this assumption was verified by the present structure determination. The CA name of the compound for the present collective-index period (1972–76) is 2-

hydroxyimidodicarbonic diamide potassium salt (2:1). By treatment of the potassium salt with a strongly acidic ion-exchange resin a compound was obtained which is presumed to be 3-HB, but the melting point of 158–160 °C decomp. does still not agree with that reported by Exner. X-Ray structure determination of this compound and of 1-HB are in progress.

## EXPERIMENTAL

The potassium acid salt of 3-HB was synthesized from hydroxyurea and potassium cyanate following the method described by Exner<sup>2</sup> for preparation of 3-HB. The yield was 48 % of a crystalline crude product. Recrystallization by precipitation with absolute ethanol from a saturated aqueous solution of the crude product yielded thin, colourless needles, m.p. 138–139 °C (decomp.). (Found: C 17.35; H 3.34; N 30.16. Calc. for  $\text{C}_4\text{H}_9\text{KN}_6\text{O}_6$ : C 17.40; H 3.28; N 30.20).

Table 1. Crystal data for the 3-HB potassium salt (2:1).

Mol. formula	$\text{C}_4\text{H}_9\text{KN}_6\text{O}_6$
Mol. weight	276.3
Melting point	138–139 °C (decomp.)
Space group	$P2_1/c$
$a$	3.868(2) Å
$b$	20.38(1) Å
$c$	12.400(7) Å
$\beta$	90.38(7)°
$V$	977.5 Å <sup>3</sup>
$Z$	4
$D_x$	1.87 g/cm <sup>3</sup>
$D_m$	1.85 g/cm <sup>3</sup>
$\mu_{\text{MoK}\alpha}$	5.78 cm <sup>-1</sup>
Crystal size	0.10 × 0.12 × 0.50 mm <sup>3</sup>
Rot. axis	$a$

Crystals suitable for X-ray work were obtained by diffusion at room temperature of absolute ethanol into a saturated solution of the salt in aqueous ethanol. Some crystal data of the compound are given in Table 1. The density was measured by flotation in a mixture of carbon tetrachloride and methyl iodide. The melting points were determined with a hot stage microscope (Mikroskop Heiztisch Ernst Leitz G.m.b.H., Wetzlar). The microanalysis was performed by Preben Hansen, Microanalytical Laboratory, University of Copenhagen.

The lattice parameters were calculated from a series of diffractometer-measured  $\theta$ -values. The intensity data were collected with a NO-NIUS 3-circle automatic diffractometer by the moving crystal-stationary detector technique, using graphite monochromatized  $MoK\alpha$ -radiation ( $\lambda = 0.71069 \text{ \AA}$ ). The scan speed was  $0.6^\circ/\text{min}$  and each reflexion was scanned over a range of  $1.2^\circ$ . The background was measured on each side of the reflexion for half the scanning time. The background count was in the data reduction set equal to twice the lowest count obtained in order to avoid weak satellite peaks, observed on one side of the strong reflexions. Intensities of reflexions were measured in the range  $2.5^\circ < \theta < 25^\circ$ , and the intensity of a reference reflexion was measured for every 25 reflexions. Thus 1663 independent reflexions were measured. A reflexion was considered unobserved and was omitted when the intensity observed was less than 2.5 times its estimated standard deviation. The number of observed reflexions was reduced to 1446. These data were corrected for Lorentz and polarization effects, but no corrections for absorption or extinction were made.

## STRUCTURE DETERMINATION

The structure was solved from the three-dimensional Patterson map by the heavy-atom method. The electron density map, based on the position of the potassium ion only, revealed the positions of all 16 non-hydrogen atoms of the two 3-HB residues in the asymmetric unit. The conventional  $R$ -value was 0.31 at this stage. Three cycles of full-matrix least-squares refinement, in which positional as well as individual atomic, isotropic thermal parameters were varied, reduced the  $R$ -value to 0.074.

The difference Fourier map, calculated at this stage of the refinement, revealed the 8 hydrogen atoms of the  $NH_2$ -groups unambiguously (cf. Fig. 1). The 9th hydrogen atom is situated between the hydroxyl oxygen atoms  $O(3)$  and  $O(3')$ , apparently either symmetrically or statistically disordered.

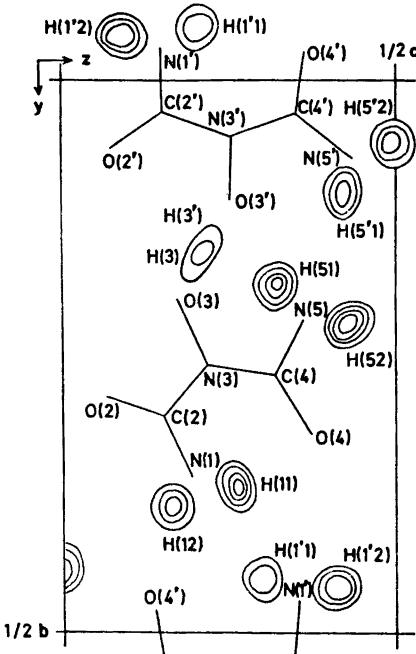


Fig. 1. Projection along the  $a$ -axis of sections of the difference Fourier map. Contours are at intervals of about  $0.1 \text{ e } \text{\AA}^{-3}$ .

In the following refinement this hydrogen atom was treated as 2 half hydrogen atoms,  $H(3)$  and  $H(3')$ , i.e. each with a population parameter of 0.5 and situated in either end of the elongated peak, about  $0.8 \text{ \AA}$  apart.

Two cycles of anisotropic least-squares refinement with all H-atom parameters fixed, reduced the  $R$ -value to 0.042. In the remaining two cycles of refinement the positional parameters of the H-atoms were also varied, but the isotropic thermal parameters of  $2.0 \text{ \AA}^2$  were still fixed. The final  $R$ -value was 0.032. Refinement of the population parameters of  $H(3)$  and  $H(3')$  was also carried out, but the shifts were not significant, and these parameters were therefore fixed on 0.5.

