

The Crystal Structures of L-Citrulline Hydrochloride and L-Homocitrulline Hydrochloride

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The crystal structures of L-citrulline hydrochloride and its homologue L-homocitrulline hydrochloride have been determined using diffractometer data. L-Citrulline hydrochloride: $\text{NH}_2\text{CONH}(\text{CH}_2)_3\text{CH}(\text{NH}_3^+)\text{COOH} \cdot \text{Cl}^-$, monoclinic, $C2$, $a=18.089$, $b=5.150$, $c=11.918$ Å, $\beta=106.9^\circ$, $R=0.040$, the average e.s.d. of the bond distances 0.005 Å. L-Homocitrulline hydrochloride: $\text{NH}_2\text{CONH}(\text{CH}_2)_4\text{CH}(\text{NH}_3^+)\text{COOH} \cdot \text{Cl}^-$, orthorhombic, $P2_12_12_1$, $a=10.518$, $b=21.627$, $c=4.916$ Å, $R=0.069$, the average e.s.d. of the bond distances 0.008 Å. There is a close similarity between the dimensions of the citrulline and homocitrulline molecules in the two crystals. Also there are several common features in the two hydrogen bond schemes. The present L-citrulline HCl structure is compared with that reported previously (Naganathan & Venkatesan, *Acta Cryst.* (1971), B26, 790), and some significant differences between the two results are pointed out.

Introduction

Citrulline is an intermediate in the ornithine cycle in mammals. In nature it is found in the water melon. Although this is not a constituent amino acid of proteins, the conformation and the size of this acid are of importance in connexion with the basic amino acids such as lysine, arginine and ornithine. The present paper deals with the crystal structures of L-citrulline hydrochloride,

$\text{NH}_2\text{CONH}(\text{CH}_2)_3\text{CH}(\text{NH}_3^+)\text{COOH} \cdot \text{Cl}^-$,
and its homologue L-homocitrulline hydrochloride,
 $\text{NH}_2\text{CONH}(\text{CH}_2)_4\text{CH}(\text{NH}_3^+)\text{COOH} \cdot \text{Cl}^-$.

When the present paper was under preparation, the crystal structure of L-citrulline hydrochloride was presented by Naganathan & Venkatesan (1971). There are, however, small but significant discrepancies in the two structures obtained independently. The unit cell dimensions and some of the bond distances in their results are a little smaller than those in ours. For a few atoms, their y parameters deviate significantly from ours. Thus our structure of L-citrulline hydrochloride is also reported here, although the primary purpose of the present paper lies in the critical comparison of the structures of citrulline and homocitrulline.

Experimental

Each crystal was obtained by recrystallization from a solution of the amino acid in a dilute hydrochloride containing a small amount of acetone. Preliminary X-ray work on the crystals was done using the photographic method. The determination of the cell size and the intensity measurements were performed on the

Rigaku Denki Computer-controlled four-circle diffractometer (AFC-II and IV). Zr-filtered Mo $K\alpha$ radiation ($\lambda=0.71069$ Å) was used with a pulse height analyser, and integrated intensities were measured by the $\omega/2\theta$ -scan method, in which the background intensity was estimated from the intensity measurements at the starting and the end points of every scan. Two or three standard reflexions were monitored after the measurement of every 50 reflexions. The intensities were corrected for Lorentz and polarization factors, but no absorption correction was made in view of the small linear absorption coefficient for Mo $K\alpha$ radiation. The crystallographic data and some details of the intensity measurement are listed in Table 1.

Structure determination

The procedures for the structure determination and the refinement were almost the same for both crystals. Most of the non-hydrogen atoms could be located from the heavy atom Fourier synthesis based on the chloride ion. The hydrogen atoms were found on the difference electron density map, and were included in the refinement. The computation was done on the NEAC 2200-500 computer in the Computing Centre of this University. The block-diagonal least-squares program *HBL5 IV* (Ashida, 1967) was used, which minimizes $\sum w(|F_o| - |F_c|)^2$. The weighting scheme was: $w=1.0$ when $F_o \neq 0$ and $w=0.5$ when $F_o=0$. The scattering factors for all the atoms were taken from *International Tables for X-ray Crystallography* (1962).

The structure of L-citrulline hydrochloride

The final R value is 0.042 for all reflexions. The final atomic parameters are listed in Tables 2 and 3, and the observed and calculated structure factors in Table 4.

Table 1. *The crystallographic data and experimental conditions*

	Citrulline hydrochloride	Homocitrulline hydrochloride
Crystallographic data		
Shape of crystals	plate	needle
Crystal system	monoclinic	orthorhombic
Space group	C2	$P2_12_12_1$
<i>a</i>	18.089	10.518 Å ($\pm 0.1\%$)
<i>b</i>	5.150	21.627
<i>c</i>	11.918	4.916
β	106.9°	
<i>Z</i>	4	4
ρ_{obs}^*	1.32	1.32 g.cm ⁻³
ρ_{calc}	1.32	1.34
Experimental		
Scan speed (2 θ /min)	2	4°
Background counting time	15	7.5 sec
Scan range (2 θ)	1.4 + 0.76 tan θ_B	2.0 + 0.7 tan θ_B †
μ (Mo <i>K</i> α)	3.62	3.45 cm ⁻¹
Crystal size	0.20 × 0.15 × 0.35	0.05 × 0.33 × 0.30 mm
Maximum 2 θ_B	55	55°
No. of reflexions	1363	1691

