

Structural Studies of Tryptophan Metabolites by X-ray Diffraction Method. II. The Crystal and Molecular Structure of Tryptamine Hydrochloride

Akio WAKAHARA, Takaji FUJIWARA, and Ken-ichi TOMITA

Faculty of Pharmaceutical Sciences, Osaka University, Toneyama, Toyonaka, Osaka 560

(Received December 18, 1972)

The crystal structure of tryptamine hydrochloride has been solved by the application of the symbolic addition method, and refined by successive Fourier syntheses and least-squares procedures. The crystals are orthorhombic, space group *Pbca*, with lattice parameters $a=8.545$, $b=10.025$ and $c=24.252$ Å. All the positions of hydrogen atoms were found in the difference Fourier map and included in the refinement. The ethylamine side chain bound to the indole ring is in a folded configuration as found in the crystal structure of serotonin picrate monohydrate. A weakly basic nitrogen atom of the indole ring takes part in the hydrogen bonding with the adjacent ion. Every chloride ion behaves as an acceptor for four hydrogen bonds. This is in line with the absence of any marked cleavage where indole rings are packed together to form a pleated sheet and a three-dimensional network structure is essentially stabilized by the hydrogen bond formation.

Tryptamine, a tryptophan metabolite, was first isolated from shoots and flowers of *Acacia* species.¹⁾ A significant biological implication of this compound lies in its role as starting material for the biosynthesis of alkaloids containing an indole ring. It has been also reported to have anti-radiation ability²⁻⁴⁾ as well as serotonin and 5-hydroxytryptophan. In order to study the correlation between the molecular structure and its pharmacodynamic action, it is necessary to elucidate the stereoconfiguration of the molecule. In the course of studies on tryptamine hydrochloride, we found that the aminoethyl side chain of this molecule is in folded configuration.⁵⁾ Extended conformation was reported in the case of a serotonin-creatinine sulphate complex.⁶⁾ The same folded configuration was found in the crystal structure of serotonin picrate monohydrate.⁷⁾ The present paper gives a more detailed account of the methods and the results of analysis. The structure is compared with that of other tryptophan metabolites, some structural features of radiation protective agents being described.

Experimental

Crystals were prepared by slow evaporation from an aqueous solution, and obtained as transparent yellow prismatic crystals. Preliminary rotation and Weissenberg photographs showed them to be orthorhombic, the space group being uniquely determined as *Pbca* from the absent spectra. The unit-cell dimensions were finally determined with a Rigaku Denki manual four-circle diffractometer at room temperature. Melting point of this crystal measured with a DSC (differential scanning calorimeter) is in agreement with the reference value.⁸⁾ The density was measured by the flotation method in a benzene-carbon tetrachloride mixture. It was confirmed that no solvent of crystalliza-

TABLE 1. CRYSTAL DATA OF TRYPTAMINE HYDROCHLORIDE

$C_{10}H_{12}N_2 \cdot HCl$, M.W. 196.68, mp 256—257°C	
Yellowish prism-shaped crystals. Orthorhombic	
$a=8.545(9)$, $b=10.025(8)$, $c=24.252(16)$ Å	
$V=2077.5$ Å ³ , $Z=8$, $F(000)=832$	
Absorption coefficient for X-rays ($\lambda=1.5418$ Å):	
$\mu=29.2$ cm ⁻¹ , $D_m=1.261$ g·cm ⁻³ , $D_x=1.258$ g·cm ⁻³	
(0kl): $k=2n+1$	
Absent spectra (h0l): $l=2n+1$	
(hk0): $h=2n+1$	
Space group; <i>Pbca</i>	

tion is contained in the crystal, the result coinciding with DSC results. The physical and crystal data obtained are given in Table 1.

Intensities of 875 independent non-zero reflections were measured visually on equi-inclination Weissenberg photographs of 0 to 7th layers around the b-axis, with the use of Ni-filtered $CuK\alpha$ radiation. Corrections were made for the Lorentz-polarization factors and spot-shape. No absorption effect was taken into account, since the crystal used was sufficiently small. The data were then adjusted to the absolute scale by means of Wilson's method. The overall temperature factor obtained at this stage was 3.35 Å². The SIGMA program was utilized to list the E -values and Σ_2 relationships with their probabilities.

