

Note

Molecular and crystal structure of 6 α -acetoxy azadirone

Jibon Kotoky^{a*}, Mahendra Kalita^a, Gautam Kumar
Sarmah^a & Babulal Das^b

^aDepartment of Life Sciences, Institute of Advanced Study in
Science and Technology, Guwahati 781 035, India

^bDepartment of Chemistry, Indian Institute of Technology,
Guwahati, India

E-mail: jkotoky@yahoo.com

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The crystal structure of the title compound, **1** has been determined by X-ray crystallographic techniques. The compound crystallizes in orthorhombic space group $P2_12_12_1$ with unit cell parameters: $a=10.699(3)$ Å, $b=15.510(4)$ Å, $c=16.626(4)$ Å, $Z=4$. The structure has been solved by direct methods and refined to $R=0.0585$ for 6681 independent reflections.

Keywords: *Chisocheton paniculatus*, Meliaceae, limonoids, 6 α -acetoxy azadirone, crystal structure

Limonoids have attracted much attention because of their marked and notable insect antifeedant^{1,2} activity. They are found in the plants of Meliaceae³ family as secondary metabolites^{4,5}. One of the most important limonoid found in the plant *Chisocheton paniculatus*⁶ Hiern of Meliaceae family is 6 α -acetoxy azadirone⁷ **1**. This compound exists as α - as well as β -isomer. The crystal structure of β -isomer is already known⁸. Herein is reported the molecular and crystal structure of its α -isomer.

Materials and Methods

Powdered fruits of the plant *Chisocheton paniculatus* were extracted with petroleum ether (60-80°C) using a soxhlet apparatus, which gave a mixture of several compounds. One compound was isolated by usual chromatographic technique. On purification by crystallization from toluene, a white crystalline solid was obtained having a melting point of 192°C. The compound was identified as 6 α -acetoxy azadirone **1** by a comparison of its physical (e.g. m.p.) and spectroscopic (IR, NMR, and MS) data with the reported values⁷ as well as by comparing with authentic samples.

Experimental Section

Single crystals of the title compound **1** were grown from toluene by slow evaporation technique. The XRD data for a good quality single crystal ($0.40 \times 0.34 \times 0.22$ mm³) was collected at 296K temperature with a Bruker 3-circle diffractometer (Bruker Nonius SMART APEX 2) equipped with CCD area detector [using graphite monochromated MoK α radiation ($\lambda = 0.71073$ Å)]. Bruker SMART software⁹ was used for data collection and also for indexing reflections and the unit cell parameters. The data were integrated using SAINT software¹⁰. The structures were solved by direct methods and refined by full-matrix least-squares calculation using SHELXL software^{11,12}. Lattice parameters were determined from θ values in the range $3.81 < \theta < 28.40$. A total of 20488 reflections were recorded of which 6681 reflections were found unique. Absorption correction was not applied.

All the non-H atoms were refined in the anisotropic approximation against F^2 of all reflections. The H-atoms except those attached to N, O and F were placed at their calculated positions and refined in the isotropic approximations. The hydrogens attached to heteroatoms (N, O and F) were located from the difference Fourier maps and refined with isotropic displacement co-efficients. The final refinement cycle converged at $R=0.0501$ and $wR(F^2) = 0.1464$. Atomic scattering factors were taken from International Tables for X-ray crystallography.

The chemical structure of **1** is given in the **Figure 1** and its ORTEP view¹³ in **Figure 2**. Crystal data and other experimental details are given in the **Table I**.