* by the flotation method

† θ_B : Bragg angleTable 2. *Citrulline hydrochloride: the final atomic coordinates (fractional) and thermal parameters ($\times 10^4$) with standard deviations in parentheses*Standard deviations are in units of the last digits. The anisotropic coefficients have the form: $\exp\{-(\beta_{11}h^2 + \beta_{22}k^2 + \beta_{33}l^2 + \beta_{12}hk + \beta_{13}hl + \beta_{23}kl)\}$

	<i>x</i>	<i>y</i>	<i>z</i>	β_{11}	β_{22}	β_{33}	β_{12}	β_{13}	β_{23}
Cl	0.1568 (1)	0.2500 (3)	0.1479 (1)	40 (0)	296 (3)	56 (1)	6 (2)	47 (1)	41 (3)
C(1)	0.1504 (2)	0.6016 (8)	0.4831 (3)	19 (1)	274 (15)	43 (2)	-1 (7)	7 (2)	9 (10)
C(2)	0.1353 (2)	0.8132 (7)	0.3917 (3)	20 (1)	216 (14)	49 (2)	-5 (6)	9 (2)	-1 (9)
C(3)	0.0495 (2)	0.8454 (8)	0.3247 (3)	19 (1)	221 (12)	58 (2)	8 (6)	11 (3)	22 (10)
C(4)	0.0092 (2)	0.6004 (8)	0.2659 (3)	18 (1)	247 (15)	68 (3)	-6 (7)	13 (3)	-17 (11)
C(5)	-0.0716 (2)	0.6725 (8)	0.1895 (3)	19 (1)	333 (17)	62 (3)	-26 (7)	8 (3)	31 (11)
C(6)	-0.1620 (2)	0.3078 (8)	0.1709 (3)	16 (1)	343 (18)	49 (2)	4 (7)	12 (2)	-39 (10)
N(1)	0.1821 (1)	0.7586 (8)	0.3105 (2)	20 (1)	263 (11)	57 (2)	-16 (7)	21 (2)	41 (11)
N(2)	-0.1155 (2)	0.4522 (9)	0.1275 (3)	24 (1)	518 (18)	45 (2)	-70 (8)	20 (2)	-47 (11)
N(3)	-0.2075 (2)	0.1385 (8)	0.0964 (3)	27 (1)	395 (15)	50 (2)	-43 (7)	20 (2)	-65 (10)
O(1)	0.1921 (1)	0.4191 (6)	0.4823 (2)	27 (1)	268 (10)	57 (2)	29 (5)	20 (2)	47 (8)
O(2)	0.1148 (2)	0.6443 (8)	0.5614 (2)	47 (1)	577 (18)	64 (2)	150 (8)	58 (3)	139 (11)
O(3)	-0.1646 (1)	0.3252 (7)	0.2735 (2)	31 (1)	480 (16)	49 (2)	-65 (7)	33 (2)	-85 (9)

The bond distances and angles are shown in Fig. 1. The averages of the standard deviations for the bond distances and angles among the C, N and O atoms are 0.005 Å and 0.3° respectively.

The lengths of the seven C-H bonds are in the range 0.96 to 1.11 Å with an average of 1.02 Å, and the lengths of the six N-H bonds range from 0.90 to 0.99 Å with an average of 0.95 Å. One O-H bond is 1.05 Å long.

As expected, the crystal structure obtained agrees essentially with that reported by Naganathan & Venkatesan (1971). The citrulline molecule consists of three planar groups, the carboxyl group (I), the aliphatic *trans* zigzag chain (II) and the carbamylamino group (III). The equations of the best planes are listed in Table 5, and the relative orientations of these planar

Table 3. *The final parameters of the hydrogen atoms*

	<i>x</i>	<i>y</i>	<i>z</i>	<i>B</i> (Å ²)	Bonded to
H(1)	0.173	0.606	0.268	2.7	N(1)
H(2)	0.176	0.892	0.255	1.8	N(1)
H(3)	0.237	0.775	0.346	2.4	N(1)
H(4)	-0.117	0.456	0.052	1.2	N(2)
H(5)	-0.202	0.112	0.020	2.7	N(3)
H(6)	-0.234	-0.003	0.125	1.8	N(3)
H(7)	0.130	0.518	0.633	5.0	O(2)
H(21)	0.155	0.974	0.431	1.5	C(2)
H(31)	0.041	0.995	0.262	1.0	C(3)
H(32)	0.023	0.918	0.376	0.5	C(3)
H(41)	0.005	0.483	0.326	2.5	C(4)
H(42)	0.043	0.524	0.212	1.0	C(4)
H(51)	-0.101	0.785	0.243	1.6	C(5)
H(52)	-0.068	0.801	0.131	1.2	C(5)

$$\langle\sigma(x)\rangle = 0.04 \quad \langle\sigma(y)\rangle = 0.05 \quad \langle\sigma(z)\rangle = 0.04 \text{ \AA}$$

$$\langle\sigma(B)\rangle = 0.9 \text{ \AA}^2$$

groups are shown in Fig. 5. These three planes are found to intersect roughly perpendicularly to each other; the dihedral angles are 88° for I and II, 77° for I and III, and 90° for II and III.