Determination and Refinement of Structure

The structure of tryptamine hydrochloride was successfully solved by using the direct method. Reflections with $|E|>1.3$ were selected, five of them being given a sign or a letter symbol as follows.

h	k	l	phase	$ E $
2	2	3	+	3.11
5	4	16	+	2.77
2	5	14	+	2.69
5	2	3	A	2.81
6	4	8	B	3.10

Application of Σ_1 formula to reflection (446) indicated that it has a positive sign of 88% probability. It is also included in the initial starting set. The sign

- 1) E. P. White, *New Zealand J. Sci. Tech.*, **25B**, 137 (1944).
- 2) H. Langendorff and R. Koch, *Strahlentherapie*, **98**, 245 (1955).
- 3) P. Alexander and Z. M. Bacq, *Radiat. Res.*, **2**, 292 (1955).
- 4) I. I. Sapezhinskii and N. M. Emanuel, *Dokl. Akad. Nauk. SSSR*, **132**, 1441 (1960).
- 5) A. Wakahara, T. Fujiwara, and K. Tomita, *Tetrahedron Lett.*, **1970**, 4999.
- 6) I. L. Karle, K. S. Dragonette, and S. A. Brenner, *Acta Crystallogr.*, **19**, 713 (1965).
- 7) U. Thewalt and C. E. Bugg, *ibid.*, **B28**, 82 (1972).
- 8) A. H. Jackson and A. E. Smith, *J. Chem. Soc.*, **1965**, 3498.

relations between a number of reflections strongly suggested $A=B=+$. At the end of the process the phases of 157 independent reflections were determined, and a Fourier synthesis was calculated with normalized structure factors as the coefficients (E -map). The three dimensional E -map showed the sites of all 13 atoms of this molecule. Another 20 spurious peaks of the same height as that of light atoms appeared. The indole ring plane enables us to distinguish true peaks from false ones. The structure was refined by the block-diagonal-matrix least-squares method, first with isotropic thermal parameters and then with anisotropic temperature factors. The estimated parameters of hydrogen atoms were computed in such a way as to give bond angles as close to trigonal or tetrahedral as possible. The distances were assumed to be 1.08 Å for C-H, 1.01 Å for N-H, and 0.97 Å for O-H. The amino nitrogen atom had to be the protonated form, and three hydrogen atoms were placed on it. This nitrogen atom has three chloride neighbors at reasonable hydrogen-bonding distances and angles. Thus, three hydrogen atoms were placed at proper positions on the $N\cdots Cl^-$ line. In order to check the expected positions of hydrogen atoms, a difference Fourier synthesis was prepared and all the hydrogen peaks were obtained near the calculated positions as shown in Fig. 1. This may indicate the correctness of the assumed positions of hydrogen atoms and the assignment of hydrogen bonding system. More refinement cycles were performed, which included refinement of the positional and overall isotropic thermal parameters of the hydrogen atoms. Refinement was stopped when parameter shifts were less than one fourth of their standard deviations, giving an R value of 0.094. The N-H and C-H bond lengths are in the ranges 0.99–1.11 Å and 1.03–1.17 Å, respectively. The final electron density map is shown in Fig. 2. The final observed and calculated structure factor amplitudes are given in Table 2 and final atomic positional and thermal parameters in Table 3. The scattering factors used throughout the refinement were taken from *International Tables for X-ray Crystallography* (1962). All calculations were done on a NEAC 2200–500 of this University, using programs written by T. Ashida and the authors.

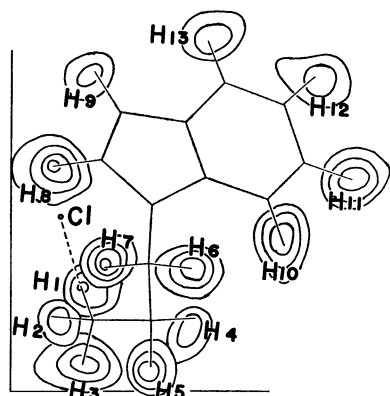


Fig. 1. The location of hydrogen atoms from a three-dimensional difference Fourier map. Contours from 0.2 $e\cdot\text{\AA}^{-3}$ at intervals of 0.2 $e\cdot\text{\AA}^{-3}$.

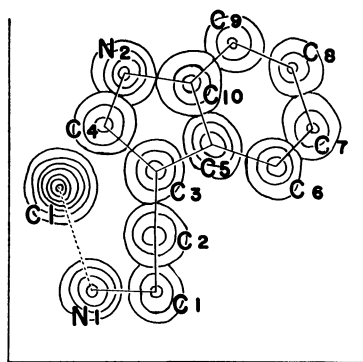


Fig. 2. A composite drawing of the final electron density map, viewed along a -axis. Contour intervals of 2.0 $e\cdot\text{\AA}^{-3}$ for light atoms and 4.0 $e\cdot\text{\AA}^{-3}$ for the chloride ion, the first contour begins at 2.0 $e\cdot\text{\AA}^{-3}$.