In another attempt of refinement the hydrogen atom  $H(3)$  was included with a position midway between the oxygen atoms  $O(3)$  and  $O(3')$  and allowed to vibrate isotropically. Only these four parameters of  $H(3)$  and the scale factor were refined, and at convergence ( $R = 0.031$ ) the hydrogen atom had moved toward the oxygen atom  $O(3')$ . The dimensions corre-

**Table 2.** Observed and calculated structure factors. The columns are  $l$ ,  $10 \times F_o$  and  $10 \times F_c$ .

0.0,L	8	318	-317	3	67	-60	-10	290	283	13	102	-109	1.9,L						
2	412	-448	10	81	76	5	126	-131	-12	209	205	-14	107	-114					
4	294	-286	11	208	-206	6	110	-109	-14	100	91	-14	113	-112					
6	439	457	12	132	-133	7	74	-79	-14	145	140		1.5,L	-1.4,L					
8	128	139	13	30	24	6	192	-189		35	33		1.5,L	-1.4,L					
10	592	-598	14	116	-98	9	96	-36	-169	10	170		1.5,L	-1.4,L					
12	123	126	11	36	-30	11	36	-15	0	541	520		1.5,L	-1.4,L					
14			0.8,L	11	36	15	0	541	520	1	112	164	-4	180	-125				
							1	713	717	-1	120	164	-4	180	-125				
0.1,L	0	422	-416	0	16,L		-1	117	120	-2	314	307	-5	170	-161				
2	231	243	0	425	0	266	-262	-2	1409	-1479	-3	226	-219	-6	253	-242			
4	300	333	20	208	-207	1	125	-121	-3	330	-337	-3	194	-188	-7	53	-52		
6	136	357	53	534	52	225	-247	-5	51	-526	-4	442	-452	-7	27	-26			
8	165	519	5	514	3	107	-109	-5	426	-426	-5	526	-521	-7	59	-58			
10	814	-812	6	129	121	4	30	-22	5	33	113	-5	167	-160	-9	152	-150		
12	97	96	7	420	-414	5	111	113	5	117	119	-5	275	-275	-9	320	-319		
14	216	-225	8	56	57	6	327	-328	6	39	37	-6	31	-28	-6	64	-66		
10	107	-105	9	146	-141	7	157	-150	-6	210	-215	-6	107	-103	-7	100	-99		
12	99	-93	10	205	200	8	158	-154	-6	213	-207	-7	136	-134	-7	120	-119		
14	168	-163	11	207	-199	9	151	-135	-6	223	-219	-8	134	-131	-7	124	-123		
			0.2,L	12	97	66	11	115	117	9	213	-213	-7	122	-121	-7	121	-120	
14	123	164	-163	11	66	-72	-10	99	-105	-9	120	-122	-7	121	-120	-7	121	-120	
							-11	337	-336	-9	155	-155	-11	151	-151	-7	151	-151	
1	329	-334	0.9,L	0	17,L		-11	431	436	10	141	-144		1.10,L					
2	50	-445	1	769	-783	1	27	6	-13	138	136	-11	109	-104	-6	45	-35		
4	206	115	2	704	-709	1	27	6	-14	56	55	-12	128	-128	-6	169	-176		
6	852	195	3	207	-208	2	210	-210	137	140	-12	98	-98	-6	137	-138			
8	361	-359	4	123	-117	4	337	335	-137	140	-13	90	-93	-1	125	-127			
10	137	-143	6	166	-164	5	148	-149		1.2,L			1.6,L			32	-32		
12	201	188	7	135	-144	6	155	-155		0	464	448	0	224	-224	-6	365	-363	
14	328	-331	8	82	70	7	53	-45		1	254	-274	1	414	-414	-4	211	-209	
10	287	-215	10	335	-339	8	83	80	-1	295	-295	-1	586	-576	-6	241	-239		
12	161	153	11	190	-189	9	71	-71	-2	188	-187	-2	389	-378	-6	240	-238		
14	268	-269	12	120	-120	10	112	-112	-2	188	-187	-2	389	-378	-6	240	-238		
			0.10,L	11	106	0.16,L	-3	889	-905	-2	127	-127	-113	263	-263	-6	66	-67	
1	0	438	-439	0	105	-97	4	113	-110	-5	61	-61	-7	75	-75	-6	61	-61	
2	493	-490	1	236	230	1	201	211	-4	589	79	-4	228	-228	-7	50	-50		
4	73	-71	3	244	242	2	190	186	-6	61	-62	-7	122	-122	-6	34	-34		
6	398	-425	4	587	-576	4	342	336	-6	119	112	-5	116	-116	-7	300	-300		
8	85	79	5	324	323	5	66	-69	7	309	-310	-6	335	-334	-10	179	-178		
10	523	-240	7	302	317	6	321	-317	-7	155	-156	-6	52	-52	-7	173	-170		
12	186	179	8	83	84	7	86	86	-8	120	-117	-7	98	-107	-12	145	-145		
14	413	-420	9	166	-168	9	279	271	-8	117	-113	-7	132	-133	-13	247	-245		