Since the hydrogen bond scheme is the same as that given in Fig. 2 of Naganathan & Venkatesan (1971), the packing diagram is omitted here. The H(3) atom of the α -amino group participates in the bifurcated hydrogen bond, in which the two acceptor atoms are located fairly symmetrically with respect to the $N^\alpha-H(3)$ bond. Some of the details around the hydrogen bonds are shown in Fig. 2.

Table 4. *Citrulline hydrochloride*:
observed and calculated structure factors

h	k	l	F _o			F _c			R	R _w
			F _o	h ² +k ² +l ²	hkl	F _c	h ² +k ² +l ²	hkl		
0	0	0	21.15	0	0	21.15	0	0.00	0.00	
1	0	0	24.46	0	0	24.46	0	0.00	0.00	
2	0	0	18.95	0	0	18.95	0	0.00	0.00	
3	0	0	15.24	0	0	15.24	0	0.00	0.00	
4	0	0	12.11	0	0	12.11	0	0.00	0.00	
5	0	0	9.47	0	0	9.47	0	0.00	0.00	
6	0	0	7.21	0	0	7.21	0	0.00	0.00	
7	0	0	5.35	0	0	5.35	0	0.00	0.00	
8	0	0	3.88	0	0	3.88	0	0.00	0.00	
9	0	0	2.79	0	0	2.79	0	0.00	0.00	
10	0	0	2.06	0	0	2.06	0	0.00	0.00	
11	0	0	1.60	0	0	1.60	0	0.00	0.00	
12	0	0	1.28	0	0	1.28	0	0.00	0.00	
13	0	0	1.02	0	0	1.02	0	0.00	0.00	
14	0	0	0.81	0	0	0.81	0	0.00	0.00	
15	0	0	0.64	0	0	0.64	0	0.00	0.00	
16	0	0	0.50	0	0	0.50	0	0.00	0.00	
17	0	0	0.39	0	0	0.39	0	0.00	0.00	
18	0	0	0.30	0	0	0.30	0	0.00	0.00	
19	0	0	0.23	0	0	0.23	0	0.00	0.00	
20	0	0	0.18	0	0	0.18	0	0.00	0.00	
21	0	0	0.14	0	0	0.14	0	0.00	0.00	
22	0	0	0.11	0	0	0.11	0	0.00	0.00	
23	0	0	0.08	0	0	0.08	0	0.00	0.00	
24	0	0	0.06	0	0	0.06	0	0.00	0.00	
25	0	0	0.04	0	0	0.04	0	0.00	0.00	
26	0	0	0.03	0	0	0.03	0	0.00	0.00	
27	0	0	0.02	0	0	0.02	0	0.00	0.00	
28	0	0	0.01	0	0	0.01	0	0.00	0.00	
29	0	0	0.01	0	0	0.01	0	0.00	0.00	
30	0	0	0.01	0	0	0.01	0	0.00	0.00	
31	0	0	0.01	0	0	0.01	0	0.00	0.00	
32	0	0	0.01	0	0	0.01	0	0.00	0.00	
33	0	0	0.01	0	0	0.01	0	0.00	0.00	
34	0	0	0.01	0	0	0.01	0	0.00	0.00	
35	0	0	0.01	0	0	0.01	0	0.00	0.00	
36	0	0	0.01	0	0	0.01	0	0.00	0.00	
37	0	0	0.01	0	0	0.01	0	0.00	0.00	
38	0	0	0.01	0	0	0.01	0	0.00	0.00	
39	0	0	0.01	0	0	0.01	0	0.00	0.00	
40	0	0	0.01	0	0	0.01	0	0.00	0.00	
41	0	0	0.01	0	0	0.01	0	0.00	0.00	
42	0	0	0.01	0	0	0.01	0	0.00	0.00	
43	0	0	0.01	0	0	0.01	0	0.00	0.00	
44	0	0	0.01	0	0	0.01	0	0.00	0.00	
45	0	0	0.01	0	0	0.01	0	0.00	0.00	
46	0	0	0.01	0	0	0.01	0	0.00	0.00	
47	0	0	0.01	0	0	0.01	0	0.00	0.00	
48	0	0	0.01	0	0	0.01	0	0.00	0.00	
49	0	0	0.01	0	0	0.01	0	0.00	0.00	
50	0	0	0.01	0	0	0.01	0	0.00	0.00	
51	0	0	0.01	0	0	0.01	0	0.00	0.00	
52	0	0	0.01	0	0	0.01	0	0.00	0.00	
53	0	0	0.01	0	0	0.01	0	0.00	0.00	
54	0	0	0.01	0	0	0.01	0	0.00	0.00	
55	0	0	0.01	0	0	0.01	0	0.00	0.00	
56	0	0	0.01	0	0	0.01	0	0.00	0.00	
57	0	0	0.01	0	0	0.01	0	0.00	0.00	
58	0	0	0.01	0	0	0.01	0	0.00	0.00	
59	0	0	0.01	0	0	0.01	0	0.00	0.00	
60	0	0	0.01	0	0	0.01	0	0.00	0.00	
61	0	0	0.01	0	0	0.01	0	0.00	0.00	
62	0	0	0.01	0	0	0.01	0	0.00	0.00	
63	0	0	0.01	0	0	0.01	0	0.00	0.00	
64	0	0	0.01	0	0	0.01	0	0.00	0.00	
65	0	0	0.01	0	0	0.01	0	0.00	0.00	
66	0	0	0.01	0	0	0.01	0	0.00	0.00	
67	0	0	0.01	0	0	0.01	0	0.00	0.00	
68	0	0	0.01	0	0	0.01	0	0.00	0.00	
69	0	0	0.01	0	0	0.01	0	0.00	0.00	
70	0	0	0.01	0	0	0.01	0	0.00	0.00	
71	0	0	0.01	0	0	0.01	0	0.00	0.00	
72	0	0	0.01	0	0	0.01	0	0.00	0.00	
73	0	0	0.01	0	0	0.01	0	0.00	0.00	
74	0	0	0.01	0	0	0.01	0	0.00	0.00	
75	0	0	0.01	0	0	0.01	0	0.00	0.00	
76	0	0	0.01	0	0	0.01	0	0.00	0.00	
77	0	0	0.01	0	0	0.01	0	0.00	0.00	
78	0	0	0.01	0	0	0.01	0	0.00	0.00	
79	0	0	0.01	0	0	0.01	0	0.00	0.00	
80	0	0	0.01	0	0	0.01	0	0.00	0.00	
81	0	0	0.01	0	0	0.01	0	0.00	0.00	
82	0	0	0.01	0	0	0.01	0	0.00	0.00	
83	0	0	0.01	0	0	0.01	0	0.00	0.00	
84	0	0	0.01	0	0	0.01	0	0.00	0.00	
85	0	0	0.01	0	0	0.01	0	0.00	0.00	
86	0	0	0.01	0	0	0.01	0	0.00	0.00	
87	0	0	0.01	0	0	0.01	0	0.00	0.00	
88	0	0	0.01	0	0	0.01	0	0.00	0.00	
89	0	0	0.01	0	0	0.01	0	0.00	0.00	
90	0	0	0.01	0	0	0.01	0	0.00	0.00	
91	0	0	0.01	0	0	0.01	0	0.00	0.00	
92	0	0	0.01	0	0	0.01	0	0.00	0.00	
93	0	0	0.01	0	0	0.01	0	0.00	0.00	
94	0	0	0.01	0	0	0.01	0	0.00	0.00	
95	0	0	0.01	0	0	0.01	0	0.00	0.00	
96	0	0	0.01	0	0	0.01	0	0.00	0.00	
97	0	0	0.01	0	0	0.01	0	0.00	0.00	
98	0	0	0.01	0	0	0.01	0	0.00	0.00	
99	0	0	0.01	0	0	0.01	0	0.00	0.00	
100	0	0	0.01	0	0	0.01	0	0.00	0.00	

