## COMPARISON OF GLYCOPROTEINS FROM OPPORTUNISTIC PATHOGENS SCEDOSPORIUM PROLIFICANS AND RELATED PSEUDALLESCHERIA BOYDII

## <u>Gorin, P.A.J.<sup>1</sup></u>, Wagner, R.<sup>1</sup>, Sassaki, G.L.<sup>1</sup>, Souza. L.M.<sup>1</sup>, Silva, M.I.D.<sup>2</sup>, Pinto, M.R.<sup>2,3</sup>, Barreto-Bergter, E.<sup>2</sup>

<sup>1</sup>Departamento de Bioquímica, UFPR, 81531-990; Curitiba-PR, <sup>2</sup>Instituto de Microbiologia, CCS, UFRJ, 21944-590; Rio de Janeiro-RJ, <sup>3</sup>Departamento de Microbiologia, ICB, USP, 05508-000, São Paulo-SP,

Scedosporium prolificans is an ubiquitous filamentous fungus, causing infection in both immunocompetent and immunocompromized patients. The structure of Its mycelial galactorhamnomannoprotein (RMP-Sp) was compared with that of the related pathogen *Pseudallescheria boydii* (RMP-Pb), which contains Rhap- $(1\rightarrow 3)$ -Rha epitopes linked (1 $\rightarrow$ 3)- to Manp [1]. Reductive, alkaline  $\beta$ -elimination gave  $\alpha$ -Rhap- $(1\rightarrow 3)$ - $\alpha$ -Rhap- $(1\rightarrow 3)$ - $\alpha$ -Manp- $(1\rightarrow 2)$ -Man-ol, substituted at O-6 with  $\alpha$ -Glcp- $(1\rightarrow 4)$ - $\beta$ -Galp. It had a hapten reduction effect of ~75% [2]. RMP-Sp contained similar carbohydrates (63%), except that both 2-O- and 3-O-subst. Rhap units were present. Reductive  $\beta$ -elimination of RMP-Sp gave principally penta- and hexasaccharides, the former being  $[\alpha$ -Rha $p]_3$ - $\alpha$ -Manp-(1 $\rightarrow$ 2)-Man-ol and the latter with β-Galp substituents at O-6 of Man-ol (ESI-MS, ESI-MS-MS, <sup>13</sup>C NMR). Partial acetolysis of RMP-Sp gave  $[\alpha$ -Rhap- $(1\rightarrow 2)]_{1-2}$ -Rha,  $\alpha$ -Rhap- $(1\rightarrow 2)$ -Man, and  $\alpha$ -Manp- $(1 \rightarrow 3)$ -Man. The structural differences between S. prolificans and P. boydi PRMs were consistent with ELISA antigenicity tests, carried out with hyperimmune rabbit antiserum against mycelial forms of *P. boydii*, which reacted strongly but less so with S. prolificans RMP-Sp

[1] M.R. Pinto et al., Microbiology, 147, 1499 (2001); [2] M.R. Pinto et al., Glycobiology, 15, 895 (2005).

Supported by CNPq, FAPERJ, FAPESP, PRONEX-1996, and Fundação Araucária