			0.19,L	10	72	64	0.19,L	-282	-287	-9	207	-207	-9	253	-253	-7	243	-241	
10	63	-59	11	111	0.11,L	1	116	114	-10	180	-177	-10	518	-518	-4	525	-525		
12	42	-22	12	145	152	-11	41	41	-11	38	-10	211	-214	-4	140	-138			
14	134	-141	13	112	105	99	240	237	-12	47	48	-11	120	-118	-1	166	-177		
			0.4,L	14	53	-41	4	168	-176	-12	189	-194	-11	101	-104	-1	93	-93	
1	577	609	15	184	-184	5	69	-69	-13	143	-142	-12	136	-136	-7	307	-307		
2	605	-633	6	271	272	8	89	87	-14	107	102	-12	136	-136	-7	216	-216		
4	445	-473	7	34	-2	9	86	84	-14	135	-135	-12	136	-136	-7	216	-216		
6	141	-137	8	35	36	10	91	28	1	1.3,L		0	286	270	-4	47	-32		
8	952	-983	9	34	34	0	115	111	1	429	-429	-1	394	-382	-5	69	-67		
10	464	-441	10	292	285	0.20,L	0	118	108	-1	130	-130	-1	394	-382	-5	69	-67	
12	287	253	11	186	-186	0	115	111	1	429	-429	-1	394	-382	-5	69	-67		
14	377	353	13	188	-188	1	216	-223	2	27	27	-1	544	-526	-7	342	-342		
			0.12,L	15	24	9	34	34	3	760	757	-3	363	-363	-7	342	-342		
1	73	-69	16	31	-26	2	34	34	3	760	757	-3	363	-363	-7	342	-342		
2	184	186	17	31	-26	3	34	34	3	760	757	-3	363	-363	-7	342	-342		
4	111	-227	18	304	-287	6	39	-26	4	206	-204	-4	537	-537	-8	162	-162		
6	342	342	20	322	323	7	34	-39	4	206	-204	-4	531	-531	-8	162	-162		
8	132	-21	21	304	-287	6	39	-26	4	206	-204	-4	537	-537	-8	162	-162		
10	216	215	22	322	323	8	36	25	4	206	-204	-4	531	-531	-8	162	-162		
12	235	233	23	323	323	9	36	25	4	206	-204	-4	531	-531	-8	162	-162		
14	235	233	24	348	348	10	53	53	4	206	-204	-4	531	-531	-8	162	-162		
			0.5,L	15	95	94	0.21,L	-373	-357	-7	238	-242	-11	154	-160	-4	160	-160	
1	25	-13	8	32	-30	1	103	97	7	206	198	-6	46	-43	-12	137	-137		
2	230	-251	9	325	322	2	251	-256	7	95	-93	-5	36	-31	-7	117	-117		
4	154	-146	10	271	274	4	45	-36	8	164	-166	-5	230	-230	-7	117	-117		
6	700	700	11	78	69	5	90	-90	9	150	-152	-5	209	-209	-7	117	-117		
8	199	200	12	129	-125	6	79	-79	9	154	-156	-5	209	-209	-7	117	-117		
10	320	331	13	129	-125	7	45	53	9	154	-156	-5	209	-209	-7	117	-117		
12	320	331	14	143	-143	8	45	37	9	154	-156	-5	209	-209	-7	117	-117		
14	100	96	15	132	-132	0	123	-117	7	44	-40	-6	135	-139	-10	117	-117		
			0.6,L	16	65	60	6	90	-84	1	144	-144	-6	222	-222	-6	167	-167	
1	392	396	17	9	35	30	0.22,L	-10	198	-154	-12	105	105	-4	223	-223	-6	164	-164
2	93	-109	18	169	-170	45	45	0.23,L	-1	661	-662	-5	222	-223	-6	164	-164		
4	323	399	19	45	-45	0.14,L	1	179	-177	2	280	-272	-3	165	-165	-8	225	-225	
6	149	155	20	487	484	2	48	51	-2	724	-713	-3	817	-805	-8	190	-190		
8	312	-301	21	270	263	2	100	-107	-3	765	768	-4	127	-123	-9	165	-165		
10	165	167	22	487	484	4	45	-36	-3	529	518	-4	226	-212	-10	169	-169		
12	793	781	23	31	26	4	100	-107	-4	529	518	-4	226	-212	-10	169	-169		
14	335	331	24	270	263	4	203	-205	-4	529	518	-4	226	-212	-10	169	-169		
			0.6,L	25	487	484	4	203	-205	-4	529	518	-4	226	-212	-10	169	-169	
1	228	235	26	311	-309	5	130	-130	-5	263	266	-4	164	-164	-7	225	-225		
2	111	195	27	139	143	6	69	-69	6	263	266	-4	164	-164	-7	225	-225		
4	128	-122	28	5	64	-35	7	137	-132	0	123	-117	-6	135	-139	-10	173	-173	
6	58	-56	29	6	39	-35	7	137	-132	0	123	-117	-6	135	-139	-10	173	-173	
8	0.7,L	274	-284	20	200	-200	-2	170	169	5	313	314	-6	248	-248	-6	226	-226	
10	116	-123	21	245	-243	-4	495	-474	9	194	-201	-10	121	-111	-7	173	-173		
12	639	647	22	174	-174	-6	670	646	-10	348	-348	-12	488	-488	-6	165			

