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Lignin-degrading activity and ligninolytic enzymes of different white-rot fungi: effects of manganese and malonate

Tamara Vares and Annele Hatakka

Abstract: Ten species of white-rot fungi, mainly belonging to the family Polyporaceae (Basidiomycotina), were studied in terms of their ability to degrade ¹⁴C-ring labelled synthetic lignin and secrete ligninolytic enzymes in liquid cultures under varying growth conditions. Lignin mineralization by the fungi in an air atmosphere did not exceed 14% within 29 days. Different responses to the elevated Mn²⁺ concentration and the addition of a manganese chelator (sodium malonate) were observed among various fungal species. This could be related with the utilization of either lignin peroxidase (LiP) or manganese peroxidase (MnP) for lignin depolymerization, i.e., some fungi apparently had an LiP-dominating ligninolytic system and others an MnP-dominating ligninolytic system. The LiP isoforms were purified from *Trametes gibbosa* and *Trametes trogii*. Isoelectric focusing of purified ligninolytic enzymes revealed the expression of numerous MnP isoforms in *Trametes gibbosa*, *Trametes hirsuta*, *Trametes trogii*, and *Abortiporus biennis* grown under a high (50-fold) Mn²⁺ level (120 μM) with the addition of the chelator. In addition, two to three laccase isoforms were detected.

Key words: white-rot fungi, lignin degradation, lignin peroxidase, manganese peroxidase, manganese, malonate.

Résumé: Les auteurs ont étudié 10 espèces de champignons de carie blanche, appartenant surtout à la famille des Polyporaceae (Basidiomycotina), afin de déterminer leur capacité à dégrader la lignine synthétique avec anneaux ¹⁴C marqués et à secréter des enzymes cellulolytiques en milieu de culture liquide, sous différentes conditions de croissance. La minéralisation de la lignine par les champignons, dans l'air atmosphérique, n'a pas dépassé 14% en 29 jours. On observe différentes réactions aux concentrations élevées de Mn²⁺ et à l'addition d'un chélateur du manganèse (malonate de sodium) parmi les différentes espèces fongiques. On peut relier ceci à l'utilisation, soit de la peroxydase de la lignine (LiP) ou soit à la peroxydase du manganèse (MnP) pour la dépolymérisation de la lignine, i.e., certains champignons possèdent apparemment un système ligninolytique dominé par la LiP et d'autres par la MnP. Les isoformes de la LiP ont été purifiées à partir du *Trametes gibbosa* et du *Trametes trogii*. La focalisation isoélectrique des enzymes ligninolytiques purifiés montre l'expression de nombreuses isoformes de la MnP chez les *Trametes gibbosa*, *Trametes hirsuta*, *Trametes trogii* et l'*Abortiporus biennis* cultivés en présence de fortes teneurs (50 fois) de Mn²⁺ (120 μM) avec addition du chélateur. On a en plus détecté la présence de deux ou trois isoformes de laccase.

Mots clés : champignon de carie blanche, dégradation de la lignine, peroxydase de la lignine, peroxydase manganique, manganèse, malonate.

[Traduit par la rédaction]

Introduction

The terms lignin-degrading or white-rot fungi designate a group of taxonomically heterogeneous higher fungi, characterized by their unique ability to depolymerize and mineralize lignin using a set of extracellular ligninolytic enzymes. These fungi are mainly saprophytes and belong to the orders Aphyllophorales and Agaricales (Basidiomycotina) (1). The origin of white-rot fungi dates back to the Carboniferous period and is closely linked with the formation of forests, since wood is the natural substrate for these species (13). The mechanism of lignin degradation by white-rot fungi has been

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Author to whom all correspondence should be addressed. e-mail: Annele.Hatakka@HELSINKI.FI intensively studied during the last 20 years in relation to biotechnical applications such as biopulping, biobleaching, treating of pulp mill effluents, and soil bioremediation (24, 28, 30, 31, 41). The molecular biology of lignin degradation was mainly determined in the white-rot fungus Phanerochaete chrysosporium (12). To date, compared with other ligninolytic microorganisms, Phanerochaete chrysosporium is much better characterized because the two ligninmodifying extracellular peroxidases, lignin peroxidase (LiP, EC 1.11.1.14) and manganese peroxidase (MnP, EC 1.11.1.13), were first isolated from this fungus (26, 49). However, other species of white-rot fungi were reported to degrade lignin as efficiently as Phanerochaete chrysosporium (5, 15, 35, 54). Moreover, several fungi show better selectivity for lignin removal (5, 30, 31, 42). Physiological conditions for lignin degradation, as well as secretion patterns of the ligninolytic enzymes, vary between different fungal species (14). Probably the most intriguing question is which enzyme or enzyme combination is responsible for lignin

Table 1. Description of the fungal strains.