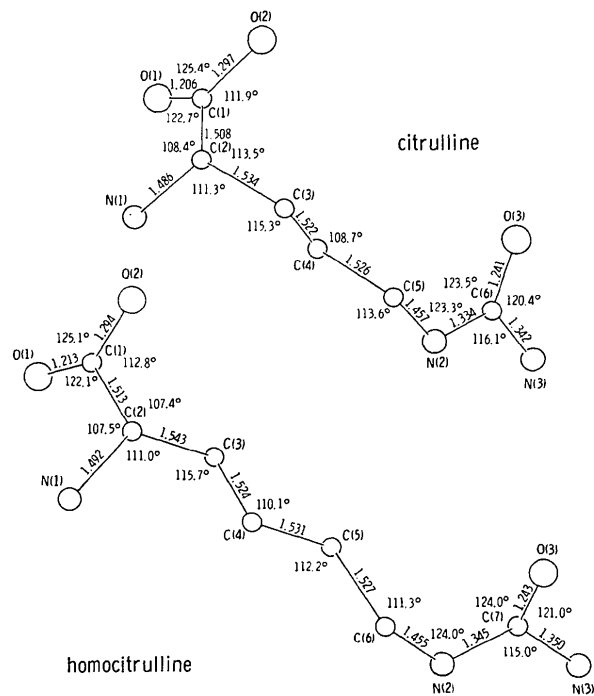


Fig. 1. Bond distances and angles. The averages of the standard deviations are 0.005 \AA and 0.3° for citrulline, and 0.008 \AA and 0.5° for homocitrulline.

The structure of L-homocitrulline hydrochloride

The final R value is 0.069 for all non-zero reflexions, and it is 0.086 if all the reflexions are included. The parameters are listed in Tables 6 and 7, and the observed and calculated structure factors in Table 8. The bond distances and angles are shown in Fig. 1. The averages of the standard deviations for the bond distances and angles are 0.008 \AA and 0.5° respectively.

The nine C-H bond distances range from 0.96 to 1.07 \AA with an average of 1.02 \AA , and the six N-H bonds from 0.85 to 1.02 \AA with an average of 0.91 \AA . One O-H distance is 0.92 \AA .

The homocitrulline molecule also consists of three planar groups, the carboxyl group (I), the *trans* zigzag aliphatic chain (II), and the carbamylamino group (III). The equations of the planes are listed in Table 9. The dihedral angles among the planes are 100° for I and II, 88° for I and III, and 96° for II and III.

The homocitrulline molecule and the chloride ion are held together in the crystal by seven hydrogen bonds. The hydrogen bond scheme is shown in Figs. 3 and 4.

