

PHYTOCHEMICAL REPORTS

HYDROXYLAGOPODIN B, A SESQUITERPENOID QUINONE FROM A MUTANT STRAIN OF *COPRINUS MACRORHIZUS* VAR. *MICROSPORUS*

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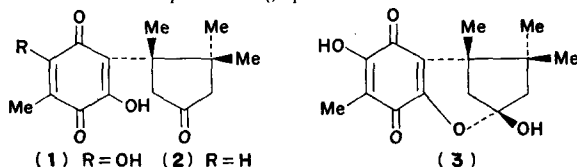
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Key Word Index—*Coprinus macrorhizus* var. *microsporus*; Basidiomycetes; mushroom; isolation; characterization, sesquiterpenoid quinone; hydroxylagopodin B.

Plant. *Coprinus macrorhizus* var. *microsporus*.
Source. Dr Philip G. Miles, Biology Department, SUNY at Buffalo.

Present work. A single gene mutant of the basidiomycete *Coprinus macrorhizus* var. *microsporus* (5074) excretes into the liquid culture medium an intense violet pigment which represents a series of possibly related compounds [1]. One of these has now been characterized as hydroxylagopodin B (1) which is clearly related to lagopodin B (2) [2], isolated from *Coprinus lagopus*.



The pigments were isolated by EtOAc extraction of the acidified filtered spent medium. The extract was evaporated *in vacuo* to dryness, the tarry residue redissolved in EtOAc and the pigments extracted into 0.1 M NaHCO₃. The bicarbonate phase was acidified (pH ~ 3) and extracted with C₆H₆. Following evaporation of this extract to near dryness, the residue was dissolved in 2% MeOH-CHCl₃ and separated on silica gel; (1) was eluted with 2% MeOH-CHCl₃ as the second purple band which yielded red-orange rhombic platelets (from C₆H₆-heptane), m.p. 184–186°. Yield: 3.5 mg/l of culture medium. Alkaline aqueous solutions of (1) were violet. Its solutions are decolorized by NaBH₄ and the color restored on aeration after the addition of NaHCO₃. UV λ_{max} (EtOH) 212, 307 and 351 nm is consistent with a 2,5 dihydroxy-3,6-dialkylbenzoquinone, e.g. helicobasidine [3].

The IR spectrum (KBr) showed OH at 3500 and 3385 cm⁻¹, characteristic (hydroxyquinone) CO absorptions at 1650 and 1627 cm⁻¹, and alkane C-H stretching bands in the region of 2900–3000 cm⁻¹. The cyclopentanone carbonyl band, which should appear at ~1750 cm⁻¹, was not present. This suggests that crystalline hydroxylagopodin B exists as a hemiketal structure (3) [3].

Hydroxylagopodin B has a molecular formula, C₁₅H₁₈O₅ (M⁺ 278), as determined by high resolution MS. Major ions in MS: *m/e* 280 (5.7), 278 (56.5), 221 (78.3), 194 (100), 166 (23.6). The strong M + 2 peak is observed which is typical of *p*-benzoquinones [3]. The fragmentation pattern confirms that hydroxylagopodin B is related to lagopodin B, the most abundant ion (100%) in the latter being at *m/e* 178. The ion *m/e* 166 arises from the expected elimination of an acetylene fragment from the benzoquinone ring.

Biological significance. The lagopodins have some antibiotic activity. No test of the antibiotic properties of hydroxylagopodin B was made.

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