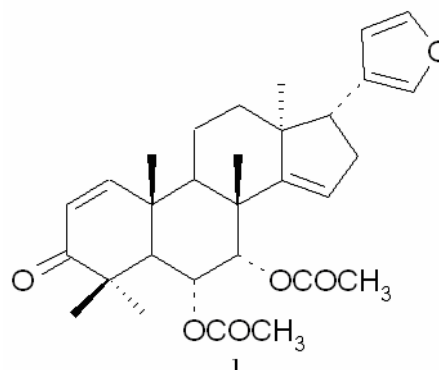
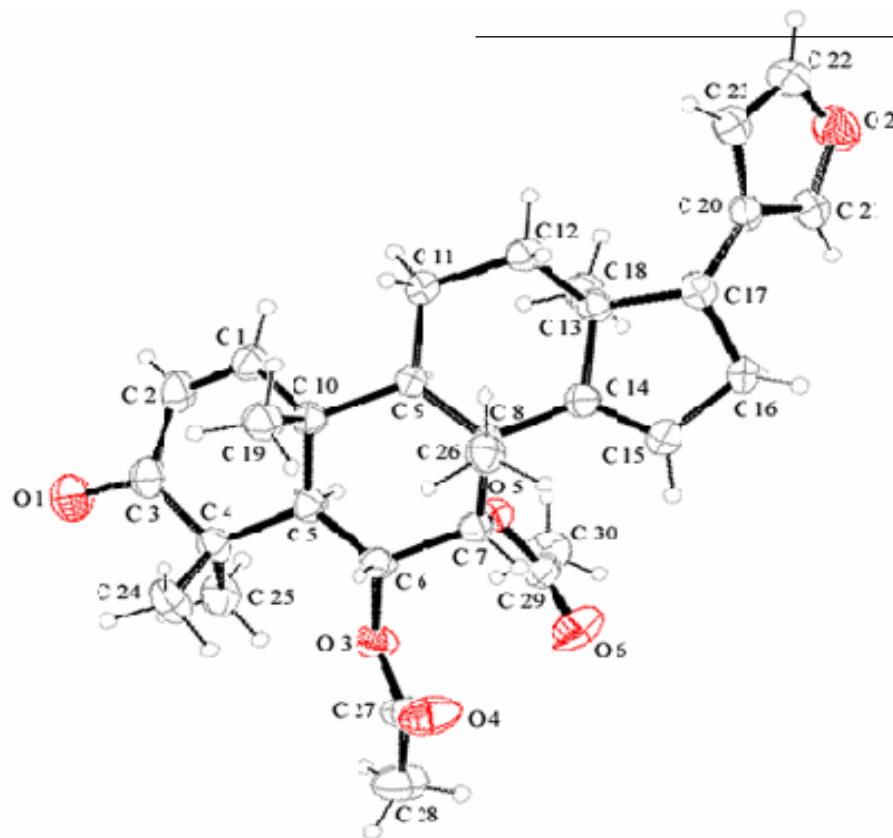


Figure 1 — Structure of 6 α -acetoxy azadirone

Figure 2 — An ORTEP view of **1** in 30% probabilityTable I — Crystal data and structure refinement for **1**

Empirical formula	: C ₃₀ H ₃₈ O ₆
Formula weight	: 494.60
Temperature	: 296(2) K
Wavelength	: 0.71073 Å
Crystal system	: Orthorhombic
Space group	: P2 ₁ 2 ₁ 2 ₁
Unit cell dimensions	: a = 10.699(3) Å α = 90° b = 15.510(4) Å β = 90° c = 16.626(4) Å γ = 90°
Volume	: 2759.0(12) Å ³
Z	: 4
Density (calculated)	: 1.191 Kg/m ³
Absorption coefficient	: 0.082 mm ⁻¹
F(000)	: 1064
Crystal size	: 0.48 × 0.34 × 0.22 mm ³
Theta range for data collection	: 3.81 to 28.40°
Index ranges	: -11 ≤ h ≤ 14, -20 ≤ k ≤ 20, -22 ≤ l ≤ 21
Reflections collected	: 20488
Independent reflections	: 6681 [R(int) = 0.0585]

Contd

Table I — Crystal data and structure refinement for **1**—Cont.

Completeness to theta = 28.40°	: 98.1%
Absorption correction	: None
Refinement method	: Full-matrix least-squares on F ²
Data / restraints / parameters	: 6681 / 0 / 332
Goodness-of-fit on F ²	: 0.998
Final R indices [I > 2σ(I)]	: R1 = 0.0501, wR2 = 0.1200
R indices (all data)	: R1 = 0.0876, wR2 = 0.1464
Absolute structure parameter	: 0.0(11)
Largest diff. peak and hole	: 0.174 and -0.180 e.Å ⁻³

