

Topochemical Studies. III. The Crystal and Molecular Structures of Crotonic Acid, $\text{CH}_3\text{CH}=\text{CHCOOH}$, and Crotonamide, $\text{CH}_3\text{CH}=\text{CHCONH}_2$

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(Received March 25, 1974)

Crotonic acid crystallizes in the monoclinic space group C2/c , with $a=15.30(3)$, $b=4.06(1)$, $c=16.18(2)$ Å, $\beta=107.4(1)^\circ$ and $Z=8$; crotonamide in the monoclinic space group $\text{P2}_1/\text{a}$, with $a=9.71(2)$, $b=6.90(2)$, $c=7.75(1)$ Å, $\beta=104.0(4)^\circ$ and $Z=4$. The structures have been solved by the Patterson method and refined by the block-diagonal least-squares method. The final R values are 0.090 for 337 observed reflections for crotonic acid, and 0.088 for 525 observed ones for crotonamide. The configuration of the molecules of crotonic acid and crotonamide is approximately planar and of *trans*-type regarding the $\text{C}=\text{C}$ bond. Every two centrosymmetrically related molecules form a dimer through hydrogen bonds of 2.645 Å ($\text{OH}\cdots\text{O}$) for crotonic acid and 2.963 Å ($\text{NH}\cdots\text{O}$) for crotonamide. In the crystals of crotonamide additional $\text{NH}\cdots\text{O}$ hydrogen bonds (2.862 Å) link the neighboring molecules related by the a -glide plane to form a sheet parallel to (001) plane. Distances between the $\text{C}=\text{C}$ bonds of the nearest neighbors are 3.86 Å for crotonic acid and 3.62 and 3.88 Å for crotonamide.

In the first paper of this series¹⁾ attention has been drawn to the crystal structure of aconitic acid as an unsaturated aliphatic acid in order to test the topochemical rule²⁾ and to examine the relationship between the molecular geometry and the hydrogen bond system.³⁾ The present paper is devoted to the examination of the degree of planarity of the molecules and the packing arrangement of the simplest α,β -unsaturated acid, crotonic acid, and its amide.

Experimental

Crotonic acid crystallized from aqueous solution as soft, colorless transparent plates elongated along the b axis and became gradually sublimate. The differential thermogram

showed a weak *endo*-peak at 44 °C and the melting peak at 72 °C. Crotonamide crystallized from ethanol solution as twinned plates or as prisms elongated along the a axis. Crystals of crotonic acid were sealed in glass capillaries. Single crystals of crotonamide were obtained by cutting the twinned plates with razor. Equi-inclination Weissenberg photographs were taken by using $\text{CuK}\alpha$ radiation, $\lambda=1.5418$ Å. Crystal data are summarized in Table 1. The space group for crotonic acid was deduced from the systematic absences to be Cc or C2/c , but the latter was assumed, and confirmed by subsequent refinement. The intensities were estimated visually, and corrected for the Lorentz and polarization factors and for spot shape.

Structure Determination and Refinement

Crotonic Acid. Two-dimensional Patterson and Fourier maps projected along the b axis were used to obtain the trial structure. The y coordinates were estimated by tilting and shifting the planar molecular model against the b axis so that the calculated structure factors may suit to the observed ones. The trial structure gave an R value of 0.27. Three cycles of block-diagonal least-squares refinement of positional and isotropic thermal parameters reduced the R value to 0.15. The reflections $20\bar{4}$, $11\bar{3}$, and $11\bar{2}$, which were considered to suffer from extinction effect, were dropped off in subsequent refinements. Including the positions of hydrogen atoms obtained from a difference synthesis and the appropriate isotropic thermal parameters, further refinements were repeated with the weighting scheme:

$$w = (16.0/F_o)^2 \quad \text{for } |F_o| > 16.0$$

$$w = 1.0 \quad \text{for } 16.0 \geq |F_o| \geq 6.7$$

and

$$w = 0.5 \quad \text{for } 6.7 > |F_o| > 0.0.$$

The final R value was 0.090 for 337 observed reflections.