Table 2. Continued.

1.13,L	-7	112	113	-2	106	-101	-8	313	-307	-4	182	176	-7	92	69		
-4	115	-120	10	51	8	134	-136	-3	362	-357	9	30	35	6	138	-139	
-5	32	-27	1	19	-1	179	-176	-1	221	-210	-10	58	57	6	142	-143	
-6	81	-20	1	19	1,19,L	-1	135	-129	-11	365	-370	-7	103	-106	2,18,L		
-7	164	165	-1	75	-79	6	207	-210	-11	390	-381	6	164	-162	1		
-8	131	135	-1	75	-79	6	227	-224	-12	154	-153	6	36	33	-101		
-9	463	-476	-2	131	130	-6	562	-560	-12	110	-106	5	107	-106	66		
-10	136	137	-2	202	207	7	313	-311	-10	153	-141	1	35	33	-56		
-11	216	-222	-3	78	-81	-7	2,7,L	-2,7,L	-10	270	-256	-2	118	-116	-54		
-12	240	248	-3	180	187	5	184	-183	-5	156	-156	16	155	156	-148		
-13	42	56	4	121	119	5	203	-213	-5	2,7,L	-2,7,L	-10	155	-153	-71		
-14	74	79	-4	42	29	9	60	-63	0	45	-44	2,12,L	-4	69	-71		
-15	51	-53	5	69	-71	-9	105	-102	-1	169	-177	6	166	-167	0		
-16	157	158	6	79	-79	10	126	-127	-2	91	-86	0	384	-388	-61		
-17	49	-45	-6	74	-73	-10	191	-184	-2	170	-170	1	54	-48	-21		
-18	154	-150	-7	218	-220	11	41	-28	3	154	-141	-1	35	-3	74		
0	182	-184	1,20,L	-8	47	-55	12	89	-85	4	168	-166	-167	0	55	-58	
-1	159	-169	1,20,L	-13	68	85	5	85	-85	5	146	-136	3	68	-66	1	
-2	477	-481	0	244	-250	2,3,L	-6	6	312	311	-4	219	-227	-1	152	-146	
-3	62	-72	1	65	-62	2,3,L	-6	6	556	-551	3	160	-155	1	145	-147	
-4	145	-147	-1	170	-170	0	153	-141	-7	43	-59	5	467	-461	-165		
-5	254	-259	3	138	-136	-6	764	-738	-8	143	-142	1	35	-35	-54		
-6	72	75	-3	66	-48	-1	581	-571	9	56	-56	10	61	-64	-137		
-7	168	160	4	54	-54	2	43	-43	-9	257	-254	11	138	-121	2,20,L		
-8	328	-337	-4	189	192	3	76	-75	10	36	-38	-11	60	57	0		
-9	43	46	5	36	-22	-3	89	-86	-10	78	79	2,13,L	0	135	-132		
-10	359	-369	-5	132	-128	4	292	-288	11	57	-44	0	104	-103	-208		
-11	16	10	6	148	-156	2	274	-272	-11	66	-59	-5	342	4	144	-140	
-12	73	83	7	125	-125	2	82	-82	12	92	-74	-1	65	-64	89	-92	
-13	177	-177	-7	90	-85	5	514	-506	-12	143	-149	1	353	-342	171	-167	
-14	124	132	6	90	82	3	2,8,L	-2,8,L	3	254	-256	-5	171	-166	-156		
-15	149	152	1,21,L	-6	49	34	3	261	-258	-5	156	-156	-5	156	-156	-156	
-16	56	52	2	106	106	7	162	-158	0	435	426	-4	53	53	2,21,L		
-17	98	-102	0	110	-106	-7	323	-324	1	500	-506	5	368	-371	2,21,L		
-18	42	-53	5	101	-97	5	24	-24	1	326	-325	1	156	-156	1		
-19	49	55	-2	180	-182	9	182	-182	-2	101	-102	6	203	-210	1		
0	1,15,L	-3	217	-213	-17	60	63	-2	42	44	7	104	-107	-1	66	73	
-1	526	-533	-3	48	-45	-11	110	-106	3	246	-242	8	41	-34	2	134	133
-2	237	243	5	100	96	13	202	-199	-4	127	-124	9	85	-81	3,0,L		
-3	49	-46	5	147	-139	-13	88	-87	5	267	-266	10	90	-84	0		
-4	123	-203	-6	197	-196	2	216	-216	-10	67	-60	0	48	-46	314	313	
-5	100	-139	1,22,L	-6	65	67	2,4,L	-2,4,L	7	161	-163	-4	272	-274	2,4,L		
-6	226	-230	1,22,L	-7	38	43	2,14,L	-2,14,L	6	161	-163	4	272	-274	2,14,L		
-7	299	295	-1	71	-72	1	126	-123	-8	416	-407	6	39	32	-4	54	46
-8	222	-221	-2	84	83	-9	1	80	-83	-432	-300	1	34	32	5	111	-106
-9	60	57	3	84	-73	2	537	-527	10	49	-49	-1	127	129	5	393	-393
-10	166	-164	4	223	-221	3	307	-301	11	65	-71	-2	166	-165	5	266	-266
-11	131	-136	-4	41	-49	-3	38	-33	-11	153	-145	3	230	-231	-1	150	-150
-12	168	-171	-4	202	-202	4	202	-201	-12	104	-96	4	168	-169	3,1,L		
-13	87	-91	1,23,L	-5	352	91	2,9,L	-2,9,L	5	64	65	0	105	-109	0		
-14	47	52	0	60	65	6	102	-97	7	161	-163	-5	203	-203	2,4,L		
-15	46	38	-2	70	63	9	136	-131	0	39	42	6	133	-132	1,24,L		
-16	59	-54	-2	74	73	10	466	-465	-1	207	-209	6	59	-58	154	148	
0	1,16,L	-3	47	-47	-7	83	83	-1	123	115	7	104	-105	120	123	123	
-1	97	-102	2,0,L	-8	119	120	3	74	-73	95	91	4	40	-36	4	97	98
-2	257	-264	0	467	-455	10	133	-135	3	338	-334	6	246	-242	-4	238	234
-3	184	183	0	275	-275	11	135	-135	4	204	-203	5	173	-169	29	125	124
-4	244	251	2	325	-325	12	135	-135	-4	103	-105	10	236	-236	5	266	-266
-5	33	-32	-2	551	-547	-11	178	-181	8	185	-183	15	179	-179	6	359	-359
-6	70	67	4	120	-118	12	166	-165	-5	137	-137	130	190	-187	5	238	239
-7	171	-168	-4	452	-440	-13	203	-202	6	197	-197	190	190	-190	7	94	96
-8	273	289	6	101	104	7	149	-145	0	104	105	7	56	-56	161	161	
-9	99	100	-6	269	-266	2,5,L	-7	180	180	2	72	65	8	62	61	161	
-10	53	-31	8	160	160	6	102	-98	8	45	-45	163	157	-8	62	55	161
-11	155	147	-8	76	-68	1	299	-277	-8	160	-161	5	257	-257	6	167	162
-12	197	199	10	86	-73	1	496	-496	9	93	90	2	236	-228	-9	182	182
-13	351	-358	-10	89	-89	-2	176	-170	-9	109	108	4	442	-431	7	77	-66
-14	47	51	-12	45	29	3	244	-234	10	196	-197	6	61	-64	-10	143	145
-15	219	-224	-12	33	-23	-3	151	-152	-10	185	-180	6	96	93	11	199	205
-16	183	183	4	496	-489	11	170	-166	-11	96	-94	7	65	-84	-11	175	-177
0	1,17,L	-4	510	510	-106	5	168	-168	7	168	-168	14	14	14	3,2,L		
-1	366	372	1	328	-320	6	130	-127	0	641	612	-9	143	133	0	153	152
-2	170	-166	-1	475	-458	6	215	-214	0	641	612	-9	143	133	-1	168	161
-3	114	119	2	86	74	7	68	-89	1	34	22	1	200	202	-1	200	202
-4	74	76	-2	85	75	-7	264	-264	-1	141	-137	2,16,L	-2,16,L	-1	195	-195	
-5	261	267	3	63	56	8	404	-397	-2	201	202	3	337	333	0	161	155
-6	132	132	-3	146	-140	10	180	-181	1	161	-163	0	113	-109	3	354	354
-7	133	134	4	146	-142	12	152	-153	3	163	-163	1	277	278	4	165	165
-8	331	334	-4	34	29	10	115	-108	-3	203	-195	-1	232	-232	5	300	303
-9	369	-380	5	325	-306	-10	159	-157	4	212	-214	2	228	-223	6	91	93
-10	77	82	5	145	133	11	81	-84	-4	442	-431	2	228	-223	6	301	303
-11	101	105	6	379	372	-11	167	-165	3	337	-336	4	135	137	-1	161	-112
-12	81	-81	-6	49	-35	12	141	-142	-5	370	-369	-3	218	217	7	99	-96
-13	47	37	-7	201	-199	-12	210	-212	6	70	-72	1	117	-117	1	156	-156
-14	89	89	9	213	-215	5	177	-175	-7	60	-58	6	237	-232	-8	38	36
-15	55	-58	9	163	156	61	48	-426	8	54	-52	7	131	-131	-9	74	72
0	1,18,L	-9	142	-139	0	113	-110	-8	182	162	-8	132	-132	10	147	-143	
-1	171	-172	-10	147	-146	-1	294	-284	-9	84	-84	-274	2,17,L	2,17,L	3,3,L		
-2	321	-322	12	146	138	2	288	-279	10	268	-268	-2	217	217	-1	208	-209
-3	127	-127	-12	100	-97	-2	353	-352	-10	42	-42	-37	2,11,L	2,11,L	-1	208	-209
-4	201	213	-13	47	48	-3	426	-415	1	230	225	-2	199	-203	1	72	-74
-5	99	-107	2,2,L	-4	48	-4	245	-241	1	37	-40	-3	103	102	-2	40	36
-6	39	-43	0	30	35	-5	52	51	-1	37	-442	4	138	-134	-1	343	-359
-7	201	202	-1	507	50	-6	367	-378	3	57	60	2					