Discussion

Conformation

The conformations of the citrulline and homocitrulline molecules in their hydrochloride crystals are compared in Fig. 5. As mentioned previously, each molecule consists of the three planar groups, the car-

Table 5. *Citrulline hydrochloride: best planes and displacements from the planes*

(a) Equations of the best planes $X = ax + cz \cos \beta$, $Y = by$, $Z = cz \sin \beta$				Atoms				
I†	$0.6553X + 0.5124Y + 0.5550Z =$	5.3359		O(1)	O(2)	C(1)	C(2)	
II†	$0.5641X + 0.2160Y - 0.7970Z =$	-2.1024		N(2)	C(2)	C(3)	C(4)	C(5)
III	$0.6604X - 0.6971Y + 0.2791Z =$	-2.7712		O(3)	N(2)	N(3)	C(6)	

boxyl group, the carbamylamino group and the aliphatic side chain. A fully extended planar conformation of the aliphatic side chain is a common characteristic of the related basic amino acids so far reported. The conformations of the two molecules are, however, different from each other owing to the internal rotation of the $C^\alpha-C^\beta$ bonds. The conformation of the carboxyl group and the aliphatic chain in citrulline is the same as that of arginine found in its dihydrate (Karle & Karle, 1964), while the conformation of the lysine part of homocitrulline is the one which is found in L-lysine.HCl.2H₂O (Wright & Marsh, 1962). The difference in the conformations may be due to the necessity of making up the effective hydrogen bond

Table 5 (cont.)

(b) Displacements (Å) of atoms from the planes									
	I		II		III				
O(1)	0.002	N(2)	0.019	O(3)	-0.001				
O(2)	0.002	C(2)	0.061	N(2)	-0.001				
C(1)	-0.006	C(3)	-0.039	N(3)	-0.001				
C(2)	0.002	C(4)	-0.072	C(6)	0.002				
N(1)*	0.083	C(5)	0.027	C(5)*	-0.211				
C(3)*	-1.202	N(1)*	1.375	H(4)*	-0.102				
H(7)*	0.139	C(1)*	-1.029	H(5)*	0.095				
		C(6)*	-1.095	H(6)*	0.210				

* Atoms with asterisks are not included in the calculation of the best planes.

† In the report of Naganathan & Venkatesan (1971) there seem to be misprints in the signs of coefficients of the equations of the planes II and III.

Table 6. *Homocitrulline hydrochloride: the final atomic coordinates and thermal parameters with standard deviations*

	<i>x</i>	<i>y</i>	<i>z</i>	β_{11}	β_{22}	β_{33}	β_{12}	β_{13}	β_{23}
Cl	0.0695 (1)	0.8285 (1)	0.0238 (2)	81 (1)	16 (0)	356 (5)	2 (1)	-56 (5)	1 (2)
C(1)	1.0819 (5)	1.0182 (2)	0.3161 (11)	65 (5)	17 (1)	321 (21)	25 (4)	-39 (19)	-18 (8)
C(2)	1.0286 (4)	0.9707 (2)	0.5109 (11)	62 (4)	14 (1)	267 (19)	9 (3)	24 (18)	-1 (8)
C(3)	0.8987 (5)	0.9497 (2)	0.3976 (10)	51 (4)	20 (1)	388 (25)	10 (4)	-36 (18)	-44 (9)
C(4)	0.8316 (5)	0.8992 (2)	0.5588 (12)	57 (4)	22 (1)	368 (25)	17 (4)	-57 (20)	-32 (10)
C(5)	0.7169 (5)	0.8750 (3)	0.3993 (13)	56 (4)	23 (1)	456 (29)	20 (4)	-61 (21)	-8 (11)
C(6)	0.6486 (5)	0.8227 (2)	0.5485 (13)	68 (5)	18 (1)	484 (29)	16 (4)	-94 (22)	-28 (11)
C(7)	0.4358 (5)	0.8172 (2)	0.3359 (12)	57 (4)	16 (1)	432 (24)	16 (4)	-26 (21)	7 (9)
N(1)	1.1201 (4)	0.9180 (2)	0.5210 (10)	56 (3)	17 (1)	332 (19)	6 (3)	-11 (17)	-18 (8)
N(2)	0.5528 (4)	0.7939 (2)	0.3767 (12)	64 (4)	15 (1)	607 (28)	4 (3)	-89 (21)	33 (9)
N(3)	0.3669 (4)	0.7883 (2)	0.1422 (12)	72 (4)	22 (1)	541 (28)	-8 (4)	-93 (21)	98 (10)
O(1)	1.1652 (4)	1.0057 (2)	0.1547 (9)	77 (4)	23 (1)	422 (19)	3 (3)	97 (16)	-46 (8)
O(2)	1.0257 (4)	1.0713 (2)	0.3348 (10)	131 (5)	18 (1)	605 (26)	-11 (4)	224 (22)	-57 (8)
O(3)	0.3909 (3)	0.8613 (2)	0.4663 (10)	76 (3)	22 (1)	587 (24)	-13 (3)	-87 (18)	102 (9)

