

Refinement of the Structure of Alclofenac, 4-Allyloxy-3-Chlorophenyl acetic acid (C₁₁H₁₁O₃Cl)

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Abstract □ The structure of alclofenac was determined by single crystal X-ray diffraction analysis. The compound was recrystallized from chloroform solution in monoclinic system, space group C2/c, with $Z=8$, $a=23,349(9)$, $b=14,295(3)$, $c=7,192(2)$ Å, $\beta=111,32(3)^\circ$, and $d_{obs}=1,29$, $d_{calc}=1,30$. The structure was solved by direct method and refined by least-squares procedure to the final R-value being 0,044 for 1055 reflections ($F \geq 6\sigma(F)$). The molecules are dimerized by OH...O type hydrogen bonds related by two-fold axis.

Keywords □ Anti-inflammatory agent, Alclofenac, X-ray diffraction, Direct method, Hydrogen bond.

The investigation of the molecular structure of related drugs and elucidation of the relationship between the molecular conformations and their physiological functions are of interest.

4-Allyloxy-3-chlorophenyl acetic acid, alclofenac (Fig.1) is a non-steroidal anti-inflammatory agent which inhibits prostaglandin synthesis (1-3). Some non-steroidal anti-inflammatory drugs show their structural similarities in the carboxyl group and it was reported that the structure of this part may relate to the drug action.

We already presented the molecular structure of this compound briefly (12). This paper deals with the precise structure analysis of alclofenac and comparison of the molecular structure with those of related compounds.

EXPERIMENTAL METHODS

The compound was recrystallized by the evaporation method from chloroform solution as colorless, transparent crystals. The space group was determined from Weissenberg photographs, and lattice constants were obtained from precise measurement on a diffractometer with MoK α radiations. The density was measured by the flotation method in a mixture of benzene and carbon tetrachloride. The crystallographic data are listed in Table I.

Intensities for 1690 independent reflections lim-

ited within $\sin \theta / \lambda = 0,601 \text{ \AA}^{-1}$ were measured on a computer-controlled automatic four circle diffractometer (Rigaku Denki Co. Ltd.) with Zr-filtered MoK α radiations by ω - 2θ scan technique, $3^\circ/\text{min}$. With a $0,2 \times 0,3 \times 0,6$ mm dimensioned crystal.

All the reflections were corrected for usual Lorentz and polarization effects, but no absorption correction being made. Of all 1690 independent reflections, 635 reflections which had $F < 6\sigma(F)$ were treated as unobserved.

The structure was solved by direct method using the program SHELX 76 (5) on VAX-11/780 computer system at Seoul National University. All the non-hydrogen atoms could be found by the successive Fourier syntheses.

It was refined by full matrix least-squares procedure with isotropic temperature factors, and R-value was decreased from 0,318 to 0,167. Further refinement with anisotropic temperature factors reduced R-value to 0,079.

A difference Fourier synthesis calculated at this

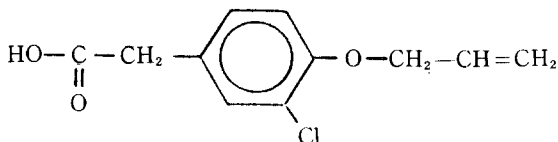


Fig.1. alclofenac.

Table I. Crystallographic data.

4-Allyloxy-3-chlorophenyl acetic acid	
Molecular formula: C ₁₁ H ₁₁ O ₃ Cl	M. W. 226. 66
Colorless transparent needle	Monoclinic
$a = 23.349(9) \text{ \AA}$	$b = 14.295(3) \text{ \AA}$
$c = 7.192(2) \text{ \AA}$	$\beta = 111.32(3)^\circ$
Volume of unit cell	2236.2 \AA^3
$d_{obs} = 1.29 \text{ g/cm}^3$	$d_{calc} = 1.30 \text{ g/cm}^3$
Space group: C2/c	Z=8

stage revealed all hydrogen atoms, and refinement including hydrogen atoms was continued.

The final R-value was 0.044 for 1055 reflections with $F \geq 6\sigma(F)$. The average and maximum ratios of parameter shift to e.s.d. after refinement were 0.007 and 0.064 for non-hydrogen atoms, 0.007 and 0.071 for hydrogen atoms, respectively.

The used atomic scattering factors were from *International Tables for X-ray Crystallography* (6).

RESULTS AND DISCUSSION

The final positional and thermal parameters are

listed in Table II and Table III, together with their estimated standard deviations. The observed and calculated structure factors are listed in Table IV.