Table 2. Continued.

	3.3.L	-6	93	-99		3.10.L	-7	98	-93	-6	75	75	2	149	150			
5	325	-323	-7	108	99		302	-298	3,14,L	4,1,L			-2	137	141			
-5	122	-123	-8	51	-1	145	167	0	103	98	0	207	-210	-3	41	42		
6	132	-130	-10	105	-165	-1	44	56	0	277	288	1	108	112	4	24	242	
-7	132	-130	-10	131	-132	-2	68	80	-1	169	168	-1	68	62	5	63	63	
-7	153	152					88	-93	-1	169	168	-1	68	62	5	49	50	
8	65	-66					195	-205	2	41	19	-2	142	-146	-6	110	-116	
-8	121	119					326	-327	-3	41	34	-3	135	140				
9	119	-123	0	43	-46	5	103	-100	5	37	20	-4	48	43				
-9	70	23	1	25	38	6	57	-55	5	159	144	-4	74	65				
10	45	48	-2	110	112	-6	132	131	6	89	-63	6	399	-407	0	106	105	
-10	135	134	-2	40	22	8	239	-232	-6	64	-62	6	212	7	110	-119	68	
-11	147	146	-3	96	98	-8	65	64	-7	209	212	-7	150	155	-3	50	51	
							178	-182	9	160	157				5	79	75	
							108	-112	-9	164	-172				6	72	74	
							399	403							-6	251	261	
-1	295	-285	5	243	58				0	143	138							
2	316	-322	-5	58	58				1	111	-110	1	124	-125				
-2	146	-149	6	44	-34	0	164	-165	2	126	129	-1	124	-125				
3	75	-70	7	198	193		56	-54	3	178	-176		80	-85				
-3	362	-368	-7	75	-77	-1	66	-69	4	112	106		129	-135	0	225	-235	
4	101	-96	8	34	-26	-2	295	-287	-4	194	192	-3	58	-54	1	67	67	
-4	83	85	9	131	134	-3	65	-65	5	64	-60	4	64	-67	-1	166	152	
5	236	-235	-9	138	-144	4	59	-54	5	71	72	-4	68	-67	-2	52	-55	
-5	33	-33	10	46	32	3	40	-39	6	36	-25	5	251	-262	3	51	51	
7	174	173	-10	200	205	-5	66	-69	-6	72	69	7	77	-76	-4	68	70	
-8	152	157				6	105	-108							-4	175	185	
9	186	187				-6	271	-273							-5	180	185	
-9	170	-171				7	65	-64										
10	131	-131	1	231	223	-7	72	78	0	77	-76	0	66	63				
-10	97	93	-1	105	100	8	45	35	1	70	76	-1	58	-59				
						2	129	-127	1	129	-128	-1	126	-125	1	65	-66	
						3	50	58	0	58	-58	-1	122	-121	-1	162	-160	
						4	100	-104	1	100	-104	-1	101	-102	-1	132	-131	
						5	53	58	4	56	-56	-4	141	-140				
						6	76	79	5	76	-76	-3	166	-165				
						7	100	-102	6	100	-102	-1	100	-102				
						8	195	-203	7	195	-203							
						9	100	-102	8	100	-102							
						10	100	-102	9	100	-102							
						11	100	-102	10	100	-102							
						12	100	-102	11	100	-102							
						13	100	-102	12	100	-102							
						14	100	-102	13	100	-102							
						15	100	-102	14	100	-102							
						16	100	-102	15	100	-102							
						17	100	-102	16	100	-102							
						18	100	-102	17	100	-102							
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						37	100	-102	36	100	-102							
						38	100	-102	37	100	-102							
						39	100	-102	38	100	-102							
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						41	100	-102	40	100	-102							
						42	100	-102	41	100	-102							
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						44	100	-102	43	100	-102							
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						59	100	-102	58	100	-102							
						60	100	-102	59	100	-102							
						61	100	-102	60	100	-102							
						62	100	-102	61	100	-102							
						63	100	-102	62	100	-102							
						64	100	-102	63	100	-102							
						65	100	-102	64	100	-102							
						66	100	-102	65	100	-102							
						67	100	-102	66	100	-102							
						68	100	-102	67	100	-102							
						69	100	-102	68	100	-102							
						70	100	-102	69	100	-102							
						71	100	-102	70	100	-102							
						72	100	-102	71	100	-102							
						73	100	-102	72	100	-102							
						74	100	-102	73	100	-102							
						75	100	-102	74	100	-102							
						76	100	-102	75	100	-102							
						77	100	-102	76	100	-102							
						78	100	-102	77	100	-102							
						79	100	-102	78	100	-102							
						80	100	-102	79	100	-102							
						81	100	-102	80	100	-102							
						82	100	-102	81	100	-102							
						83	100	-102	82	100	-102							
						84	100	-102	83	100	-102							
						85	100	-102	84	100	-102							

Table 3. Fractional atomic coordinates.