IN THE STANDARD STANDARD SAN	Type of wood substrate in nature*
Polyporaceae*	The state of the second second
Abortiporus biennis (Bull. ex Fr.) Sing.†	Gymnosperms and angiosperms
Cerrena unicolor (Fr.) Murr.‡	Angiosperms
Junghuhnia separabilima (Pouz.) Ryv.§	Angiosperms
Trametes gibbosa (Pers.) Fr.§	Angiosperms, preferably Fagus
Trametes hirsuta (Fr.) Pilat†	Angiosperms
Trametes pubescens (Fr.) Pilat [†]	Angiosperms
Trametes trogii Berk.§	Mainly on Populus tremula
Trametes versicolor (L. ex Fr.) Pilat PRL 5721	Angiosperms
Trichaptum abietinum (Fr.) Ryv.§	Preferably on Picea
Ganodermataceae*	
Ganoderma applanatum (S.F. Gray) Pat.§	Angiosperms

^{*}According to Ryvarden and Gilbertson (45, 46).

depolymerization, particularly with regard to the preliminary steps. In this respect, the role of LiP, MnP, and laccase (EC 1.10.3.2, benzenediol:oxygen oxidoreductase) was widely discussed and they were suggested to be the main extracellular enzymes involved in lignin modification (14, 24). Different white-rot fungi produce various combinations of these enzymes. Recently, on the basis of extracellular enzyme patterns, three groups of lignin-degrading fungi were proposed: LiP-MnP, MnP-laccase, and LiP-laccase producing groups (14). MnP was found in many fungal species (14) and manganese is considered to be an important factor in lignin biodegradation (2, 4, 7, 21, 32, 38-40, 54). Free manganous ion Mn²⁺ is the primary substrate in the catalytic cycle of MnP (12). In the presence of an appropriate manganese chelator, a highly reactive Mn3+-chelated complex is formed that is capable of oxidizing phenolic lignin substructures (39, 56). Accumulation of manganese in wood, decayed by several white rot fungi under natural conditions, was described (2, 4).

In this paper we report the ability of 10 fungal species to mineralize and solubilize synthetic lignin (dehydrogenative polymerizate, DHP) and produce ligninolytic enzymes in liquid cultures. The influence of such regulative factors as manganese and its chelator, malonate (56), was studied. Ligninolytic enzymes (LiP, MnP, and laccase) were purified and characterized in a number of species.

Materials and methods

Fungi

Ten different species of white-rot fungi, known to be active wood-degraders in natural environment (3, 4, 6, 45, 46), were selected for investigation. These fungi cause white rot of wood and are therefore considered lignin-degrading species. Table 1 shows the taxonomical position, the natural substrate, and the origin of the fungi. Recent application of DNA sequencing methods in fungal systematics allows the phylogenetic classification of wood-decaying

fungi based on both molecular and morphological characters (17). However, since this classification is not yet complete (17), here we refer to the traditional systematics based mainly on the morphological characters (1, 6, 45, 46). Species and substrate types are designated according to Ryvarden and Gilbertson (45, 46). Strains were maintained on malt agar (2%, w/v) slants.