Selected bond distances and bond angles are given in Table II. The furan ring in the molecule at the position 17 shows normal geometry and lies in the same plane with those of the two-acetyl groups at position 6 and 7. Both the acetyl groups lying in the same side represent a specific geometry to the molecule (α -isomer). The furan ring at the position 17 is found to be twisted with respect to the D ring of the

steroidal structure. The existence of the two acetyl groups at positions 6 and 7 in the molecule is expected

Table II — Bond lengths [Å] and angles [°] for **1**

O(5)-C(29)	1.347(3)
O(5)-C(7)	1.447(3)
O(3)-C(27)	1.341(3)
O(3)-C(6)	1.462(3)
O(2)-C(21)	1.360(3)
O(2)-C(22)	1.363(4)
O(1)-C(3)	1.217(3)
O(4)-C(27)	1.199(3)
O(6)-C(29)	1.198(3)
C(10)-C(1)	1.511(3)
C(10)-C(19)	1.541(3)
C(10)-C(5)	1.563(3)
C(10)-C(9)	1.571(3)
C(7)-C(6)	1.529(3)
C(7)-C(8)	1.537(3)
C(8)-C(14)	1.529(3)
C(8)-C(26)	1.541(3)
C(8)-C(9)	1.545(3)
C(5)-C(6)	1.523(3)
C(5)-C(4)	1.564(3)
C(13)-C(12)	1.513(3)
C(13)-C(14)	1.521(3)
C(13)-C(18)	1.542(3)
C(13)-C(17)	1.568(3)
C(17)-C(20)	1.498(3)
C(17)-C(16)	1.529(4)
C(14)-C(15)	1.314(3)
C(9)-C(11)	1.542(3)
C(1)-C(2)	1.327(3)
C(11)-C(12)	1.539(3)
C(3)-C(2)	1.454(4)
C(3)-C(4)	1.525(3)
C(29)-C(30)	1.482(4)
C(15)-C(16)	1.511(3)
C(4)-C(25)	1.523(4)
C(4)-C(24)	1.553(4)
C(27)-C(28)	1.483(4)
C(23)-C(22)	1.336(4)
C(23)-C(20)	1.410(4)
C(20)-C(21)	1.362(4)
C(29)-O(5)-C(7)	119.51(17)
C(27)-O(3)-C(6)	116.4(2)
C(21)-O(2)-C(22)	105.9(2)
C(1)-C(10)-C(19)	105.25(19)
C(1)-C(10)-C(5)	106.24(19)
C(19)-C(10)-C(5)	114.11(17)
C(1)-C(10)-C(9)	108.56(17)
C(19)-C(10)-C(9)	114.50(18)
C(5)-C(10)-C(9)	107.73(17)
O(5)-C(7)-C(6)	107.45(17)
O(5)-C(7)-C(8)	109.28(16)
C(6)-C(7)-C(8)	111.88(19)
C(14)-C(8)-C(7)	109.58(19)
C(14)-C(8)-C(26)	106.29(17)
C(7)-C(8)-C(26)	108.36(18)
C(14)-C(8)-C(9)	108.62(17)
C(7)-C(8)-C(9)	108.97(16)

C(26)-C(8)-C(9) 114.92(19)

