

# Comparative genomics of *Ceriporiopsis subvermispora* and *Phanerochaete chrysosporium* provide insight into selective ligninolysis

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Efficient lignin depolymerization is unique to the wood decay basidiomycetes, collectively referred to as white rot fungi. *Phanerochaete chrysosporium* simultaneously degrades lignin and cellulose, whereas the closely related species, *Ceriporiopsis subvermispora*, also depolymerizes lignin but may do so with relatively little cellulose degradation. To investigate the basis for selective ligninolysis, we conducted comparative genome analysis of *C. subvermispora* and *P. chrysosporium*. Genes encoding manganese peroxidase numbered 13 and five in *C. subvermispora* and *P. chrysosporium*, respectively. In addition, the *C. subvermispora* genome contains at least seven genes predicted to encode laccases, whereas the *P. chrysosporium* genome contains none. We also observed expansion of the number of *C. subvermispora* desaturase-encoding genes

putatively involved in lipid metabolism. Microarray-based transcriptome analysis showed substantial upregulation of several desaturase and MnP genes in wood-contai