X-ray Structure Analysis Online 2012, VOL. 28 2012 © The Japan Society for Analytical Chemistry

X-ray Structure Analysis Online

Multiple Molecules in the Crystallographic Asymmetric Unit of (*Z*)-3-(3-Chlorophenyl)-2-phenyl acrylic acid

Rajnikant,* Vivek K. GUPTA,* Kamini KAPOOR,* Sunil KUMAR,** Kanahya Lal DHAR**

* X-ray Crystallography Laboratory, Post-Graduate Department of Physics, University of Jammu, Jammu Tawi - 180 006, India

**Department of Pharmaceutical Chemistry, I. S. F. College of Pharmacy, Moga, 142 001, India

[†] To whom correspondence should be addressed.

* rkvk paper11@gmail.com

The ¹H-NMR spectrum of the compound displayed resonance signals at 7.90 (s, 1H) and 9.8 (br s, exch.D₂O, 1H) due to the protons of the propenoic acid, in addition to the expected resonance signals of the aryl groups. The ¹³CNMR signals at δ 173.9 (CO₂H), 106(C-2) and 154.1(C-3) confirmed the propenoic acid moiety in the compound. Normally, distinction between *cis* and *trans* isomers of stilbenes is done by comparing the extinction coefficients in the UV Spectra (*trans* showing~28000and *cis* ~11000-12000), or by the coupling constant of the vinylic protons, if the double bond contains no other substituent. However, X-ray studies are accurate and give the exact geometry of compounds.

X-ray intensity data of a well-defined rod-shaped light-yellow crystal $(0.3 \times 0.2 \times 0.1 \text{ mm}^3)$ were collected at room temperature (293K) by using a CCD area-detector diffractometer (X'calibur system – Oxford diffraction make, U.K.) equipped with graphite monochromated MoK α radiation (λ =0.71073 Å). A total of 29559 reflections were collected, of which 8186 were treated as observed (I > 2 σ (I)). Data were corrected for Lorentz, polarization and absorption factors. The structure was solved by direct methods using SHELXS-97, and refined by full-matrix least-squares procedures using SHELXL-97 software. All of the hydrogen atoms were geometrically fixed and allowed to ride on the corresponding non-H atoms with C-H = 0.93 Å, O-H = 0.82 Å and U_{iso}(H) = 1.2U_{eq}(C), U_{iso}(H) = 1.5U_{eq}(O) for H atoms. The final refinement cycle yielded an R- factor of 0.0781 [wR (F²) = 0.1813] for the observed data. The residual electron density ranged from -0.372 to 0.455 eÅ⁻³. Atomic-scattering factors were taken from International Tables for X-ray Crystallography (1992, Vol. C, Tables 4.2.6.8 and 6.1.1.4).

	angle (e.s.d.)		angle (e.s.d.)
Molecule A	71.9(2)	Molecule B	77.9(2)
Molecule C	69.0(2)	Molecule D	72.1(2)
Molecule E	75.8(5)	Molecule F	70.9(3)
Molecule G	77.5(3)	Molecule H	73.9(3)
Molecule I	77.1(3)	Molecule J	70.5(4)