Production of extracellular ligninolytic enzymes

To elucidate the production of extracellular ligninolytic enzymes by the fungi, preliminary studies were carried out under three different culture conditions, with or without addition of lignin-related aromatic compounds, which previously were shown to enhance secretion of lignin-modifying enzymes, particularly LiP (24, 29, 53) and laccase (29). Fungi were cultivated in 100-mL flasks using basal low-nitrogen asparagine ammonium nitrate dimethylsuccinate medium (2.4 µM of manganese as Mn+2), supplemented with 56 mM glucose, pH 4.5 (15). A-flasks contained stationary cultures (total volume of 10 mL) without any inducer, cultures in the B-flasks were agitated (rotary shaker at 180 rpm) with the addition of 0.2 mM veratryl alcohol (Aldrich, Steinheim, Germany) and 0.05 % (w/v) Tween 80 (Fluka Chemie, Buchs, Switzerland) in a total volume of 25 mL, and the C-flasks were stationary cultures with addition of 1.0 mM veratric acid (Fluka Chemie, Buchs, Switzerland) in a total volume of 10 mL (53). Extracellular ligninolytic enzyme activities were followed in two replicate cultures starting from 2-4 days of growth. Cultures were inoculated with 4% (v/v) mycelial suspension (15) and grown at 28°C.

Degradation of synthetic lignin (DHP)

¹⁴C ring-labelled synthetic lignin (DHP) with molecular mass of 4000 to 10 000 (10, 54) was added to 100-mL Erlenmeyer flasks (ca. 50 000 dpm in a total volume of 10 mL) and evolution of ¹⁴CO₂ was monitored as described previously (15, 25) using a liquid scintillation counter (model 1411, Wallac Oy, Turku, Finland). To study the influence of manganese and its chelator, malonate, on DHP degradation, the following cultivation conditions were employed: (*i*) basal Mn²⁺ level (2.4 μM); (*ii*) 50-fold Mn²⁺ level (120 μM); (*iii*) basal Mn²⁺ with addition of 10 mM malonate; (*iv*) 50-fold Mn²⁺ with addition of 10 mM malonate.

[†] Fungal strains were obtained from the culture collection of the Institute of Microbiology, Academy of Science of Belarus, Minsk, Belarus.

[‡] Fungal strains were collected in Finland, isolated at the Department of Applied Chemistry and Microbiology, University of Helsinki, Helsinki, Finland.

[§] Fungal strains were collected in Russia, isolated at the Department of Applied Chemistry and Microbiology, University of Helsinki, Helsinki, Finland.

¹ Trametes versicolor PRL 572 was a gift from T. Johansson (University of Lund, Lund, Sweden).

Nonlabelled DHP (3 mg per flask; by S. Galkin, Department of Applied Chemistry and Microbiology, University of Helsinki, Finland, and G. Brunow, Department of Chemistry, University of Helsinki, Finland) was added to enhance ligninolytic activity. Cultures were grown without agitation under constant flow of air at 28°C. At the end of the cultivation, culture liquids were filtered through Whatman No. 4 (Whatman, Maidstone, U.K.) on glass sinters and mycelia were washed with a total of 15 mL H₂O. Radioactivity recovered as water-soluble products of DHP degradation, i.e., the portion of DHP not retained on the filter, was measured in liquids using a liquid scintillation counter. Mycelia and residual lignin were combusted in a combustion chamber (Junitek Oy, Turku, Finland) and the trapped ¹⁴CO₂ was quantified (15). Results are expressed as means of four replicates. Standard deviations are indicated in Fig. 2.

Enzyme assays

Ligninolytic enzyme activities were measured spectrophotometrically (Shimadzu 160A UV/visible spectrophotometer, Shimadzu, Kyoto, Japan) as described previously (29). LiP was assayed with veratryl alcohol (vacuum distilled prior to use) (50) and MnP with phenol red (Merck, Darmstadt, Germany) by monitoring the increase of absorbance at 520 nm in the presence of Mn²⁺ (29) and in the absence of Mn²⁺ for the peroxidase control. Laccase activity was measured with syringaldazine (Aldrich, Steinheim, Germany) (29). Protein concentration was determined with the Bradford assay (8) using a protein dye kit (Bio-Rad, Richmond, Calif.).