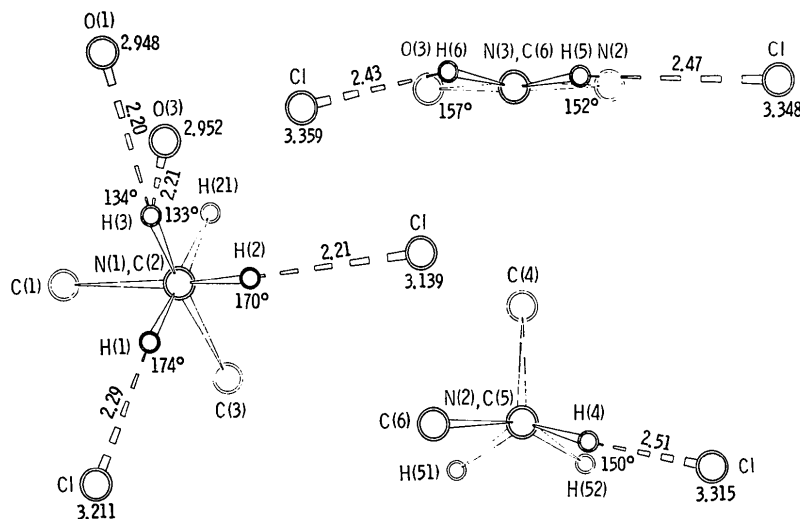


Fig. 2. Citrulline hydrochloride: The environments of the hydrogen bonds viewed down the N-C bonds. The donor-acceptor distances are shown near by the acceptor atoms. The H-acceptor distances and the donor-H-acceptor angles are also shown. The H(3) atom participates in a bifurcated hydrogen bond. One more hydrogen bond in the crystal is $O(2) \cdots O(3) = 2.520$ Å, $H(7) \cdots O(3) = 1.49$ Å, $\angle O(2)-H(7) \cdots O(3) = 169^\circ$.

Table 9. *Homocitrulline hydrochloride: best planes and displacements from the planes*

(a) Equations of the best planes		Atoms				
The coordinates (X, Y, Z) are in Å.						
I	$-0.6790X - 0.2794Y - 0.6789Z = -8.8774$	O(1)	O(2)	C(1)	C(2)	
II	$-0.5334X + 0.6811Y + 0.5016Z = -4.8597$	C(2)	C(3)	C(4)	C(5)	C(6)
III	$0.3641X + 0.6317Y - 0.6844Z = -1.9639$	O(3)	N(2)	N(3)	C(7)	

and 1.523 Å, for the C^α-C' and C^α-C^β distances respectively. The average bond distances of the C-C single bonds in the side chains are 1.527 Å for citrulline and 1.531 Å for homocitrulline. These values are in close agreement with those found in the related compounds mentioned above. The shape and the size of the carbamylamino group in both crystals coincide well with those reported for urea (Vaughan & Donohue, 1952) and those for the peptide linkage given by Corey & Pauling (1952). The bond distances of the C(2)-N(2) bonds, 1.455 and 1.457 Å, are in close agreement with an average value of 1.455 Å for the N-C^α bonds in the peptide linkages (Marsh & Donohue, 1967).

Hydrogen bonds

The overall hydrogen bond schemes in the two crystals are very different, resulting in a difference in the packing in the crystals. The 'local' hydrogen bond schemes are, however, essentially the same for both crystals, although there is not a bifurcated hydrogen bond in the homocitrulline crystal (Figs. 3 and 5). All the hydrogen atoms of the carbamylamino nitrogen atoms form hydrogen bonds to the chloride ions, the mean distance of the six hydrogen bonds in both crystals being 3.338 Å. Each α-amino group forms two hydrogen bonds to the chloride ions, the mean distance of the four bonds being 3.175 Å. Thus the hydrogen bond between the α-amino group and the chloride ion is significantly stronger than the one between the carbamylamino group and the chloride ion. It is interesting to remember that the hydrogen bonds in urea are 'weak', the distances of the two N-H...O bonds being 2.99 and 3.04 Å (Vaughan & Donohue, 1952). The lengthening of these hydrogen bonds is due to the smaller formal positive charge on the nitrogen atoms in the carbamylamino group and in urea.

The other hydrogen atom of the α-amino group forms a bond of 2.873 Å to the oxygen atom of the carboxyl group in homocitrulline, while in citrulline it forms a bifurcated bond to an oxygen atom of the carboxyl group (2.948 Å) and to the oxygen atom of the carbamylamino group (2.952 Å). The donor-acceptor distance in a simple hydrogen bond is shorter than that in a bifurcated one.

In the crystal there is one more hydrogen bond of the type O-H...O, which connects the carboxyl group with the carbamylamino group of the neighbouring molecules, the distance being 2.520 Å in citrulline and 2.485 Å in homocitrulline. Both the hydrogen bonds are fairly strong.

Table 9 (cont.)

(b) Displacements (Å) of atoms from the planes					
I		II		III	
O(1)	0.005	C(2)	-0.083	O(3)	-0.002
O(2)	0.005	C(3)	0.058	N(2)	-0.002
C(1)	-0.014	C(4)	0.087	N(3)	-0.002
C(2)	0.004	C(5)	-0.019	C(7)	0.006
N(1)*	-0.365	C(6)	-0.039	C(6)*	0.180
H(7)*	0.085	N(2)*	-0.349	H(4)*	-0.123
		C(1)*	-0.163	H(5)*	0.109
				H(6)*	-0.019

* Atoms with asterisks are not included in the calculation of the best planes.