Description and Discussion of the Structure

Molecular Dimensions. The bond lengths and angles are shown in Fig. 3 and Table 4 together with those of the other tryptophan metabolites. The numbering of other metabolites corresponds to that of tryptamine in Fig. 3. We see some interesting structural features of radiation protective agents containing indole ring (indicated with asterisk in Table 4). The C(3)–C(4) bond, 1.342–1.387 Å, is apparently shorter than a normal aromatic C–C bond (1.395 Å). The corresponding bond of four radiation protectors is longer than that of other indole derivatives, and is compatible to 1.381 Å of yohimbane hydrobromide⁹ and 1.395 Å of reserpine.¹⁰ On the other hand, the C(2)–C(3) distance of the protector is slightly shorter than that of the other related compounds. The corresponding length of yohimbane and reserpine is 1.489 Å and 1.444 Å, respectively. It is of biological in-

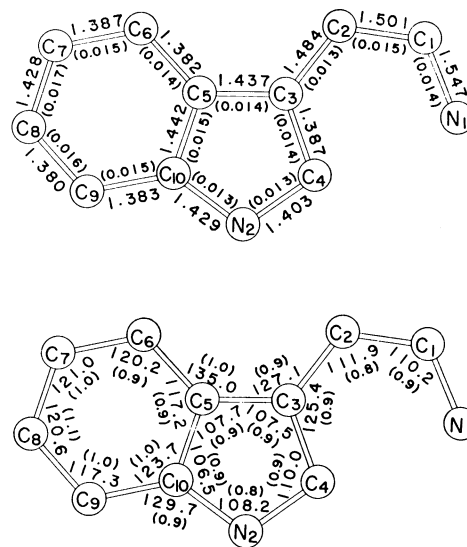


Fig. 3. Bond lengths (Å) and valency angles (°). The standard deviations are given in parentheses.

- 9) J. P. Fennessey and W. Nowacki, *Z. Kristallogr.*, **131**, 342 (1970).
- 10) I. L. Karle and J. Karle, *Acta Crystallogr.*, **B24**, 81 (1968).