Crotonamide. The trial structure was obtained from the Patterson maps projected along the b and a axes, and corrected by the Fourier synthesis projected along the b axis. A three-dimensional Fourier map

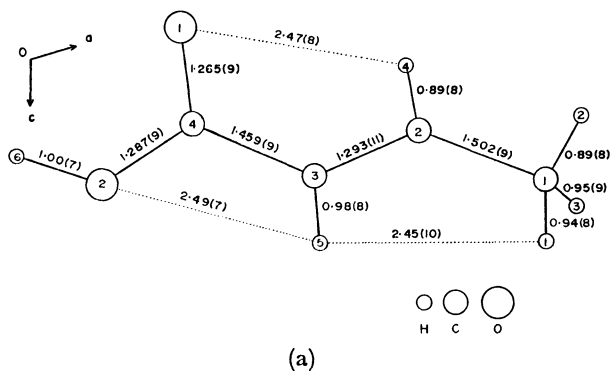
TABLE 1. CRYSTAL DATA

	Crotonic acid	Crotonamide
Chemical formula	$\text{CH}_3\text{CH}=\text{CHCOOH}$	$\text{CH}_3\text{CH}=\text{CHCONH}_2$
Molecular weight	86.09	85.12
Melting point	72 °C	158 °C
Space group	Monoclinic, C2/c	Monoclinic, $\text{P2}_1/\text{a}$
Systematic absences	hkl for $h+k$ odd $h0l$ for l odd	$h0l$ for h odd $0k0$ for k odd
Cell constants: a	15.30(3) Å	9.71(2) Å
b	4.06(1)	6.90(2)
c	16.18(2)	7.75(1)
β	107.4(1)°	104.0(4)°
Z	8	4
D_c	1.19 g/cm ³	1.12 g/cm ³
D_m	1.19	1.11
$F(000)$	368	184
μ for $\text{CuK}\alpha$ ($\lambda=1.5418$ Å)	8.18 cm ⁻¹	6.72 cm ⁻¹
Dimension of crystals used	$0.5 \times 1.5 \times 0.2$ mm $0.7 \times 0.5 \times 0.2$ mm	$0.3 \times 0.1 \times 0.1$ mm
Layers photographed	$h0l$ to $h2l$ $0kl$ to $7kl$	$0kl$ to $7kl$
Observed reflections	337	525

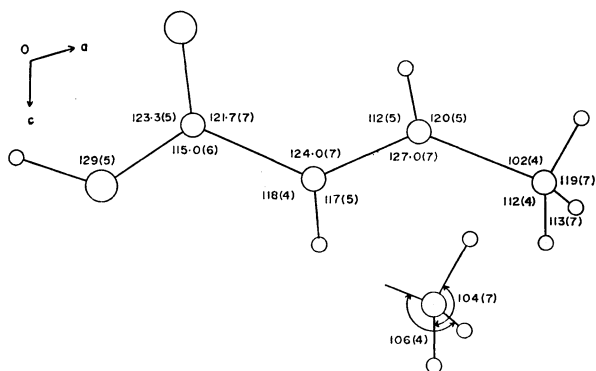
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TABLE 5. FINAL FRACTIONAL COORDINATES ($\times 10^3$) AND ISOTROPIC THERMAL PARAMETERS ($\text{\AA}^2$) FOR HYDROGEN ATOMS
Their estimated standard deviations are in parentheses.

	x/a	y/b	z/c	B		x/a	y/b	z/c	B
Crotonic acid					Crotonamide				
H(1)	383(5)	294(22)	224(4)	6.3(2.0)	H(1)	46(6)	895(10)	308(8)	8.1(1.6)
H(2)	409(4)	231(19)	147(4)	5.2(1.8)	H(2)	95(6)	764(8)	506(8)	6.1(1.3)
H(3)	405(5)	-73(24)	206(5)	8.2(2.5)	H(3)	196(4)	758(5)	401(6)	3.7(1.0)
H(4)	278(6)	-63(21)	78(5)	7.2(2.1)	H(4)	-72(5)	535(7)	268(7)	5.0(1.2)
H(5)	216(5)	329(19)	181(5)	6.0(1.9)	H(5)	247(4)	456(5)	252(5)	2.7(0.8)
H(6)	-10(5)	162(20)	62(5)	6.7(2.1)	H(6)	138(5)	12(6)	20(6)	4.4(1.1)
					H(7)	282(4)	164(7)	111(6)	4.5(1.1)