Purification of extracellular protein

For the purification of extracellular protein, Abortiporus biennis, Junghuhnia separabilima, Trametes hirsuta, Trametes gibbosa, and Trametes trogii were selected, since these species mineralized DHP relatively well and except for J. separabilima (52), their ligninolytic enzymes have not been studied in detail. Fungi were cultivated without agitation in 1000-mL Erlenmeyer flasks (total volume of 150 mL). To enhance the production of MnP, which obviously played an important role in DHP degradation by some of the selected species, basal medium supplemented with 1 mM veratryl alcohol, 120 µM Mn²⁺, and 10 mM sodium malonate was used in these cultivations. Extracellular culture liquids were harvested (ca. 1100 mL per fungus) on day 19 of growth and maintained at -20°C prior to protein purification. After thawing, culture liquids were filtered through Whatman No. 4 and Whatman GF/C glass microfiber filters (Whatman, Maidstone, U.K.) and concentrated to the final volume of 10 mL using an Amicon (Beverly, Mass.) ultrafiltration unit (YM membrane, 10-kDa cutoff). Concentrated samples were dialyzed overnight against 25 mM sodium acetate buffer (pH 5.5) and applied to a Mono-Q HR column (Pharmacia Biotech., Uppsala, Sweden) equilibrated with 25 mM sodium acetate buffer (pH 5.5). Proteins were eluted with gradient NaCl using an FPLC system (Pharmacia). Effluent was monitored at 280 nm (protein) and 405 nm (heme) and enzyme activities were assayed in eluted fractions.

Gel electrophoresis and isoelectric focusing

Purified ligninolytic enzymes were characterized by several gel techniques. Sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) and native polyacrylamide gel electrophoresis (native PAGE) (27) were carried out on 10% gels (1 mm thick) using a Protean II electrophoresis unit (Bio-Rad) with gel dimensions of 16 by 20 cm. Isoelectric focusing (IEF) was carried out using 0.5 mm thick polyacrylamide gels on a Multiphor II electrophoresis system (Pharmacia). The generated pH gradient was obtained by proportional mixing of ampholines with a pH range of 3.5 to 10.0 and pharmolytes with a pH range of 2.5 to 5.0 (Pharmacia). Gels were visualized by silver staining with Silver Stain and

Silver Stain Plus Kits (Bio-Rad) for total protein patterns and by specific activity stainings for ligninolytic enzymes, i.e., LiP and MnP isoenzymes were stained with phenol red at pH 4.5 in the absence and presence of Mn²⁺, respectively (37, 54, 55), and laccase activity was visualized with phenol red (pH 4.5) in the absence of H₂O₂ or with guaiacol.

Results

Production of ligninolytic enzymes by the fungi in low-Mn²⁺ medium supplemented with aromatic compounds

Laccase was produced by many species in relatively high titers. Secretion of this enzyme increased in cultures supplemented with aromatic compounds (veratryl alcohol or veratric acid) in comparison with non-induced cultivations, e.g., the level of laccase production in the most efficient laccase producer, Cerrena unicolor, reached 10 nkat·mL⁻¹. In A. biennis and J. separabilima cultures, maximum titers of laccase activity were 1.0 and 1.6 nkat·mL⁻¹, respectively. Trametes gibbosa and Trametes trogii produced 0.5 nkat·mL⁻¹ of this enzyme. On the other hand, in Trametes hirsuta, laccase activity was not detectable under the conditions studied, but laccase was detected during subsequent purification of extracellular proteins (shown later).

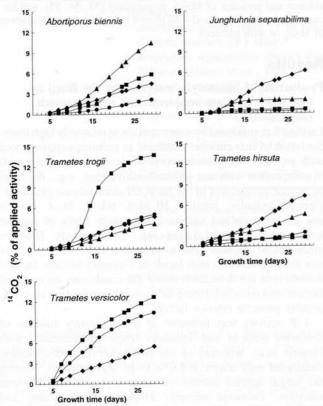
LiP activity was followed in the stationary cultures of *Trametes gibbosa* and *Trametes trogii* supplemented with veratric acid, whereas in the two other growth-condition treatments only traces of LiP activity was detected. Among the fungal species studied, cultures of *A. biennis, Cerrena unicolor, Trametes hirsuta, Trichaptum abietinum*, and *Ganoderma applanatum* grown under various conditions did produce traces of LiP-like activity as detected by the veratryl alcohol oxidation method. However, since the presence of LiP was not confirmed by the other analysis (protein purification or IEF), production of LiP by these fungi needs further investigation. Three LiP isoforms from *J. separabilima* were previously purified by us (52), and multiple LiP isoenzymes of *Trametes versicolor* PRL 572 were reported by other authors (18, 19).