As shown in Fig.2, the bond lengths and angles

Table III. Fractional coordinates of hydrogen atoms ($\times 10^3$). The isotropic temperature factors are expressed in the form of $\exp(-8\pi^2U)\sin^2\theta/\lambda^2$

Atom	bound to	x/a	y/b	z/c	U
H1	C2	323(2)	222(3)	330(6)	79(13)
H2	C4	347(2)	-44(3)	455(5)	66(12)
H3	C5	246(2)	-89(3)	318(6)	81(13)
H4	C7	401(2)	178(3)	567(6)	83(15)
H5	C7	417(2)	79(3)	616(6)	76(14)
H6	C9	142(2)	-108(3)	228(6)	85(14)
H7	C9	164(2)	-109(3)	34(6)	89(14)
H8	C10	60(3)	-52(5)	-125(9)	183(31)
H9	C11	39(3)	-157(4)	178(9)	152(27)
H10	C11	-13(3)	-133(5)	-10(10)	195(33)
H11	O3	532(2)	129(3)	396(7)	143(18)

Table II. Final positional and thermal parameters with their estimated standard deviations in parentheses ($\times 10^4$). The anisotropic temperature factors are expressed in the form of $\exp(-2\pi^2(U_{11}a^2h^2 + U_{22}b^2k^2 + U_{33}c^2l^2 + 2U_{12}a^*b^*hk + 2U_{13}a^*c^*hl + 2U_{23}b^*c^*kl))$

Atom	x/a	y/b	z/c	U_{11}	U_{22}	U_{33}	U_{23}	U_{13}	U_{12}
CL	1776(1)	2178(1)	1278(2)	962(8)	589(6)	990(9)	42(6)	91(7)	136(6)
C1	2312(2)	1331(3)	2386(5)	741(29)	632(25)	510(22)	5(20)	166(21)	82(22)
C2	2897(2)	1599(3)	3365(6)	864(35)	652(30)	586(26)	-14(22)	193(24)	-51(27)
C3	3339(2)	937(3)	4239(5)	736(29)	859(31)	439(22)	-3(21)	158(20)	-29(26)
C4	3160(2)	10(3)	4099(6)	794(34)	824(33)	520(26)	145(23)	182(24)	177(30)
C5	2574(2)	-270(3)	3113(6)	832(34)	597(27)	548(25)	93(22)	205(24)	48(25)
C6	2135(2)	390(3)	2245(5)	722(28)	593(25)	476(22)	4(20)	197(20)	31(23)
C7	3982(2)	1219(5)	5294(7)	842(33)	1057(43)	553(27)	-34(31)	96(24)	-3(32)
C8	4422(2)	1254(3)	4074(6)	756(30)	745(27)	525(24)	-46(22)	75(23)	-17(22)
C9	1369(2)	-777(3)	1052(8)	880(38)	658(30)	745(33)	15(25)	260(31)	-22(26)
C10	723(3)	-837(4)	-58(10)	905(41)	858(37)	1101(47)	95(34)	114(36)	-184(32)
C11	316(4)	-1318(5)	391(14)	1022(53)	1042(47)	1695(75)	172(47)	322(52)	-100(41)
O1	1545(1)	195(2)	1255(4)	737(19)	594(17)	733(18)	23(14)	126(16)	-8(14)
O2	4267(1)	1309(2)	2327(4)	709(18)	1423(28)	476(16)	-2(18)	52(14)	-155(18)
O3	5006(1)	1256(4)	5022(4)	716(19)	1487(30)	555(15)	54(18)	-14(16)	148(19)

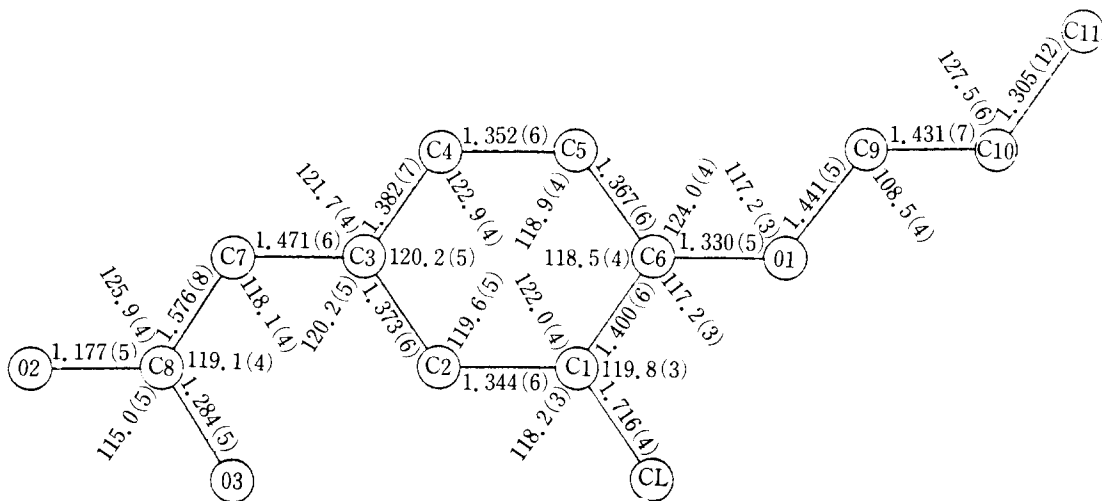


Fig.2. Bond lengths (Å) and angles (°) of alclofenac.