L	FO	FC	L	FO	FC	L	FO	FC	L	FO	FC	L	FO	FC	L	FO	FC	L	FO	FC	L	FO	FC	L	FO	FC	L	FO	FC
H,K=	0	0	2	117	111	12	16	-15	11	66	-61	24	22	25	13	22	-22	12	27	-34	9	30	28	8	45	-40	-40	-40	-40
4	123	-121	2	124	-120	14	37	-38	13	20	-22	25	27	-25	15	29	-35	14	20	-25	11	35	-30	11	38	-36	-36	-36	
6	233	-241	3	233	-268	H,K=	9	1	17	20	-22	27	13	-13	16	40	-40	H,K=	1	4	16	36	33	13	41	-38	-38	-38	
10	173	-19	5	19	-26	2	27	28	8	9	19	20	6	-9	19	20	6	-9	19	20	6	-9	19	20	6	-9	19	20	
10	173	-174	3	21	23	21	23	21	29	29	H,K=	4	3	18	10	-31	8	12	-16	H,K=	8	5	15	21	-17	-17	-17	-17	
12	79	-80	8	93	97	6	30	-32	H,K=	6	2	0	8	10	20	35	-39	H,K=	10	4	0	30	-32	H,K=	8	6	-6	-6	
14	124	-116	9	42	39	8	58	-51	0	48	-42	1	21	17	22	25	-25	5	15	16	4	24	-25	0	24	-24	-24		
16	33	-30	10	40	-35	15	17	-21	2	49	-41	2	19	20	25	21	20	H,K=	1	5	6	17	-15	1	10	-14	-14		
18	40	-39	11	73	71	H,K=	10	1	3	47	-40	3	64	70	H,K=	2	4	1	35	-39	8	26	33	2	35	-36	-36		
20	20	-17	12	20	0	54	-54	4	16	16	4	142	163	12	16	68	-7	2	16	1	34	34	8	26	34	8	26		
24	51	50	15	122	119	3	23	-24	5	56	-58	5	118	118	1	40	-39	3	68	78	11	34	35	9	16	-35	-35		
26	66	58	14	50	46	5	56	-37	7	65	-92	7	27	33	2	28	23	6	67	-77	14	22	21	10	10	6	6		
28	27	25	15	17	-19	H,K=	0	2	9	84	-81	8	76	82	3	6	-2	8	69	-79	16	10	13	H,K=	9	6	9	6	
H,K=	1	0	17	45	-36	2	143	-144	12	42	36	9	74	62	4	127	118	9	49	-56	H,K=	9	5	9	25	25	25	25	
2	74	-64	18	65	-56	3	44	38	13	20	23	10	44	5	53	-63	10	12	11	4	24	25	18	1	9	-7	-7		
4	20	-24	20	99	-89	4	34	38	17	46	48	13	57	58	64	125	27	6	23	28	6	23	28	1	9	-7	-7		
6	370	-371	22	28	30	5	144	-149	19	44	13	12	16	16	7	63	-61	15	16	10	1	17	2	9	9	9	9		
8	196	-196	H,K=	2	1	6	62	-65	23	13	15	13	10	9	8	49	-46	16	37	39	12	24	2	4	28	-37	-37	-37	
10	25	-22	0	384	331	7	176	-179	H,K=	7	2	15	19	-20	9	26	-29	18	21	28	H,K=	0	6	5	29	27	27	27	
12	50	-51	1	136	-146	8	23	16	1	57	-48	16	27	27	10	72	-68	19	12	-8	2	106	104	9	26	27	27		
14	25	-25	2	87	98	9	149	-160	2	10	-11	17	43	-47	14	33	-29	20	16	19	3	25	28	10	58	63	63		
16	62	55	3	289	279	10	41	-45	3	57	-38	19	45	-47	15	39	-18	23	30	24	4	86	96	11	34	-11	-11		
20	82	82	5	143	130	12	101	120	6	35	26	22	30	-30	16	33	-34	26	9	-13	5	26	27	14	38	-36	-36		
22	13	-13	6	46	38	13	21	20	8	23	23	23	41	-34	18	33	-39	0	91	-89	8	76	-83	H,K=	2	7	7		
30	12	-12	7	141	132	14	44	50	11	51	60	H,K=	5	3	19	27	-28	1	36	33	9	65	69	0	32	-26	-26		
H,K=	2	0	8	54	50	15	35	37	13	54	61	1	45	40	22	33	36	2	36	-82	10	68	-78	1	18	16	16		
0	196	212	9	18	-20	16	13	-14	15	37	37	3	43	43	3	36	-3	7	-9	-11	7	70	2	31	24	-32	-32		
4	219	-256	11	28	32	18	29	-33	23	16	-17	4	78	74	H,K=	3	4	6	33	-29	13	41	-40	4	28	28	28		
6	87	-80	12	122	-106	19	52	53	H,K=	8	2	5	19	17	1	34	33	8	29	24	14	36	-41	5	92	77	77		
8	36	-33	13	55	-49	21	31	33	0	19	20	6	15	15	3	39	-29	9	37	-41	16	20	19	6	29	26	26		
10	65	61	14	140	-120	23	70	65	1	13	-16	7	36	31	4	32	-28	10	42	42	17	36	-40	7	62	54	54		