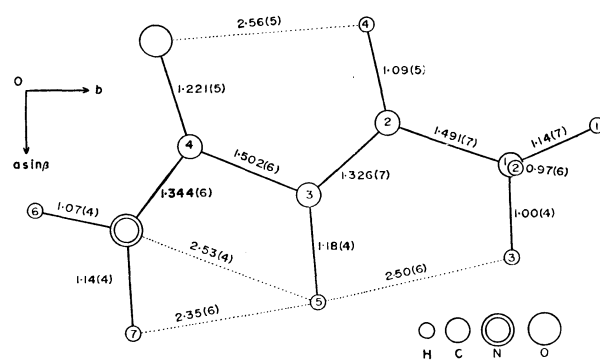


(a)

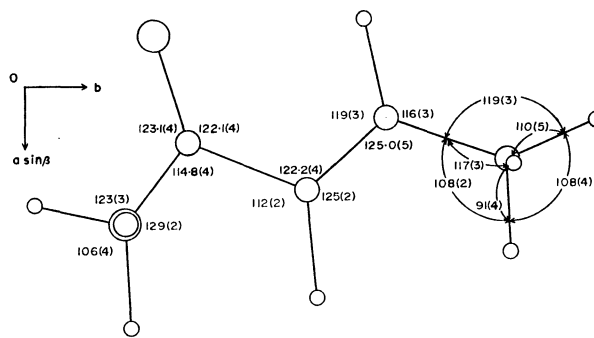


(b)

Fig. 1. The molecular structure of crotonic acid.
(a) Bond lengths ($\text{\AA}$) and numbering of atoms.
(b) Bond angles ($^\circ$). The estimated standard deviations are given in parentheses.



(a)



(b)

Fig. 2. The molecular structure of crotonamide.
(a) Bond lengths ($\text{\AA}$) and numbering of atoms.
(b) Bond angles ($^\circ$). The estimated standard deviations are given in parentheses.

TABLE 6. LEAST-SQUARES PLANES

Each plane is represented by $lX + mY + nZ = p$, where X , Y and Z (in $\text{\AA}$) are referred to the crystallographic axes.

Crotonic acid				Crotonamide			
(1) Best plane through the ethylenic group		(2) Best plane through the carboxyl group		(1) Best plane through the ethylenic group		(2) Best plane through the amide group	
l	0.2068		0.1931		0.0153		0.0597
m	0.8316		0.8390		-0.4587		-0.4781
n	-0.5536		-0.5431		0.8584		0.8358
p	0.0336		0.0208		0.1541		0.1146
Deviation ($\text{\AA}$)							
C (1)	-0.004	O (1)	0.0002	C (1)	-0.006	O	-0.001
C (2)	0.004	O (2)	0.0002	C (2)	0.007	N	-0.001
C (3)	0.003	C (3)	0.0002	C (3)	0.006	C (3)	-0.001
C (4)	-0.004	C (4)	-0.0006	C (4)	-0.006	C (4)	0.003

Table 3; the final atomic parameters in Tables 4 and 5. Bond lengths and angles along with the numbering scheme for atoms are shown in Fig. 1 for crotonic acid, and in Fig. 2 for crotonamide.

Molecular Shape. The equations of best planes and the deviations of the atoms from each plane are listed in Table 6. Dihedral angle between the ethylenic and carboxylic planes of crotonic acid is only 1.0° , and that between the ethylenic and amide planes of crotonamide is 2.9° , which are rather smaller than the corresponding value (3.6°) found in *trans*- β -2-furylacrylamide.⁶⁾ Thus the configurations of both crotonic acid and crotonamide are nearly planar. The bond length $C(2)=C(3)$ of crotonic acid is rather shorter. The difference between the bond lengths $C(4)=O(1)$ and $C(4)-O(2)$ in crotonic acid is very small as observed in β -methyl-*cis*-cinnamic acid and β -chloro-*cis*-cinnamic acid.⁷⁾ On the other hand, the $C(4)=O$ and $C(4)-N$ distances in crotonamide are very close to the values (1.225 and 1.349 Å, respectively) predicted from hybridization effect.⁸⁾ The $C=O$ bonds of both car-