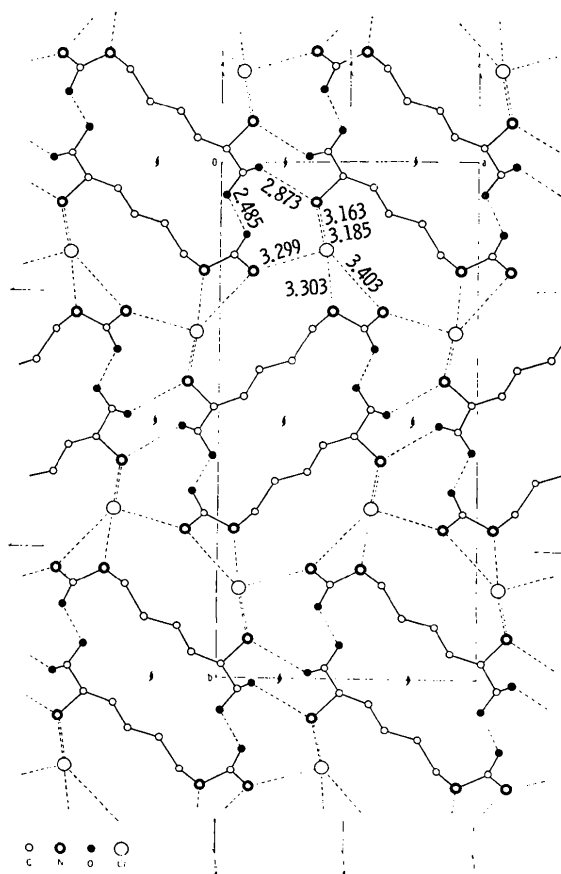


Fig. 3. Homocitrulline hydrochloride: the structure seen along the *c* axis, the *c* axis downwards. Hydrogen bonds are shown by broken lines.

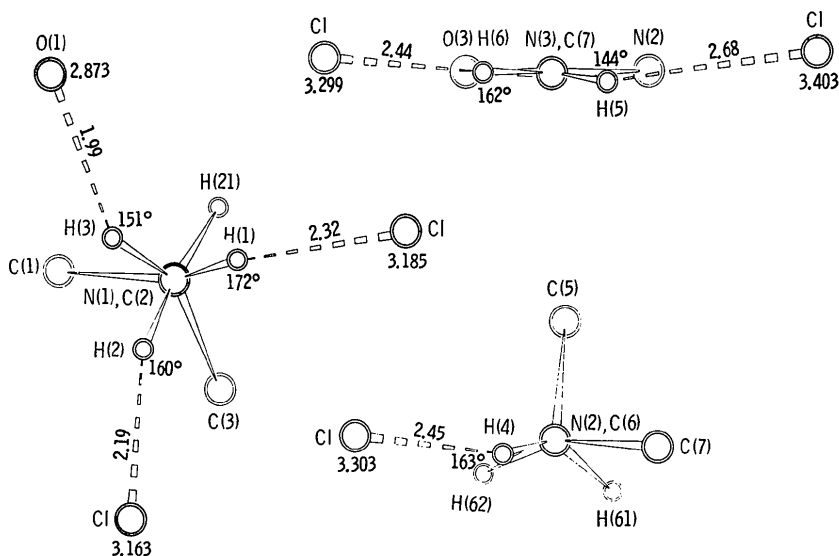


Fig. 4. Homocitrulline hydrochloride: The environments of the hydrogen bonds. One more hydrogen bond is $O(2)-H(7)\cdots O(3)$; $O(2)\cdots O(3)=2.485$ Å, $H(7)\cdots O(3)=1.57$ Å, $\angle O(2)-H(7)\cdots O(3)=180^\circ$.

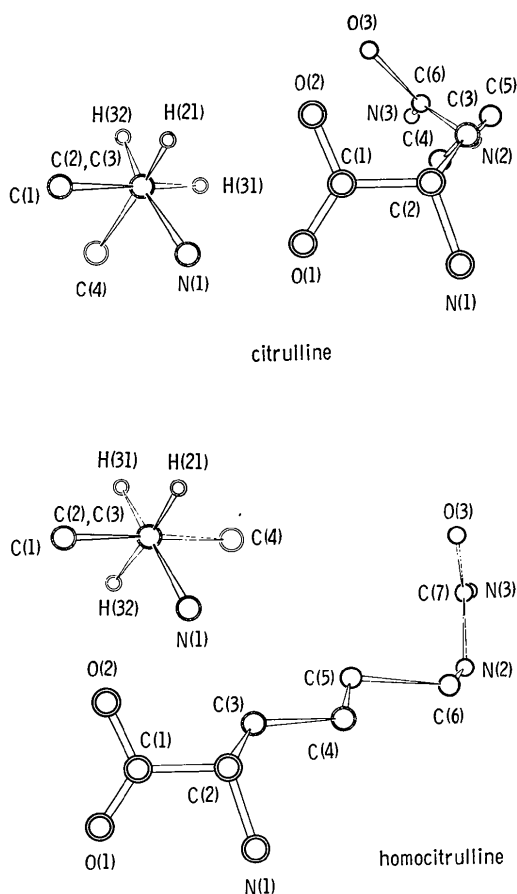


Fig. 5. A comparison of the conformations of citrulline and homocitrulline molecules in their hydrochlorides. One is a view of the molecule projected onto the plane of $C'-C_\alpha-N_\alpha$, while the another shows the internal rotation of the $C_\alpha-C_\beta$ bond.

