X RAY STRUCTURE OF BETHANCOROL, A NEW COUMARIN FROM CNEORUM TRICOCCUM

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Further to our papers ^{1,2}, we are now reporting bethancorol, a new coumarin $(\underline{1a})$ mp 190-192°, $C_{15}H_{12}O_7$ (M⁺ 304), λ_{max}^{EtOH} 230, 262, 336 nm. Its UV and IR spectra show a coumarin nucleus. In the NMR (CDCl₂), two doublets at 6.38 and 7.58 δ (J 10Hz each) indicate a coumarin ring, singlets at 6.94 and 7.12 δ , aromatic protons in C-5 and C-8, another singlet at 3.80 (3H), a methoxyl group and those at 5.26 and 4.58 (2H each), -O-CH, and -CH,-OH groups. The signal at 4.58 shifted to 5.28 on adding a drop of TAL. 3 These data are compatible with structure la, 2a, 1b or 2b for bethancorol, determined as la by x ray analysis. Bethancorol crystallizes in a triclinic cell with $\underline{a} = 8.201$ (1), $\underline{b} = 12.186(2)$, c = 7.118 (1) Å, x = 96.84 (3), $\beta = 107.53$ (2), $\gamma = 96.61$ (3)^o, z = 2. A fully automatic diffractometer with monochromated MoK $_{
m sci}$ radiation was used to collect the intensities of 3839 independent reflexions for $\theta \leq 30^{\circ}$. 2092 were observed $I > 2\sigma(I)$ and used for the structure refinement. Calculated values for $< |E^2 - 1| >$ and $\langle |E| \rangle$, 1.074 and 0.748, show a hypercentric diffraction compared with the theoretical values for a centric distribution: 0.968 and 0.798 respectively. This hypersymmetry stems from the high pseudosymmetry in both planar molecules related by a crystallographic centre. The space group P1 was confirmed by the least squares refinement. This hypercentric structure's special diffraction pattern made it difficult to solve by direct methods. Using the Multan 74 System ⁴, a good molecular fragment was found, but shifted from its real posi≈ tion. A Pl difference map sited the fragment in relation to the crystallo= graphic inversion centre. The whole molecule was anisotropically refined to



an R of 0.096 4 , standard deviations in bond distances and angles being 0.007 %and 0.4 $^{\circ}$. The figure shows the molecular structure, double bonds being C(2)O(2) = 1.20, C(3)C(4) = 1.33, C(14)C(15) = 1.34 and C(18)O(19) = 1.19 %. The mole= cule may be described as a natural isoprene C(13)C(14)C(15)C(16)C(18) attached to the coumarin nucleus through O(11) and O(12). Both fragments are almost planar with a dihedral angle of 24 $^{\circ}$. All bond distances and angles appear normal. The molecules are linked by weak hydrogen bonds O(17)H(17)...O(2) of 2.95 %.



REFERENCES

- 1 AG González, BM Fraga and R Torres. Anal Quim 70 91 (1974)
- 2 AG González, BM Fraga and O Pino. Phytochemistry <u>13</u> 2305 (1974); Real Acad Ciencias (Madrid) <u>69</u> 347 (1975)
- 3 IR Trehan, C Monder and AK Bose Tetrahedron Lett 67 (1968)
- 4 Crystallographic programme systems used:

P Main, MM Woolfson, L Lessinger, G Germain and JP Declercq: MULTAN 74 - A system of computer programmes for the automatic solution of crystal structures from x ray diffraction data Univs: York (UK)/Louvain (Belgium) JM Stewart, FA Kundell & JC Foldwin The X Ray System of Crystallographic Programs Computer Science Centre, University of Maryland, USA (1970)