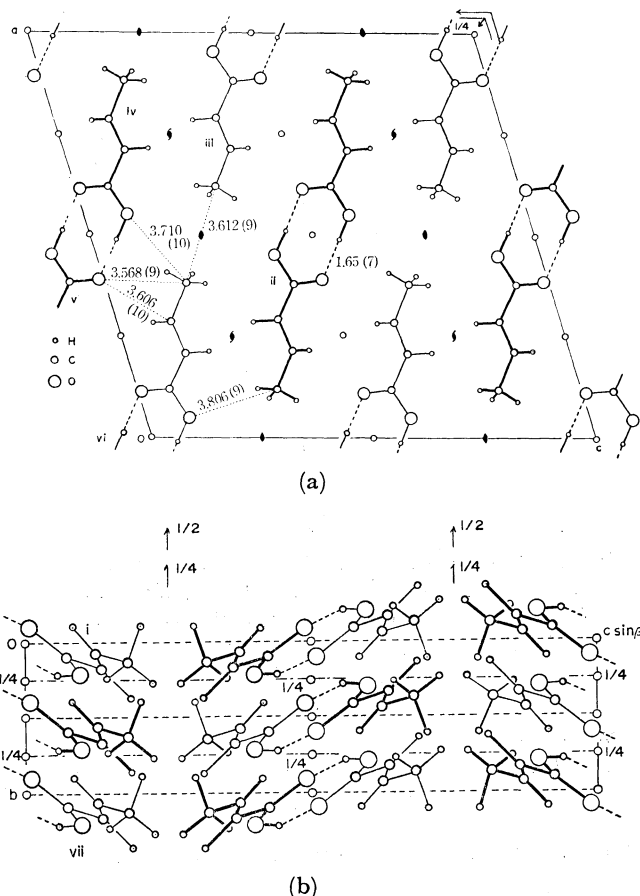


Fig. 3. Projections of the crystal structure of crotonic acid.

- (a) Viewed along the b axis.
(b) Viewed along the a axis.

Broken lines show hydrogen bonds, and dotted lines intermolecular contacts.

Symmetry code: i $x/a, y/b, z/c$; ii $1/2-x/a, 1/2+y/b, 1/2-z/c$; iii $1-x/a, y/b, 1/2-z/c$; iv $1/2+x/a, 1/2+y/b, z/c$; v $1/2-x/a, 1/2-y/b, -z/c$; vi $-x/a, -y/b, -z/c$; vii $x/a, 1+y/b, z/c$.

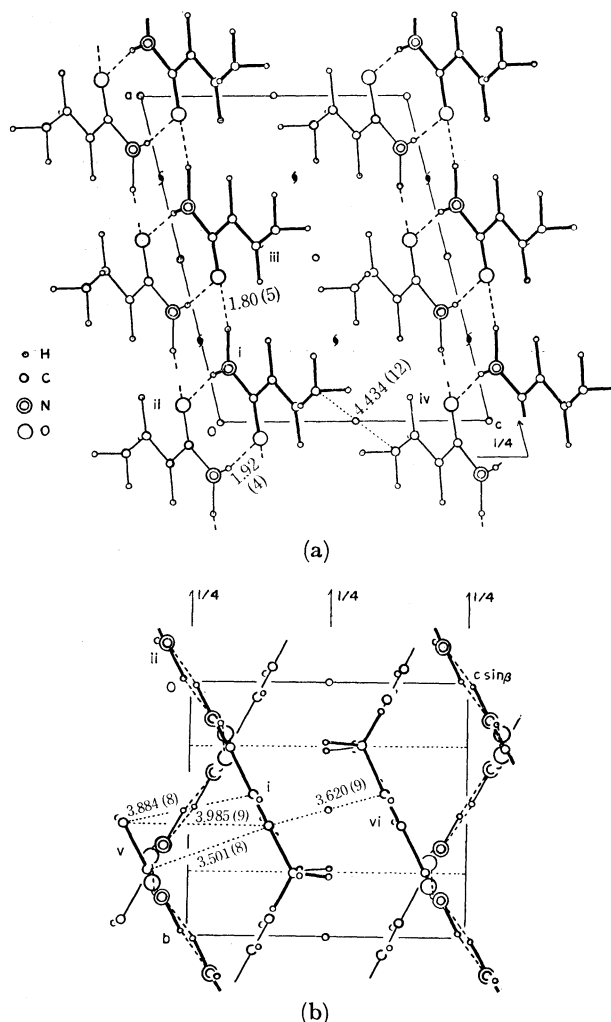


Fig. 4. Projections of the crystal structure of crotonamide.

- (a) Viewed along the b axis.
(b) Viewed along the a axis.

