

Structural elucidation of fungal polysaccharides isolated from the cell wall of *Plectosphaerella cucumerina* and *Verticillium* spp.

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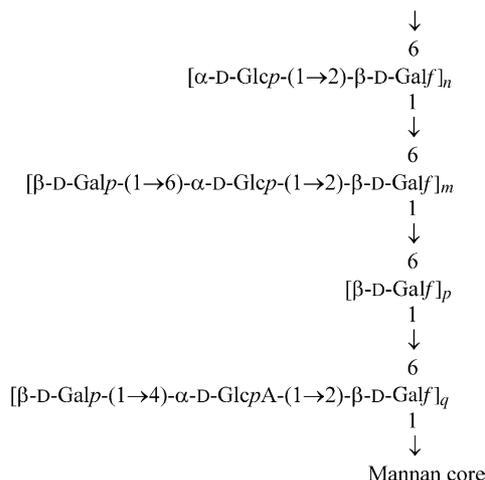
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Abstract—The structure of acidic fungal polysaccharides isolated from the cell wall of *Plectosphaerella cucumerina*, *Verticillium dahliae*, and *V. albo-atrum* has been investigated by chemical analysis, methylation analysis, and 1D and 2D ¹H and ¹³C NMR spectroscopy. The polysaccharides have an idealized repeating block of the type:



linked to a small mannan core (<15%), where $n = 13$, $m = 13$, $p = 5$, and $q = 8$ for *P. cucumerina*, and $n = 16$, $m = 16$, $p = 6$, and $q < 1$ for both *Verticillium* species.

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1. Introduction

The alkali-extractable and water-soluble fungal polysaccharides (FISS), which are minor components of the cell wall (2–8%), differ in composition and structure among genera and, in certain cases, among groups of species of a genus.¹ Polysaccharide moieties similar to that of the FISS polysaccharides have been shown to occur in

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glycoproteins.^{2,3} The complex carbohydrates of these molecules are antigenically relevant^{4–8} and serve different biological functions, one of the most important of which is its participation in cell–cell and/or cell–host recognition phenomena.⁹

The cosmopolitan fungus *Plectosphaerella cucumerina* can become a severe plant pathogen under appropriate conditions. Nevertheless, its systematics, as well as that of *Plectosporium tabacinum*, its anamorph, are problematic.^{10,11} Historically, *P. cucumerina* has been included in the Hypocreales^{12,13} but Uecker,¹¹ based on ascoma development, concluded that it should be placed in the Sordariales.

The polyphyletic genus *Verticillium* is the anamorph of several genera belonging to the Hypocreales.^{14,15}

Here we report on the novel structure of the polysaccharides FISS of *P. cucumerina*, *V. dahliae*, and *V. albo-atrum* and discuss the taxonomic placement of these species on the basis of their structures.

2. Results

The alkali-extracted water-soluble cell-wall polysaccharides (FISS) amounted to 5–7% of the dry cell-wall material in the three species. They are composed of mannose (6%), galactose (60%), and glucose (26%) in *P. cucumerina*, as shown by gas–liquid chromatography (GLC), and mannose (14%), galactose (54%), and glucose (32%) in both species of *Verticillium*. In addition, 6–10% of uronic acids were detected by the carbazole test. Absolute configuration analysis showed the D configuration for all of the sugars.

In *P. cucumerina*, methylation analysis of the reduced polysaccharide gave the derivatives corresponding to terminal glucopyranose and galactopyranose, 6-O- and 2,6-di-O-substituted galactofuranose, 6-O-substituted glucopyranose, and 6-O-substituted mannopyranose, and 4-O-substituted glucopyranuronic acid (Table 1).

Methylation analyses were also performed on both species of *Verticillium*, giving similar components to those found in *P. cucumerina*, although traces of the derivative corresponding to terminal glucopyranuronic acid were identified (<1%), instead of that of 4-O-substituted GlcpA (Table 1).

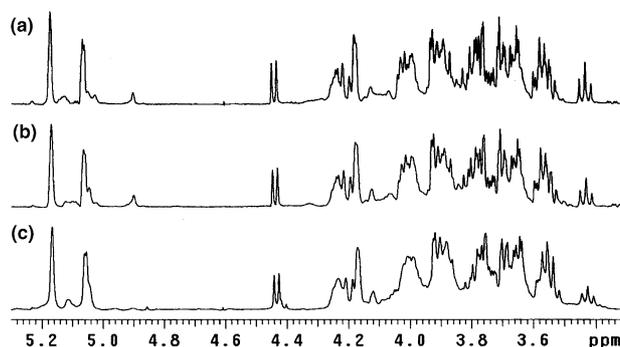


Figure 1. ¹H NMR spectra (D₂O, 40 °C, 300 MHz) of alkali-extracted water-soluble cell-wall polysaccharides FISS isolated from: (a) *V. dahliae*, (b) *V. albo-atrum*, and (c) *P. cucumerina*.

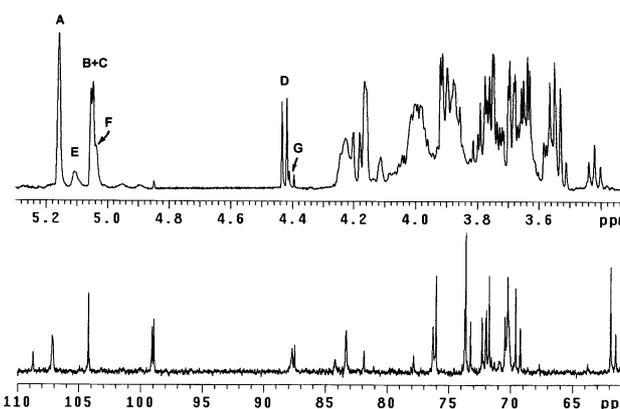


Figure 2. (a) ¹H NMR spectrum (D₂O, 40 °C, 500 MHz) and (b) ¹³C NMR spectrum (D₂O, 40 °C, 125 MHz) of FISS polysaccharide isolated from *P. cucumerina*. The anomeric peaks in the first spectrum have been labeled A–G.

The ¹H NMR spectra of the polysaccharides in the acid form were very similar in the three species (Fig. 1). Accordingly, the polysaccharide from *P. cucumerina* was then selected for further studies.

The high-resolution ¹H NMR spectrum (Fig. 2a) contained at least three major (5.16, 5.06, and 4.43 ppm) and three minor anomeric signals (5.12, 5.04, and 4.41 ppm).

The ¹³C NMR spectrum (Fig. 2b) exhibited four major (107.1, 104.2, 99.0, and 98.9 ppm) and two minor

Table 1. Linkage types deduced from methylation analysis of FISS polysaccharides

Linkage type	<i>V. albo-atrum</i> CECT 2693	<i>V. dahliae</i> CECT 2694	<i>P. cucumerina</i> CBS 137.33
Glc _p -(1→	16.2	16.0	17.6
Glc _p A-(1→	0.8	0.9	—
Gal _p -(1→	22.7	20.6	22.3
→4)-Glc _p A-(1→	—	—	8.3
→6)-Glc _p -(1→	17.8	23.6	13.6
→6)-Gal _f -(1→	4.5	2.3	2.0
→2,6)-Gal _f -(1→	38.0	36.6	36.2

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