Rather high levels of MnP was detected in the stationary cultures of *Trametes hirsuta* with and without addition of aromatic compounds. However, due to the low-Mn²⁺ culture medium and insensitive MnP assay used (see Materials and methods), levels of MnP measured in culture liquids of the other fungi were not high, and in many cases it was difficult to distinguish between Mn²⁺-dependent (MnP) and Mn²⁺-independent peroxidase activity. Subsequent purification of extracellular proteins from the high-Mn²⁺ cultures of five fungal species showing DHP-degrading capacity showed production of multiple MnP isoforms by these fungi (described later).

Degradation of synthetic ¹⁴C ring-labelled lignin by the fungi under altered manganese concentrations with and without malonate addition

Among the 10 fungal species investigated, 5 fungi displayed notable DHP-degrading ability under the growth conditions used. Figure 1 shows mineralization of ¹⁴C ring-labelled DHP by these species grown under low (basal) and high (50-fold) Mn²⁺ levels, with or without malonate addition, during the 29-day culture period in an air atmosphere. In

Fig. 1. Mineralization of DHP by different fungi cultivated under basal Mn^{2+} (\blacksquare), 50-fold Mn^{2+} (\blacktriangle), basal Mn^{2+} with malonate (\bullet), and 50-fold Mn^{2+} with malonate (\bullet).



general, the percentage of DHP mineralized to CO₂ by the fungi did not exceed 14% under all cultivation conditions. Results obtained with the five other fungi, which showed very low or negligible ability to mineralize DHP, are not presented in graphs. Species such as Cerrena unicolor, Trametes gibbosa, Trametes pubescens, and Trichaptum abietinum showed vigorous growth, but their capacity to mineralize DHP was very low. Ganoderma applanatum showed moderate growth and also a very low ability to degrade DHP under the conditions studied.

Fungi showed differential response to both manganese and malonate supplementation. The effect of Mn²⁺ addition on lignin mineralization was a stimulated evolution of ¹⁴CO₂ in cultures of many species. However, mineralization and solubilization of DHP by *Trametes trogii* was greater at the basal Mn²⁺ level compared with the high Mn²⁺ level (Figs. 1 and 2). In the *Trametes versicolor* cultures, Mn²⁺ seemed to repress DHP mineralization (Fig. 2). A slight increase in ¹⁴CO₂ evolution in high Mn²⁺ *Trametes gibbosa* culture was observed, but even here only about 2% of applied activity was detected as ¹⁴CO₂ (results not shown).

The supplementation of the manganese chelator sodium malonate to the low- and high-manganese cultivations induced species-specific effects on DHP degradation. In the case of *A. biennis*, malonate addition inhibited ¹⁴CO₂ evolution both in cultures grown under basal and 50-fold Mn²⁺ concentrations (Fig. 1) but did not notably affect the amount of water-soluble products (Fig. 2). Malonate addition clearly inhibited lignin mineralization by *Trametes trogii* in cultures supplemented with low basal Mn²⁺ content, but the level of

water-soluble products did not notably change. Patterns in *J. separabilima* and *Trametes hirsuta* were similar, with the highest mineralization rate occurring under 50-fold Mn²⁺ and malonate (Figs. 1 and 2). In these fungi, radioactivity recovered as water-solubles was also much higher in high-Mn²⁺ malonate cultures than in high-Mn²⁺ cultures without chelator (Fig. 2). In *Trametes versicolor*, malonate caused a slight inhibitory effect on the DHP degradation in low-Mn²⁺ cultures (Figs. 1 and 2). In high-Mn²⁺ cultures of *Trametes versicolor* supplemented with malonate, the mineralization rate was low (Fig. 1), although radioactivity recovered as water-soluble products was as high as 18% of totally applied (Fig. 2).