12	91	81	15	22	-22	29	18	-14	3	24	24	59	-59	39	-49	59	60	18	52	-53	8	59	59	5	94	-53	-53		
14	112	109	15	22	-22	29	18	-14	3	24	24	59	-59	39	-49	59	60	18	52	-53	8	59	59	5	94	-53	-53		
18	41	36	17	24	-25	1	102	-87	7	20	19	10	46	-37	7	40	42	15	36	45	21	32	-33	13	32	-31	-31		
22	52	-50	21	28	-27	2	31	-26	11	45	45	11	44	-37	8	22	-21	16	16	20	H,K=	1	6	14	33	28	28		
26	13	-9	23	24	-21	3	185	-219	14	26	-31	12	25	27	10	75	-77	22	21	-24	1	66	63	16	26	-25	-25		
H,K=	3	0	26	33	42	4	64	62	17	14	-14	13	51	-48	12	46	-51	23	11	13	2	64	-57	17	20	-32	-32		
2	10	9	H,K=	3	5	35	-32	19	23	-23	13	39	41	40	-24	41	-40	3	3	5	4	71	-61	H,K=	3	7	7		
4	61	-61	168	147	11	145	11	-15	3	28	28	17	20	-21	H,K=	4	4	1	42	-34	6	43	-39	1	23	25	25		
8	42	-42	3	28	29	8	57	56	H,K=	10	2	20	21	-23	0	32	-27	2	122	129	8	59	-60	2	30	-30	-30		
10	80	71	6	189	-226	9	60	56	4	12	-10	25	19	20	1	8	10	4	113	113	9	29	-30	3	89	93	93		
12	95	85	8	120	-127	10	57	47	6	5	4	H,K=	6	3	2	32	31	5	16	14	10	26	-22	4	34	-38	-38		
14	16	-17	9	50	-52	11	93	92	2	170	201	3	30	-31	5	38	-35	6	85	76	8	52	-52	5	31	24	-34		
18	44	-43	11	30	-28	15	57	56	2	7	-6	6	10	-17	6	72	61	10	58	58	13	24	-37	6	29	31	31		
22	36	-37	12	27	24	17	61	59	3	95	92	7	48	-50	7	69	-69	11	17	-19	14	29	30	11	66	-77	-77		
24	43	-45	13	29	-19	23	33	-34	4	61	60	8	41	40	8	27	-25	12	37	35	15	25	-27	12	22	-22	-22		
28	20	24	15	63	-54	25	33	-34	5	80	-67	14	25	-23	9	25	-25	14	49	-48	16	36	36	13	55	-61	-61		
H,K=	4	0	16	60	56	29	12	-11	6	26	-27	H,K=	7	3	30	-24	16	45	-45	17	30	-30	15	28	-35	28	-38	-38	
0	29	-35	17	73	73	H,K=	2	7	13	28	20	9	19	12	32	-32	17	32	31	18	16	1	17	22	22	23	23		
2	36	-31	18	25	31	0	229	204	9	124	-108	4	45	47	13	20	22	18	28	-27	H,K=	2	6	21	19	20	20		
4	58	-50	19	25	20	1	48	-43	11	50	-52	8	35	-40	14	24	-24	19	12	-12	0	67	-68	24	10	7	7		
6	13	-14	20	29	27	2	43	44	12	18	17	11	26	-28	17	32	30	20	39	-28	2	12	-12	H,K=	4	7	7		
8	36	-33	22	29	31	3	290	307	14	35	35	12	24	24	H,K=	5	4	22	29	-36	3	97	-83	0	23	30	30		
10	58	53	4	38	4	130	-121	18	38	-81	14	30	-29	2	49	-43	H,K=	4	5	6	28	-29	3	37	-34	-34			
12	70	69	H,K=	4	115	115	20	83	-28	23	15	21	21	3	57	-52	0	107	112	7	46	-43	4	14	17	-17	-17		
14	57	-60	2	60	-55	7	39	40	24	21	23	H,K=	8	3	5	39	-34	1	12	16	8	15	18	5	64	-61	-61		
26	11	-14	3	41	-31	8	23	-24	25	26	27	0	20	-22	7	40	40	2	80	72	9	20	-22	6	12	-13	-13		
28	19	-20	4	508	-106	9	32	33	27	18	18	2	21	21	8	78	-71	3	20	17	11	22	22	7	31	-34	-34		
H,K=	5	0	5	26	24	11	13	15	29	8	6	3	30	32	10	56	-58	5	21	20	16	50	52	11	44	-45	-45		
4	57	58	7	66	-68	12	22	22	24	0	108	107	7	20	-24	11	29	30	6	24	19	H,K=	3	6	14	26	26		
6	8	59	75	9	70	-82	16	36	-40	1	34	-31	8	20	-20	12	55	-58	7	8	7	2	30	31	15	28	27		
10	17	18	10	60	64	19	67	-58	2	74	63	9	20	-22	13	12	16	8	92	-101	3	23	-18	17	40	39	39		
12	55	53	11																										