	<i>x</i>	<i>y</i>	<i>z</i>
K <sup>+</sup>	0.2825(1)	0.1554(1)	0.0242(1)
N(1)	0.4725(7)	0.3586(1)	0.1933(2)
C(2)	0.5465(7)	0.3006(1)	0.1491(2)
O(2)	0.4649(6)	0.2852(1)	0.0571(1)
N(3)	0.7314(6)	0.2564(1)	0.2146(1)
O(3)	0.8180(5)	0.1970(1)	0.1667(1)
C(4)	0.8479(7)	0.2660(1)	0.3193(2)
O(4)	0.7810(5)	0.3168(1)	0.3709(1)
N(5)	1.0356(6)	0.2167(1)	0.3594(2)
N(1')	0.9760(7)	-0.0274(1)	0.1482(2)
C(2')	0.8047(6)	0.0298(1)	0.1502(2)
O(2')	0.7428(5)	0.0618(1)	0.0683(1)
N(3')	0.6879(5)	0.0520(1)	0.2504(1)
O(3')	0.4831(5)	0.1083(1)	0.2475(1)
C(4')	0.7963(7)	0.0300(1)	0.3509(2)
O(4')	0.9568(5)	-0.0224(1)	0.3610(1)
N(5')	0.7118(7)	0.0680(1)	0.4339(2)
H(11)	0.517(10)	0.366(1)	0.262(3)
H(12)	0.345(10)	0.386(1)	0.158(3)
H(3)	0.717(22)	0.166(4)	0.198(6)
H(51)	1.078(10)	0.182(2)	0.317(3)
H(52)	1.113(09)	0.217(1)	0.430(3)
H(1'1)	1.036(10)	-0.044(1)	0.208(3)
H(1'2)	1.063(10)	-0.038(1)	0.087(3)
H(3')	0.574(24)	0.140(4)	0.219(7)
H(5'1)	0.604(10)	0.107(2)	0.420(3)
H(5'2)	0.795(10)	0.057(1)	0.497(3)

*The 3-HB anions.* Fig. 3 shows the geometry of the anion together with the bond lengths and angles. No significant differences in the dimensions of the two non-equivalent 3-HB residues were found, and in addition the bond lengths and angles are in agreement with literature values for comparable molecules, e.g. biuret,<sup>8-9</sup> and hydroxyurea.<sup>10</sup>

An analysis of planarity of the 3-HB residues was carried out by calculation of the best plane through the non-hydrogen atoms of each residue and in addition through the N, C, and O atoms of each half of the residues (cf. Table 5). The analysis clearly shows that the atoms defining plane 1 is more coplanar than the atoms defining plane 2. In residue 1 there is a twist of only 3.4° between the two H<sub>2</sub>N-(C=O)-N-moieties; the corresponding angle of residue 2 is 13.2°. Both residues have adapted the conformation that gives rise to an *intramolecular* NH···O hydrogen bond (cf. Fig. 2). The same conformation was found in the biuret molecule<sup>8</sup> and the perdeuterated biuret molecule,<sup>9</sup> where the length of the *intramolecular* hydrogen bond was found to be 2.72 Å and the twisting angle 6.3°.

*The environment of the potassium ion.* The potassium ion has eight oxygen neighbours belonging to six different 3-HB residues, with K-O distances in the range 2.663–3.026 Å (cf. Fig. 4). These values are in good agreement

Table 4. Thermal parameters for the non-hydrogen atoms.  
T<sub>i</sub>=exp [-1/4 (B<sub>11</sub>h<sup>2</sup>a\*<sup>2</sup>+...+2B<sub>23</sub>klb\*c\*)].

	B <sub>11</sub>	B <sub>22</sub>	B <sub>33</sub>	B <sub>12</sub>	B <sub>13</sub>	B <sub>23</sub>
K <sup>+</sup>	1.62(02)	2.12(02)	1.81(03)	0.19(02)	-0.07(02)	0.09(02)
N(1)	3.56(11)	1.95(09)	2.56(06)	0.65(10)	-0.44(10)	-0.14(09)
C(2)	1.71(11)	1.79(10)	1.76(11)	-0.22(06)	0.12(09)	0.09(09)
O(2)	3.75(11)	2.71(10)	1.91(09)	0.21(08)	-0.84(08)	-0.10(07)
N(3)	1.95(10)	1.51(09)	1.48(09)	0.13(08)	-0.03(07)	-0.35(07)
O(3)	2.55(09)	1.77(08)	1.86(08)	0.44(07)	0.28(07)	-0.38(07)
C(4)	1.58(11)	2.00(12)	1.72(11)	-0.59(10)	0.20(09)	-0.01(07)
O(4)	2.81(09)	2.15(09)	1.81(08)	-0.08(07)	-0.13(07)	-0.60(07)
N(5)	2.63(12)	2.37(11)	1.95(10)	0.25(09)	-0.47(09)	-0.02(09)
N(1')	3.28(09)	2.01(11)	1.89(10)	0.89(10)	0.27(09)	0.08(09)
C(2')	1.60(11)	1.69(11)	1.88(11)	-0.18(09)	-0.07(09)	-0.17(09)
O(2')	2.89(09)	2.07(08)	1.52(08)	0.50(07)	0.02(07)	0.27(06)
N(3')	1.82(10)	1.58(09)	1.69(09)	0.33(08)	0.05(08)	0.22(07)
O(3')	2.15(09)	1.94(09)	2.15(09)	0.70(07)	0.29(07)	0.13(07)
C(4')	1.77(11)	1.80(12)	1.86(11)	-0.30(09)	0.00(08)	0.13(09)
O(4')	3.17(10)	1.98(09)	2.03(08)	0.44(08)	-0.29(07)	0.20(07)
N(5')	3.44(13)	2.62(12)	1.72(10)	0.68(10)	-0.04(09)	-0.10(09)

