

STRUCTURE OF XANTHOMONADIN I, A NOVEL DIBROMINATED ARYL-POLYENE PIGMENT  
PRODUCED BY THE BACTERIUM XANTHOMONAS JUGLANDIS

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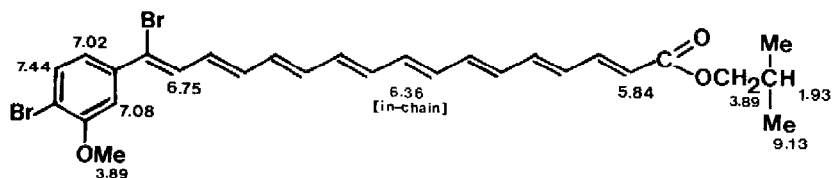
Only marine organisms are generally considered to command the capacity to concentrate, fix and incorporate bromine in organic compounds. The two reported exceptions to the marine connection involve a human spinal fluid component, 1-methylheptyl- $\gamma$ -bromoacetoacetate<sup>2</sup>, and the aromatic brominated polyene esters isolated by ourselves from the bacterial organism Xanthomonas juglandis strain XJ103<sup>3,4</sup>, a pathogen of a terrestrial plant (walnut). These latter compounds were incompletely characterized and we now report the detailed structure of one component, xanthomonadin I (1), as determined by x-ray crystallographic methods.

Treatment of crude, cellular extracts of XJ103 containing pigment-lipid complexes<sup>3</sup> with strong base for a short time (10 min., R.T.) gave, in effect, transesterification products. Thus, <sup>-</sup>OMe/MeOH or iso-<sup>-</sup>OBu/iso-BuOH yielded mixtures of methyl and iso-butyl brominated polyene esters, respectively. The superior solubility characteristics of iso-butyl esters vs. the methyl esters facilitated structural studies.

Extractive isolation of iso-<sup>-</sup>OBu treated extracts of XJ103 followed by Si gel column chromatography (developed with mixtures of petroleum ether-benzene and benzene-CHCl<sub>3</sub>) gave one major and several minor pigments. The major and least retentive pigment, the orange-red iso-butyl xanthomonadin I (1; ca. 1 mg from 540 g wet cells), crystallized from dilute acetone solution; m.p. 169-170°. The following physical properties were recorded:  $\lambda_{\max}$  (423), 445 ( $\epsilon=147,500$ ), 473 ( $\epsilon=122,900$ ) in acetone; (430), 453 ( $\epsilon=131,500$ ), 482 ( $\epsilon=109,500$ ) in CHCl<sub>3</sub>; IR (KBr) 3010, 2995, 1708, 1625, 1602, 1575, 1490, 1398, 1382, 1240, 1233, 1215, cm<sup>-1</sup>; m/e 572/574/576 (M<sup>+</sup>, C<sub>28</sub>H<sub>30</sub>Br<sub>2</sub>O<sub>3</sub>);  $\delta$ (CDCl<sub>3</sub>), see assignments on structure 1.

Isobutyl xanthomonadin I [1; isobutyl 17-(4-bromo-3-methoxyphenyl)-17-bromo-heptadeca-2,4,6,8,10,12,14,16-octaenoate] crystallizes in the monoclinic space group P2<sub>1</sub>/c with cell constants a = 42.46, b = 7.77, c = 7.87 Å,  $\beta$  = 92.47°, four molecules in the unit cell. The intensities of 2571 reflections (Cu K $\alpha$ , max  $2\theta$  = 100°) were measured at 85 K on a Picker automatic diffractometer. The crystals were extremely thin and of poor quality, resulting in irregular and broad (>2°)  $\omega$  scans. In order to enhance counting statistics, the intensity measurements were performed as 1.5°  $\omega$  scans (1° min<sup>-1</sup>) over the most intense part of the peaks.

The structure was solved by heavy atom Patterson and Fourier methods. Refinement is complicated by disorder, with about 20% of the molecules related to the major structure through a non-crystallographic mirror plane. Details will be reported elsewhere<sup>5</sup>.



ISOBUTYL XANTHOMONADIN I (1)

Oxidation of 1 by  $\text{KMnO}_4$  in *tert*-BuOH- $\text{H}_2\text{O}$  followed by esterification gave the expected methyl 4-bromo-3-methoxybenzoate, which after comparative chromatography (GLC and TLC) could not be separated from authentic material and thus supports the structure assigned by x-ray crystallographic methods.

Xanthomonadin I (1) and related compounds<sup>6</sup> have now been identified in all studied *Xanthomonas* species and pathotypes<sup>7</sup>.

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