Contd

Table II — Bond lengths [Å] and angles [°] for **1**—*Cont.*

C(6)-C(5)-C(10)	108.96(18)
C(6)-C(5)-C(4)	114.77(17)
C(10)-C(5)-C(4)	115.97(19)
C(12)-C(13)-C(14)	109.80(19)
C(12)-C(13)-C(18)	112.01(18)
C(14)-C(13)-C(18)	110.85(19)
C(12)-C(13)-C(17)	114.04(18)
C(14)-C(13)-C(17)	99.93(16)
C(18)-C(13)-C(17)	109.60(18)
C(20)-C(17)-C(16)	116.1(2)
C(20)-C(17)-C(13)	117.09(18)
C(16)-C(17)-C(13)	103.43(17)
C(15)-C(14)-C(13)	111.03(19)
C(15)-C(14)-C(8)	127.7(2)
C(13)-C(14)-C(8)	120.80(19)
C(11)-C(9)-C(8)	112.28(18)
C(11)-C(9)-C(10)	113.81(18)
C(8)-C(9)-C(10)	114.54(16)
C(2)-C(1)-C(10)	122.0(2)
C(12)-C(11)-C(9)	115.38(19)
O(1)-C(3)-C(2)	120.4(2)
O(1)-C(3)-C(4)	119.9(2)
C(2)-C(3)-C(4)	119.76(18)
O(3)-C(6)-C(5)	108.98(18)
O(3)-C(6)-C(7)	106.88(19)
C(5)-C(6)-C(7)	111.60(16)
O(6)-C(29)-O(5)	122.4(3)
O(6)-C(29)-C(30)	125.5(3)
O(5)-C(29)-C(30)	112.1(2)
C(13)-C(12)-C(11)	113.85(18)
C(14)-C(15)-C(16)	112.2(2)
C(25)-C(4)-C(3)	108.5(2)
C(25)-C(4)-C(24)	109.5(2)
C(3)-C(4)-C(24)	103.59(19)
C(25)-C(4)-C(5)	114.62(19)
C(3)-C(4)-C(5)	110.17(18)
C(24)-C(4)-C(5)	109.9(2)
O(4)-C(27)-O(3)	123.4(2)
O(4)-C(27)-C(28)	126.0(3)
O(3)-C(27)-C(28)	110.6(3)
C(22)-C(23)-C(20)	107.8(3)
C(21)-C(20)-C(23)	105.1(2)
C(21)-C(20)-C(17)	128.0(2)
C(23)-C(20)-C(17)	126.9(2)
C(1)-C(2)-C(3)	123.0(2)
C(15)-C(16)-C(17)	101.13(19)
C(23)-C(22)-O(2)	110.4(3)
O(2)-C(21)-C(20)	110.8(3)

Symmetry transformations used to generate equivalent atoms

to stabilize the three dimensional assembly of the molecules in the solid state through C-H...O weak

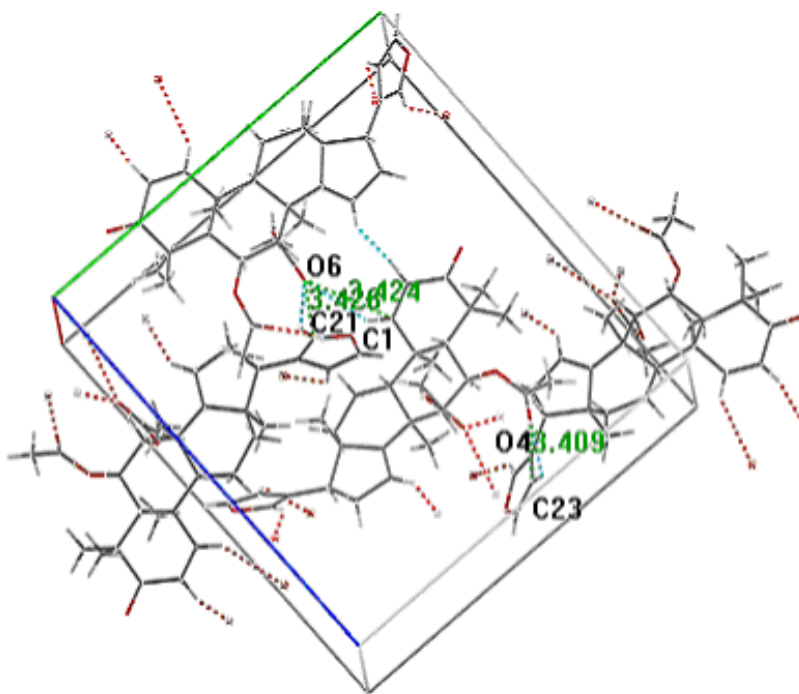


Figure 3 — Packing structure with C-H...O interactions

interaction involving C₂₃-H...O₄, C₂₁-H...O₆ and C₁-H...O₆ interactions ($d_{C23 \cdots O4}$ 3.40, $d_{C21 \cdots O6}$ 3.42 Å and $d_{C1 \cdots O6}$ 3.42 Å) respectively and van der Waals interactions as shown in the **Figure 3**. These distances are well within the reported range for C-H...O interaction¹⁴. The length of the double bond C3=O1 is slightly larger (1.217 Å) than the normal C=O double (1.20 Å) bond that could be attributed to the resonance caused by the α , β -unsaturated system present in the molecule.

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