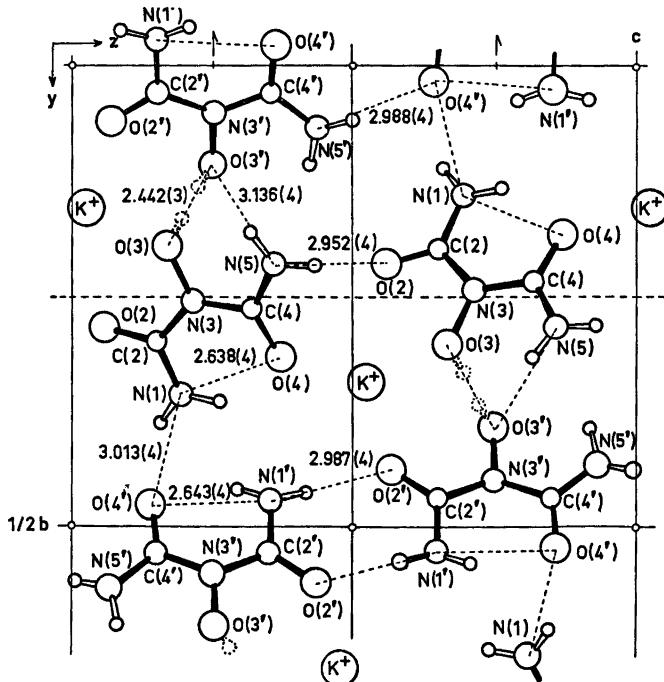


Fig. 2. The structure viewed along the  $a$ -axis.

Table 5. Least-squares planes. The equations of the planes can be expressed as  $Px + Qy + Rz = S$  in direct space.

	Plane 1	Plane 2	Plane 3	Plane 4	Plane 5	Plane 6
$P$	3.304	3.253	3.343	3.227	3.326	3.337
$Q$	8.054	11.019	7.584	8.706	10.059	10.099
$R$	-4.258	0.083	-4.270	-4.391	1.546	-1.314
$S$	3.591	2.939	3.472	3.650	3.201	2.494

Distances in Å of the atoms from the planes.

	Plane 1	Plane 2	Plane 3	Plane 4	Plane 5	Plane 6
N(1)'	0.03	N(1') -0.05	N(1) 0.001	N(3) 0.000	N(1') -0.003	N(3') -0.002
C(2)'	0.00	C(2') 0.02	C(2) -0.003	C(4) 0.000	C(2') 0.008	C(4') 0.005
O(2)'	0.00	O(2') 0.16	O(2) 0.001	O(4) 0.000	O(2') -0.003	O(4') -0.002
N(3)'	-0.02	N(3') -0.11	N(3) 0.001	N(5) 0.000	N(3') -0.002	N(5') -0.002
O(3)'	-0.01	O(3') -0.15				
C(4)'	-0.01	C(4') 0.01				
O(4)'	-0.04	O(4') -0.04				
N(5)'	0.05	N(5') 0.16				

Additional atoms:

Angles between normals to planes:

H(11)	-0.04	H(1'1)	-0.04	1:2	21.9°
H(12)	-0.01	H(1'2)	0.10	3:4	3.6°
H(51)	0.09	H(5'1)	0.25	5:6	13.2°
H(52)	0.01	H(5'2)	0.32		
H(3)	-0.72	H(3')	0.50		

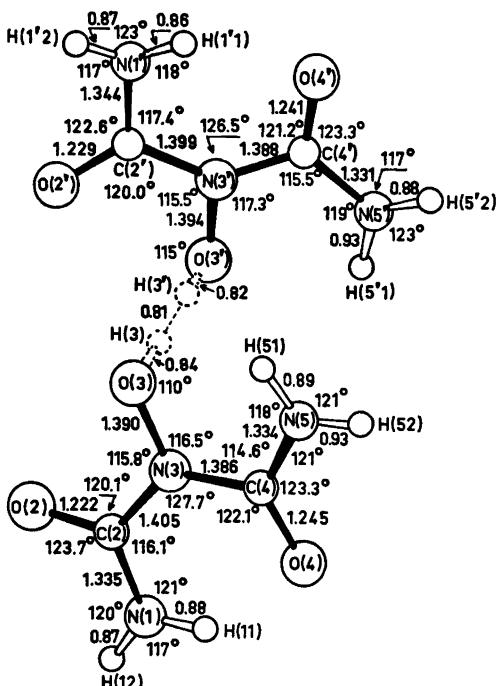


Fig. 3. Bond lengths and valency angles. The standard deviations on bonds between non-hydrogen atoms are 0.003–0.004 Å, and on the angles 0.2–0.3°. The standard deviations on bonds to hydrogen atoms are 0.03–0.04 Å and on angles including hydrogen atoms 2–6°.

with those earlier reported for eight-coordinated potassium,<sup>3</sup> e.g. in the structure of potassium hydrogen diformate.<sup>11</sup> As in this structure there is no regular geometry of the arrangement (cf. Table 6). It seems likely that the reason for

this is the intensive *intermolecular* hydrogen bonding in the structure.

Two of the coordinating oxygen atoms, O(2) and O(3), belong to residue 1, and other two, O(2') and O(3'), belong to residue 2 (cf. Fig. 4), but the planes through the chelate ligands do not pass through the central potassium ion. The displacements of K<sup>+</sup> from the planes defined by O(2)–C(2)–N(3)–O(3) and O(2')–C(2')–N(3')–O(3') are –1.51 and –0.47 Å, respectively. A similar situation was found in the potassium acid salts of isonitrosoacetophenone,<sup>12</sup> and the cyclic hydroxamic acid 5-bromo-3-hydroxy-6-methyl-uracil,<sup>13</sup> and is not a matter for surprise considering the electrostatic nature of the bonding. Nevertheless, the chelate rings in the structures of the potassium acid salts of carboxylic acids are often found to be planar.<sup>11</sup>

The number of ionic contacts to potassium is different for the two 3-HB residues: residue 1 has five and residue 2 has three contacts to K<sup>+</sup>. This may mean that there is a tendency towards less ionization in residue 2 than in residue 1.

The distance between the layers of the potassium ions is about 6 Å ( $\sim \frac{1}{2}c$ ). The nearest potassium-potassium contact is 3.868 Å (=a).

*The short hydrogen bond.* In most structures with very short hydrogen bonds, e.g. the acid salts of carboxylic acids, the short hydrogen bond has to be symmetric, because the acidic residues are crystallographically symmetry related. In the present structure the hydrogen bond, which links the 3-HB residues to a “dimer”, has no crystallographical symmetry.