- 12) R. A. Pasternak, *ibid.*, **9**, 341 (1956).
- 13) G. Falkenberg and D. Carlström, *ibid.*, **B27**, 411 (1971).

TABLE 3. FINAL POSITIONAL COORDINATES AND THERMAL PARAMETERS WITH THEIR STANDARD DEVIATIONS ($\times 10^4$)
Anisotropic thermal parameters are in the form of $\exp[-(B_{11}h^2 + B_{22}k^2 + B_{33}l^2 + B_{12}hk + B_{13}hl + B_{23}kl)]$.

Atom	x	y	z	B_{11}	B_{22}	B_{33}	B_{12}	B_{13}	B_{23}
Cl	0.1715 (3)	0.6462 (3)	0.0394 (1)	0.0118 (3)	0.0081 (3)	0.0019 (0)	-0.0002 (6)	-0.0025 (2)	-0.0003 (2)
N(1)	0.4492 (8)	0.8548 (9)	0.0646 (3)	0.0091 (11)	0.0126 (13)	0.0019 (2)	0.0026 (20)	0.0006 (7)	-0.0054 (8)
N(2)	0.4851 (9)	0.4124 (7)	0.0908 (3)	0.0152 (13)	0.0039 (11)	0.0019 (2)	0.0090 (18)	-0.0029 (8)	-0.0009 (6)
C(1)	0.5584 (11)	0.8604 (11)	0.1155 (3)	0.0197 (18)	0.0152 (17)	0.0006 (1)	-0.0251 (30)	0.0004 (8)	0.0002 (8)
C(2)	0.6687 (10)	0.7437 (9)	0.1152 (4)	0.0104 (12)	0.0035 (12)	0.0022 (2)	0.0002 (21)	-0.0007 (9)	0.0014 (7)
C(3)	0.5841 (9)	0.6147 (9)	0.1172 (3)	0.0066 (11)	0.0097 (14)	0.0011 (1)	0.0027 (18)	0.0002 (7)	0.0017 (7)
C(4)	0.5750 (9)	0.5229 (10)	0.0745 (4)	0.0074 (12)	0.0106 (15)	0.0020 (2)	0.0028 (20)	-0.0022 (8)	-0.0020 (9)
C(5)	0.4960 (9)	0.5616 (10)	0.1629 (3)	0.0076 (11)	0.0120 (15)	0.0010 (1)	0.0050 (21)	-0.0003 (7)	0.0017 (7)
C(6)	0.4573 (9)	0.6068 (9)	0.2150 (4)	0.0099 (13)	0.0035 (13)	0.0022 (2)	0.0036 (19)	-0.0014 (8)	0.0010 (7)
C(7)	0.3646 (10)	0.5292 (12)	0.2493 (4)	0.0099 (14)	0.0168 (18)	0.0020 (2)	0.0044 (24)	0.0006 (9)	0.0031 (11)
C(8)	0.3033 (12)	0.4042 (12)	0.2310 (4)	0.0145 (18)	0.0166 (20)	0.0025 (2)	0.0002 (29)	0.0017 (10)	0.0063 (11)
C(9)	0.3381 (10)	0.3564 (10)	0.1790 (4)	0.0096 (12)	0.0071 (14)	0.0026 (2)	-0.0017 (22)	-0.0002 (9)	0.0037 (9)
C(10)	0.4302 (9)	0.4359 (11)	0.1456 (4)	0.0086 (12)	0.0109 (15)	0.0018 (2)	0.0045 (22)	-0.0014 (8)	0.0016 (9)
H(1)	0.363	0.793	0.056						
H(2)	0.522	0.863	0.031						
H(3)	0.401	0.956	0.054						
H(4)	0.499	0.866	0.152						
H(5)	0.635	0.950	0.117						
H(6)	0.737	0.763	0.149						
H(7)	0.734	0.753	0.079						
H(8)	0.616	0.540	0.033						
H(9)	0.461	0.340	0.065						
H(10)	0.504	0.703	0.232						
H(11)	0.339	0.555	0.291						
H(12)	0.239	0.348	0.260						
H(13)	0.294	0.249	0.165						

Final thermal parameter for hydrogen atoms: Overall isotropic; $3.3 \text{ \AA}^2$

TABLE 4. THE MOLECULAR DIMENSIONS OF TRYPTOPHAN METABOLITES

	a*	b*	c*	d*	e	f	g
(A) Bond lengths (in Å)							
N(1)–C(1)	1.547	1.491	1.486	1.51	1.505	1.492	
C(1)–C(2)	1.501	1.502	1.513	1.53	1.539	1.532	
C(2)–C(3)	1.484	1.462	1.494	1.48	1.530	1.500	1.514
C(3)–C(4)	1.387	1.374	1.369	1.38	1.344	1.350	1.342
C(3)–C(5)	1.437	1.442	1.439	1.47	1.451	1.435	1.470
C(4)–N(2)	1.403	1.386	1.362	1.39	1.377	1.371	1.401
C(10)–N(2)	1.429	1.381	1.370	1.39	1.391	1.370	1.385
C(5)–C(10)	1.442	1.414	1.407	1.40	1.382	1.412	1.407
C(5)–C(6)	1.382	1.371	1.403	1.41	1.412	1.393	1.434
C(6)–C(7)	1.387	1.375	1.370	1.42	1.397	1.379	1.409
C(7)–C(8)	1.428	1.400	1.401	1.38	1.386	1.400	1.396
C(8)–C(9)	1.380	1.352	1.375	1.40	1.399	1.368	1.409
C(9)–C(10)	1.383	1.378	1.392	1.43	1.400	1.390	1.422
(B) Valency angles (in degree)							
N(1)–C(1)–C(2)	110.2	109.7	110.9	108	109.7	111.4	
C(1)–C(2)–C(3)	111.9	114.9	114.7	108	114.2	112.5	114.7
C(2)–C(3)–C(4)	125.4	128.4	127.8	131	128.0	125.8	127.2
C(2)–C(3)–C(5)	127.1	125.0	126.4	125	126.5	127.5	125.2
C(4)–C(3)–C(5)	107.5	106.6	105.7	104	105.5	106.6	107.1
C(3)–C(4)–N(2)	110.0	110.4	110.7	113	111.5	110.5	110.3
C(4)–N(2)–C(10)	108.2	108.0	108.8	106	107.4	108.7	108.2
N(2)–C(10)–C(5)	106.5	108.5	107.8	110	107.8	107.5	108.2
N(2)–C(10)–C(9)	129.7	130.3	130.8	128	129.0	130.8	128.2
C(5)–C(10)–C(9)	123.7	121.2	121.4	122	123.2	122.0	123.5
C(3)–C(5)–C(10)	107.7	106.6	106.9	107	107.7	106.8	106.1
C(6)–C(5)–C(10)	117.2	120.3	119.3	121	121.1	119.2	119.8
C(3)–C(5)–C(6)	135.0	133.0	133.7	132	131.2	134.2	134.1
C(5)–C(6)–C(7)	120.2	118.5	118.6	117	114.6	118.7	116.7
C(6)–C(7)–C(8)	121.0	119.7	121.7	122	124.8	121.2	122.2
C(7)–C(8)–C(9)	120.6	123.1	120.5	122	119.7	121.8	122.5
C(8)–C(9)–C(10)	117.3	117.0	118.4	116	116.4	117.7	115.2