The estimated standard deviations are shown in parentheses.

are reasonable values compared with those of other compounds such as naproxen (7), flurbiprofen (8), indomethacin (9), (2-phenyl-4(*p*-chlorophenyl)-5-thiazolyl) acetic acid (10) and ibuprofen (11). The bond lengths and angles of the acetyl group, the noticed part, in alclofenac and the related compounds are listed in Table V. These values are very similar to those of other compounds.

The benzene ring and carboxyl group are planar. The equations of the two least-square planes and the deviations of the individual atoms from these planes are listed in Table VI. The dihedral angle between these planes is 97.87° . The skeletal conformation of C6, O1, C9, C10 and C11 is trans-trans form. The torsion angles around four bonds are shown in Fig.3.

The molecules are dimerized through the O3-H...O2 hydrogen bonds (2.798Å) related by two

Table V. Comparison of bond lengths and angles of alclofenac with related compounds.

	Alclofenac	Ibuprofen	Flurbiprofen	Naproxen
Lengths (Å)				
C7-C8	1.567	1.509	1.515	1.539
C8-O2	1.177	1.221	1.206	1.200
C8-O3	1.284	1.305	1.288	1.301
Angles (°)				
C7-C8-O2	125.9	122.9	122.8	123.3
C7-C8-O3	119.1	114.5	114.0	111.5
O2-C8-O3	115.0	122.6	123.2	125.2

Table VI. The equations of the least-square planes of benzene ring and carboxyl group and the deviations of individual atoms from these planes (Å).

Equations: Plane A (benzene ring)			
$0.52567X - 0.08442Y - 0.84649Z = 0.99428$			
Plane B (carboxyl group)			
$-0.01719X + 0.99825Y + 0.05161Z = 1.78381$			
Deviations:	Atom	from Plane A	from Plane B
	CL	-0.0223	-1.3037
	C1*	-0.0017	-0.1165
	C2*	0.0024	-0.5138
	C3*	-0.0042	0.4147
	C4*	0.0050	1.7362
	C5*	-0.0042	2.1509
	C6*	0.0027	1.2251
	C7**	-0.0160	-0.0032
	C8**	-1.4114	0.0121
	C9	-0.0386	2.9067
	C10	-0.0350	3.0098
	C11	0.7229	3.6625
	O1	0.0058	1.5184
	O2**	-2.4454	-0.0043
	O3**	-1.4600	-0.0043

* : Atoms used for the calculation of plane A.

** : Atoms used for the plane B.

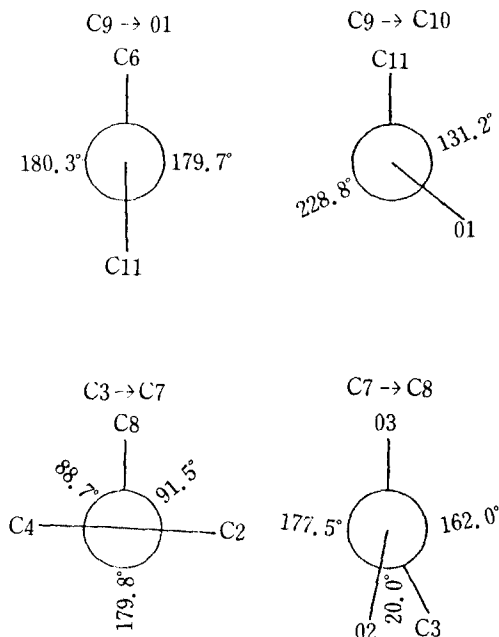


Fig.3. Newman projections of atoms around four bonds.

The arrow indicates the bond direction viewing down from front to back of paper.

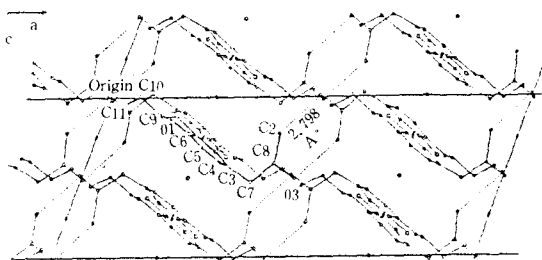


Fig.4. Molecular packing for alclofenac projected along b-axis.

The broken lines indicate hydrogen bonds.

-fold axis, and the intermolecular contacts between adjacent molecules are kept by van der Waals distances. The molecular packing is shown in Fig.4.

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