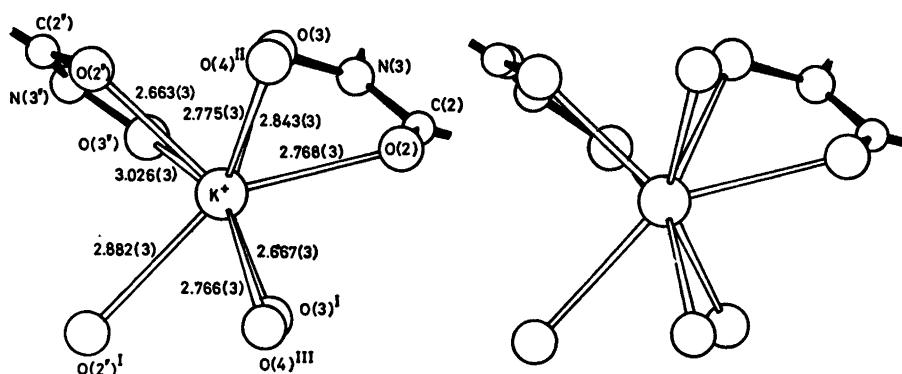


Fig. 4. Stereoscopic view of the potassium ion with its eight oxygen neighbours. In addition the chelate ligands of each 3-HB residue are indicated.

Table 6. The 28 independent O—K—O angles ( $^{\circ}$ ). The standard deviations are 0.05–0.07 $^{\circ}$ .

O(2)—K <sup>+</sup> —O(3)	55.9	O(2')—K <sup>+</sup> —O(3) <sup>I</sup>	123.0
O(2)—K <sup>+</sup> —O(2')	119.0	O(2')—K <sup>+</sup> —O(2') <sup>I</sup>	88.4
O(2)—K <sup>+</sup> —O(3')	96.0	O(2')—K <sup>+</sup> —O(4) <sup>II</sup>	79.6
O(2)—K <sup>+</sup> —O(3') <sup>I</sup>	76.7	O(2')—K <sup>+</sup> —O(4) <sup>III</sup>	138.8
O(2)—K <sup>+</sup> —O(2') <sup>I</sup>	142.1	O(3')—K <sup>+</sup> —O(3) <sup>I</sup>	70.4
O(2)—K <sup>+</sup> —O(4) <sup>II</sup>	74.3	O(3')—K <sup>+</sup> —O(2') <sup>I</sup>	78.4
O(2)—K <sup>+</sup> —O(4) <sup>III</sup>	94.8	O(3')—K <sup>+</sup> —O(4) <sup>II</sup>	121.1
O(3)—K <sup>+</sup> —O(2')	66.6	O(3')—K <sup>+</sup> —O(4) <sup>III</sup>	150.2
O(3)—K <sup>+</sup> —O(3')	49.1	O(3) <sup>I</sup> —K <sup>+</sup> —O(2') <sup>I</sup>	66.0
O(3)—K <sup>+</sup> —O(3) <sup>I</sup>	89.1	O(3) <sup>I</sup> —K <sup>+</sup> —O(4) <sup>II</sup>	149.7
O(3)—K <sup>+</sup> —O(2') <sup>I</sup>	127.2	O(3) <sup>I</sup> —K <sup>+</sup> —O(4) <sup>III</sup>	85.3
O(3)—K <sup>+</sup> —O(4) <sup>II</sup>	81.9	O(2') <sup>I</sup> —K <sup>+</sup> —O(4) <sup>II</sup>	140.5
O(3)—K <sup>+</sup> —O(4) <sup>III</sup>	150.6	O(2') <sup>I</sup> —K <sup>+</sup> —O(4) <sup>III</sup>	76.1
O(2')—K <sup>+</sup> —O(3')	54.4	O(4) <sup>II</sup> —K <sup>+</sup> —O(4) <sup>III</sup>	88.6

Roman numerals as superscripts denote the following equivalent positions relative to the reference molecule of  $x, y, z$ : I  $x - 1, y, z$ ; II  $x, -y + \frac{1}{2}, z - \frac{1}{2}$ ; III  $x - 1, -y + \frac{1}{2}, z - \frac{1}{2}$ .

Table 7. Interatomic angles ( $^{\circ}$ ) concerning the hydrogen bonding system.

N(1)—H(11)—O(4)	131(3)
N(1)—H(12)—O(4') <sup>IV</sup>	156(3)
O(3)—H(3)—O(3')	172(9)
O(3)—H(3')—O(3')	170(9)
N(5)—H(51)—O(3') <sup>V</sup>	148(3)
N(5)—H(52)—O(2) <sup>VI</sup>	158(3)
N(1')—H(1'1)—O(4')	134(4)
N(1')—H(1'2)—O(2') <sup>VII</sup>	176(4)
N(5')—H(5'2)—O(4') <sup>VIII</sup>	172(4)

Roman numerals as superscripts denote the following equivalent positions relative to the reference molecule of  $x, y, z$ : IV  $-x + 1, y + \frac{1}{2}, -z + \frac{1}{2}$ ; V  $x + 1, y, z$ ; VI  $x + 1, -y + \frac{1}{2}, z + \frac{1}{2}$ ; VII  $-x + 2, -y, -z$ ; VIII  $-x + 2, -y, -z + 1$ .

But the short O···O distance of 2.442(3) Å, the favourable N—O···O angles of 109.8(3) $^{\circ}$  and 108.5(2) $^{\circ}$ , respectively, and the almost identical geometry of the 3-HB-residues being connected, are consistent with the existence of a virtually symmetrical hydrogen bond.<sup>14–16</sup> Symmetry due to statistical disorder is also possible. The rather elongated peak between the oxygen atoms of the difference map could perhaps be interpreted as an unresolved peak due to two half H-atoms about 0.4 Å on either side of the midpoint. The final refinement ( $\sim R = 0.032$ ) was based on this assumption.

But the X-ray data were also consistent with an asymmetric hydrogen bond. Refinement of the positional parameters of the hydrogen atom, with a start position midway between the

oxygen atoms, led to the following, perhaps most reliable, arrangement of the system: O(3)—H(3)=1.31(4) Å, O(3')—H(3)=1.13(4) Å and the angle O(3)—H(3)—O(3')=179(3) $^{\circ}$ . In fact the O(3')—H(3) bond has just the length corresponding to an O···O distance of 2.44 Å, using the curve, given by Lundgren,<sup>16</sup> of O—H distance as a function of O···O distance.

The conclusion must be that it is not possible from the X-ray data to establish whether this hydrogen bond is symmetric or slightly asymmetric.

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