a) Tryptamine hydrochloride, this work, b) 5-Hydroxy-DL-tryptophan¹⁴⁾, c) Serotonin picrate monohydrate¹⁵⁾, d) Serotonin-creatinine sulfate complex¹⁶⁾, e) L-Tryptophan hydrochloride¹⁶⁾, f) DL-Tryptophan formate¹⁶⁾, g) 3-Indolylacetic acid¹⁷⁾.
* Radiation protective agents.

extent due to the difference in the localization of electron density distribution or the difference in the resonance structure of a five-membered ring of radiation protective agents compared with other indole derivatives.^{18–20)} The mean C–C distance in the benzene ring of tryptamine hydrochloride is 1.400 Å, while that of a pyrrole ring 1.422 Å. In all indole compounds, the C(3)–C(5) bond is evidently longer than the standard aromatic C–C bond.

Configuration. Aminoethyl side chain is in a

folded configuration. The orientation of the C(2)–C(3) bond is *gauche* with respect to C(1)–NH₃⁺ bond, and the torsion angles as defined by Klyne and Prelog²¹⁾ are –60.6° (*syn-clinal*) and +110.9° (*anti-clinal*) for the atoms N(1)–C(1)–C(2)–C(3) and C(1)–C(2)–C(3)–C(4), respectively. From conformational studies on tryptamine derivatives applying normal stereochemical rules, Chothia and Pauling proposed two possible values, 180° (*anti-periplanar*) and ±60° (*syn-clinal*), for the former torsion angle, and 0° (*syn-periplanar*) and ±90° (center of *clinal* region) for the latter one.²²⁾ The configurations of 5-hydroxytryptophan and serotonin picrate resemble that of tryptamine. On the other hand, *anti-periplanar* conformation about C(1)–C(2) bond was observed in a serotonin-creatinine sulfate complex (+172.5°) and 5-methoxy-*N,N*-dimethyltryptamine hydrochloride (+179.2°). The best plane was computed for the atoms in the indole

21) W. Klyne and V. Prelog, *Experientia*, **16**, 521 (1960).

22) C. Chothia and P. Pauling, *Proc. Nat. Acad. Sci. Wash.*, **63**, 1063 (1969).

14) Preceding paper; A. Wakahara, M. Kido, T. Fujiwara, and K. Tomita, *This Bulletin*, **46**, 2475 (1973).

15) T. Takigawa, T. Ashida, Y. Sasada, and M. Kakudo, *ibid.*, **39**, 2369 (1966).

16) E. Bye, A. Mostad, and C. Römmeing, *Acta Chem. Scand.*, **25**, 364 (1971).

17) I. L. Karle, K. Britts, and P. Gum, *Acta Crystallogr.*, **17**, 496 (1964).

18) A. Szent-Gyorgyi and I. Isenberg, *Proc. Nat. Acad. Sci. Wash.*, **46**, 1334 (1960).

19) J. P. Green and J. P. Malrieu, *ibid.*, **54**, 659 (1965).

20) R. Foster and C. A. Fyfe, *J. Chem. Soc., B*, **1966**, 926.