Comments on the two citrulline hydrochloride structures

The crystal structure of L-citrulline hydrochloride reported by Naganathan & Venkatesan (1971) shows systematic variations from the present result. Their a , b and c are all about 1.2% smaller than ours. They found some of the bond distances to be shorter than usual: for example, $C(1)-O(2)$, 1.251 Å (1.297 Å); $C(2)-N(1)$, 1.460 (1.486); $C(3)-C(4)$, 1.491 (1.522); $C(5)-N(2)$, 1.414 (1.457); $N(1)-H(2)\cdots Cl$, 3.074 (3.139), where the values in the parentheses are the present ones. These distances in our result agree well with those usually found in the related compounds. The shorter bond distances obtained by N & V seem to be mainly attributable to their small unit cell dimensions.

The x and z parameters in both the results show a satisfactory agreement, while the parameters such as y , β_{22} , β_{12} and β_{23} show fairly large discrepancies. The largest deviation is found with respect to $N(2)$, for which y differs by 9σ , β_{22} by 6σ and β_{12} by 16σ (σ being those in our result). The systematic discrepancies are mostly found in the y parameters, so that all the C, N and O atoms are unidirectionally displaced with a mean displacement $\langle \Delta y \rangle = 0.0052$ (ranging from 0.0014 to 0.0084 with the same sign, 0.0052 corresponding to 6σ).

At the refinement stage, N & V used the full-matrix least-squares procedure. In order to make a comparison appropriate, therefore, a full-matrix least-squares refinement procedure was also applied to our final result. However, shifts exceeding $\sigma/4$ did not occur. In view of this fact, it is likely that the discrepancies in parameters found between the two results are mainly due to the differences in the intensity measurement and data processing. Incidentally, N & V's work was done

using Weissenberg photographs around one axis, with scaling by layerwise Wilson plots.

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The Crystal and Molecular Structures of the 1:1 Addition Compounds of Piperidine with *p*-Bromobenzoic Acid and *p*-Chlorobenzoic Acid

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The structures of the two 1:1 addition compounds of piperidine with *p*-bromobenzoic acid, $\text{BrC}_6\text{H}_4\text{CO}_2\text{H} \cdot \text{C}_5\text{H}_{11}\text{N}$, and with *p*-chlorobenzoic acid, $\text{ClC}_6\text{H}_4\text{CO}_2\text{H} \cdot \text{C}_5\text{H}_{11}\text{N}$, have been determined from three-dimensional X-ray diffraction data. Both compounds crystallize in the orthorhombic space group *Pbca* with eight formula units in the unit cell. The cell dimensions are: $a = 10.70$, $b = 27.30$, $c = 9.04$ Å for the *p*-bromo compound, and $a = 10.68$, $b = 26.37$, $c = 9.01$ Å for the *p*-chloro compound. The structures were refined by the block-diagonal least-squares method, with anisotropic temperature factors. The final *R* values for observed reflexions are 0.086 and 0.123 for the *p*-bromo and *p*-chloro compounds respectively. Both structures are isotypic. The structure is composed of an alternate sequence of acid and base moieties held together by $\text{N}^+ - \text{H} \cdots \text{O}^-$ hydrogen bonds. The $\text{N}^+ - \text{H} \cdots \text{O}^-$ distance ranges from 2.67 to 2.76 Å. The hydrogen bonds form an infinite chain around the twofold screw axis parallel to the *c* axis. These molecular chains are packed intimately by van der Waals forces to make up a whole crystal. In either compound, the piperidine ring takes a normal chair form, and the carboxyl group is twisted out of the plane of the benzene ring at an angle of about 8°.

Introduction

The formation of the solid 1:1 and 2:1 addition compounds of benzoic acid and piperidine and those of some *p*-substituted benzoic acids and piperidine have previously been reported (Kashino, 1967; Kashino, Kanei & Hasegawa, 1971). The infrared spectra of the 1:1 compounds were very similar to those of the corresponding sodium benzoates. Thus, it was concluded that the acid moiety in the 1:1 compounds is almost fully ionized. However, the symmetric carboxylate stretching of the adducts showed significantly lower values of $1385\text{--}1375\text{ cm}^{-1}$ compared with the values of $1427\text{--}1406\text{ cm}^{-1}$ for the corresponding sodium benzoates.

It may be desirable to obtain information about the spatial configuration of individual molecules in these crystals and more information about the nature of the bond between the acid and the base components. As a

first step, the crystal structure determinations of the present compounds were undertaken.

Experimental

The 1:1 addition compound of *p*-bromobenzoic acid and piperidine was prepared by dissolving equimolar quantities of *p*-bromobenzoic acid and piperidine in dry benzene. The compound crystallized as transparent tabular plates having well developed {010}. As the compound was hygroscopic and gradually decomposed to its acid and base components, the specimens cut from larger crystals were sealed in glass capillaries. Two different crystals with approximate cross-sections of 0.12×0.18 mm and 0.10×0.15 mm were used for the X-ray analysis, the former for *a* axis photographs and the latter for *c* axis photographs. Using $\text{Cu K}\alpha$ radiation ($\lambda = 1.5418$ Å), multiple-film equi-inclination Weissenberg photographs were taken for the layers from