Broken lines show hydrogen bonds, and dotted lines intermolecular contacts.

Symmetry code: i $x/a, y/b, z/c$; ii $-x/a, -y/b, -z/c$; iii $1/2+x/a, 1/2-y/b, z/c$; iv $-x/a, 2-y/b, 1-z/c$; v $-x/a, 1-y/b, -z/c$; vi $-x/a, 1-y/b, 1-z/c$.

boxyl and amide groups adopt the *cis*-configuration with respect to the ethylenic $C=C$ bonds. Some intramolecular contacts are shown in Figs. 1 and 2.

A significant difference in the $C=O$ bond lengths has been pointed out between the carboxyl (1.209 Å) and the amide (1.247 Å) groups in fumaramic acid,⁹⁾ but the situation has reversed in the case of crotonic acid and crotonamide.

Molecular Packing. Projections of the crystal structure of crotonic acid viewed along the b and a axes and some intermolecular contacts are shown in Fig. 3 and those of crotonamide in Fig. 4. Geometry of the hydrogen bonds is listed in Table 7. The crystal structure of crotonic acid consists of two identical stacks of centrosymmetrically hydrogen-bonded pairs of the molecules, which are held together by van der Waals interactions between the methyl groups at a distance of 3.61 Å to form infinite chains along the

TABLE 7. HYDROGEN-BOND DISTANCES (Å) AND ANGLES (°)
Symmetry code is given in Figs. 3 and 4.

Crotonic acid			
O(1)···O(2 ^{vi})	2.645(6)	C(4)–O(2)···O(1 ^{vi})	115.6(4)
O(2)–H(6)	1.00(7)	C(4)–O(2)–H(6)	129(8)
H(6)···O(1 ^{vi})	1.65(7)	O(2)–H(6)···O(1 ^{vi})	174(8)
		C(4)–O(1)···O(2 ^{vi})	121.1(4)
		C(4)–O(1)···H(6 ^{vi})	123(8)
Crotonamide			
N···O ⁱⁱ	2.963(8)	C(4)–N···O ⁱⁱ	117.3(3)
N–H(6)	1.07(4)	C(4)–N–H(6)	123(3)
H(6)···O ⁱⁱ	1.92(4)	N–H(6)···O ⁱⁱ	165(4)
		C(4)–O···N ⁱⁱ	119.5(3)
		C(4)–O···H(6 ⁱⁱ)	122(2)
N···O ⁱⁱⁱ	2.862(7)	C(4)–N···O ⁱⁱⁱ	118.9(3)
N–H(7)	1.14(4)	C(4)–N–H(7)	129(2)
H(7)···O ⁱⁱⁱ	1.80(5)	N–H(7)···O ⁱⁱⁱ	153(4)
		N···O ⁱⁱⁱ –C(4 ⁱⁱⁱ)	142.7(3)
		H(7)···O ⁱⁱⁱ –C(4 ⁱⁱⁱ)	133(2)
		O ⁱⁱ ···N···O ⁱⁱⁱ	123.6(2)
		H(6)–N–H(7)	106(4)

[201] direction. The corresponding contact in α -tetronic acid¹⁰ is 3.81 Å, and that in *p*-toluic acid¹¹ 3.50 Å. The hydrogen-bonded carboxyl groups are not fully coplanar, being offset 0.04 Å.

Crotonamide molecules related by a center of symmetry are linked by N–H···O hydrogen bonds to form pairs at (000) and (1/2, 1/2, 0), which are held together side by side through the a-glide plane by additional N–H···O hydrogen bonds. This packing mode belongs to the glide-plane packing type.¹² The intrapair amide groups are distinctly not planar, the separation between the two corresponding parallel planes being 0.23 Å. The inclination of the amide pair to the

glide plane is 61.5° and the nearest methyl-methyl distance is 4.43 Å. The interpair angles C–N···O and C=O···N are 118.9 and 142.7°, respectively, which are close to those found in furylacrylamide.⁶

The structures of crotonic acid and crotonamide show short enough distances between ethylenic groups (C(3)···C(2^{vii}) 3.86 and C(3)···C(3^v) 3.88 Å, respectively) to dimerize or polymerize on irradiation with ultraviolet light, and the parallel alignment of double bonds considered as a pre-requisite for cycloaddition occurs in both compounds.

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