TABLE 5. DISTANCES OF ATOMS FROM THE LEAST-SQUARES PLANE THROUGH THE INDOLE RING (in Å)

Atom	Distance	Atom	Distance
C(3)	0.003	C(4)	0.013
C(5)	-0.022	C(6)	0.006
C(7)	-0.002	C(8)	0.009
C(9)	-0.004	C(10)	0.008
N(2)	-0.009	C(2) ^{a)}	0.024
C(1) ^{a)}	1.324	N(1) ^{a)}	2.498

a) Excluded from calculation of the plane.

ring and expressed by the equation, $0.8146X - 0.4580Y + 0.3559Z = 2.2578$ in rectangular coordinates in Ångström. The indole part of tryptamine molecule is planar, the mean deviation of the atoms from this plane being 0.008 Å and the maximum deviations are +0.0125 and -0.0217 Å for the atoms C(4) and C(5), respectively. The individual deviations of the atoms from this least-squares plane are given in Table 5. The plane formed by N(1), C(1) and C(2) atoms makes a dihedral angle of 83.9° with that of the indole plane. The corresponding angle in serotonin-creatinine sulfate is 12.6°.

The Molecular Environment. The crystal structure viewed along a-axis is given in Fig. 4 and intermolecular distances less than 3.75 Å including hydrogen bonds in Table 6. The side chains of tryptamine are in a folded configuration, and the packing is dominated by a three-dimensional network of hydrogen bonds. The three hydrogen atoms attached to an amino nitrogen atom form three hydrogen bondings with adjacent chloride ions. These three chloride atoms are approximately at three vertices of a regular tetrahedron centered at the N(1) atom, with the C(1) atom directed toward the fourth vertex, the average N-H...Cl

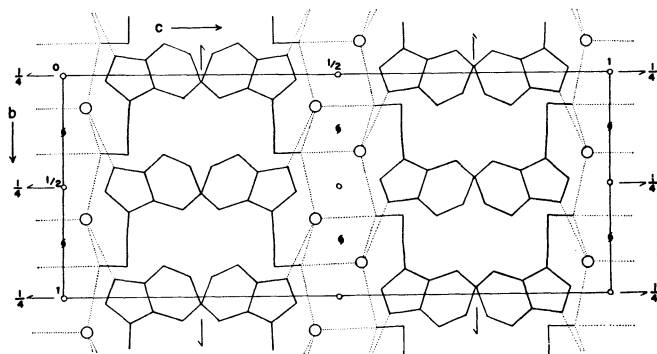


Fig. 4. The crystal structure of tryptamine hydrochloride projected along a-axis.

TABLE 6. INTERMOLECULAR SHORT CONTACTS LESS THAN 3.75 Å

Symmetrycode			
Original molecule	x	y	z
(A)	$-1/2+x$	$3/2-y$	$-z$
(B)	$1/2-x$	$-1/2+y$	z
(C)	$1/2-x$	$1/2+y$	z
(D)	$1/2+x$	y	$1/2-z$
(E)	$1-x$	$-1/2+y$	$1/2-z$
(F)	$3/2-x$	$-1/2+y$	z

(a) Hydrogen bonds			
Donor	Acceptor (Original molecule)	Distance	Angle of C-Donor-Acceptor
Original N(1)-H(1)	Cl	3.221 Å	128.2°
(A) N(1)-H(2)	Cl	3.157	105.9
(B) N(1)-H(3)	Cl	3.158	108.5
(C) N(2)-H(9)	Cl	3.235	111.0

(b) Other short contacts			
Atom (i)	Atom (j)	Distance (d _{ij})	
C(9)	N(1) (B)	3.705 Å	
C(9)	C(1) (B)	3.722	
C(9)	C(6) (B)	3.659	
C(8)	C(6) (B)	3.742	
C(6)	C(7) (D)	3.671	
C(7)	C(1) (E)	3.748	
N(2)	C(2) (F)	3.459	
C(4)	C(2) (F)	3.688	
C(4)	C(1) (F)	3.668	

distance being 3.179 Å. Another hydrogen bond is found between the N(2) atom of indole ring and chloride ion, whose distance is 3.235 Å. As a whole, each chloride atom links with the four neighboring tryptamine cations. N(1)-H...Cl hydrogen bonds form infinite columns around the center of symmetry or along the screw axes, and molecules are arranged in a double layered system parallel to *ab*-plane. In the polar part of the molecules, the hydrogen bond formation is substantial to stabilize the molecular packing. The indole rings are packed by van der Waals forces into the non-polar layer. Unusual short contacts are not observed (Table 6).

The authors wish to thank Drs S. Kobayashi and W. Nakamura, National Institute of Radiological Sciences, Chiba, for discussions and interest throughout the course of this investigation. The work was partly supported by a research grant from the Ministry of Education