Table 2S	Dihedral	angles	between	the	pheny	1 ring	planes

D-HA	D-H(Å)	DA(Å)	HA(Å)	D-HA(°)
C5B-H5BCg13	0.93	3.935(8)	3.0543	158.68
C5J-H5JCg14	0.93	3.818(10)	2.9241	161.79
C6A-H6ACg19 ⁱ	0.93	3.754(10)	3.0572	133.05
C6C-H6CCg14	0.93	3.615(7)	2.8038	146.39
C6D-H6DCg11 ⁱⁱ	0.93	3.663(10)	2.8465	147.25
C6F-H6FCg15 ⁱⁱⁱ	0.93	3.636(1)	2.8445	143.73
C6G-H6GCg13 ^{iv}	0.93	3.745(11)	3.1009	127.92
C8B-H8BCg20	0.93	3.851(11)	3.2376	125.34
C8H-H8HCg16 ^v	0.93	3.577(12)	2.7409	150.09
C8J-H8JCg18 ⁱⁱ	0.93	3.641(13)	2.8949	138.21
C9C-H9CCg16 ^{vi}	0.93	3.959(8)	3.0810	158.11
C9G-H9GCg18	0.93	3.822(10)	2.9358	159.74
C11A-H11ACg1	0.93	3.694(8)	2.8636	149.31
C11B-H11BCg2	0.93	3.656(8)	2.8184	150.34
C11C-H11CCg3	0.93	3.656(6)	2.8198	150.26
C11D-H11DCg4	0.93	3.704(8)	2.8733	149.34
C11E-H11ECg5	0.93	3.693(7)	2.8410	152.99
C11F -H11FCg6	0.93	3.708(8)	2.8849	148.28
C11G-H11GCg7	0.93	3.637(10)	2.7831	153.15
C11H-H11HCg8	0.93	3.696(10)	2.8848	146.50
C11I-H11ICg9	0.93	3.674(11)	2.8255	152.19
C11J-H11JCg10	0.93	3.651(11)	2.8334	147.37
C14A-H14ACg10 ^{vii}	0.93	3.838(9)	2.9912	152.19
C14B-H14BCg4 ^v	0.93	3.814(8)	2.9807	149.95
C14C-H14CCg5 ^v	0.93	3.835(7)	3.0317	145.64
C14D-H14DCg2	0.93	3.839(9)	3.0220	147.54

Table 3S O-H...O and C-H... π hydrogen-bonding geometry (e.s.d.'s in parentheses). Cg1-Cg10 represent the center of gravity of the phenyl ring of ten molecules (A-J) and Cg11-Cg19 represent the center of gravity of the chlorophenyl ring of ten molecules (A-J).

C14E-H14ECg3	0.93	3.829(8)	2.9944	150.09
C14F-H14F Cg7	^{viii} 0.93	3.830(9)	3.0065	148.46
C14G-H14GCg6	0.93	3.829(10)	3.0188	146.55
C14H-H14HCg9	o ^{vii} 0.93	3.875(11)	3.0811	144.36
C14I-H14ICg8 ^{ix}	0.93	3.845(12)	2.9993	151.88
C14J-H14JCg1 ⁱⁿ	x 0.93	3.811(11)	3.0040	145.97
O2A-H2A O1H ⁱ	0.82	2.597(8)	1.785	170.6
O2D-H2DO2F	0.82	2.606(8)	1.799	167.7
O2J-H2JO1I ⁱⁱ	0.82	2.611(10)	1.800	169.8
O2C-H2C O1E ⁱⁱⁱ	0.82	2.578(7)	1.776	165.3
O2E-H2EO1C ^{vi}	0.82	2.608(7)	1.804	166.6
O2I-H2IO1J ^{vii}	0.82	2.610(10)	1.801	168.8
O2B- H2BO1G ^x	0.82	2.619(9)	1.815	166.4
O2G -H2GO1B ^{xi}	0.82	2.586(9)	1.783	166.1
O2H - H2HO1A ^x	ⁱⁱ 0.82	2.610(8)	1.799	169.8
Symmetry code :	i) - <i>x</i> ,1/2+ <i>y</i> ,1- <i>z</i> iv) - <i>x</i> , 1/2+ <i>y</i> , - <i>z</i> vii) 1- <i>x</i> ,1/2+ <i>y</i> ,1- <i>z</i> x) <i>x</i> ,- 1+ <i>y</i> , <i>z</i>	 ii) 1-x,-1/2+y,1-z v) -1+x, y, z viii) 1+x, y, z xi) x,1+y,z 	 iii) 1-x,1/2+y,-z vi) 1-x, -1/2+y,-z ix) -x,-1/2+y,1-z xii) -x,-1/2+y,1-z 	