Purification and characterization of ligninolytic enzymes produced in high-Mn²⁺ medium supplemented with veratryl alcohol and malonate

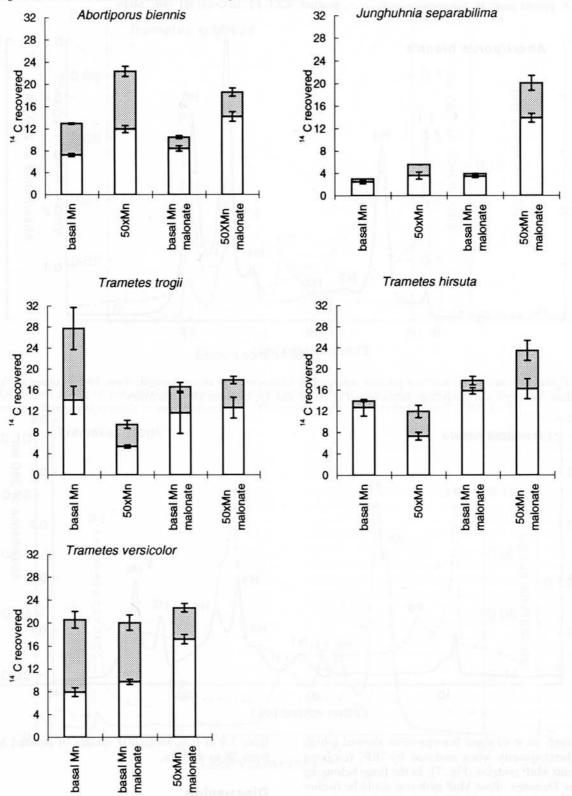
Profiles of extracellular proteins of four fungi grown under the 50-fold high Mn^{2+} level, with addition of 1 mM veratryl alcohol and 10 mM malonate, are shown in Figs. 3-6. Extracellular enzymes were also purified from J. separabilima cultures, but because of the low extracellular protein level produced by this fungus, the results are described in the text.

Enzyme activities were first assayed in eluted fractions and then the enzymes were analyzed by SDS-PAGE, native PAGE, and IEF using various staining procedures. Multiple MnP and laccase isoforms of A. biennis, Trametes hirsuta, Trametes gibbosa, and Trametes trogii were detected by enzyme activity staining of IEF gels. Isoenzyme patterns of Trametes gibbosa and Trametes trogii are shown in Fig. 7. In all elution profiles a distinct protein peak, designated P1 (Figs. 3-6), was observed, consisting of several unknown proteins the activity of which has not yet been defined. An absorption peak at 405 nm (hemeprotein) corresponding to protein peak P1 was also detected in all profiles. SDS-PAGE separation of P1 showed the presence of several protein bands with molecular masses of the major proteins being 33 and 23 kDa. H₂O₂-producing activity, assayed with methylglyoxal, glucose, and cellobiose as substrates (22), was not found in the protein peak P1. Laccase activity corresponded to the protein peaks P2 (found in all fungi) and P3 (Trametes hirsuta). Two to three laccase isoforms with molecular masses of 68 to 64 kDa and pIs of 4.0 to 3.7 were detected. In the IEF analysis, laccase but not MnP isoforms were stained with phenol red irrespective of the presence or absense of manganese in the substrate mixture (Fig. 7).

When comparing LiP and MnP isoenzyme profiles, fungal specific patterns were detected (Figs. 3-6). Two LiP isoforms with molecular masses of 44 and 41 kDa and pIs of 3.7 and 3.8 were found and characterized in *Trametes trogii* (corresponding peaks H5 and H4 in Fig. 6). LiP activity was detected in the protein elution profile of *Trametes gibbosa* (corresponding peak H1 in Fig. 5). However, the lack of LiP staining in the IEF analysis was probally due to the low enzyme amount and overlapping laccase activity (H1 in Fig. 7A). In the protein elution profiles of *Trametes hirsuta* and *A. biennis*, LiP was not identified. One possible reason for this could be the low levels of this enzyme expressed. The poor extracellular protein production in *J. separabilima* did not enable the detection of LiP in the elution profile. How-

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Fig. 2. (Evolution of ${}^{14}\text{CO}_2$ (\blacksquare) and production of ${}^{14}\text{C}$ -labelled water-soluble compounds (\square) by day 29 of DHP degradation by different white-rot fungi. Bars show standard deviations of four replicate cultures.



ever, under different culture conditions (high nitrogen and low manganese) three LiP isoforms were reported earlier in this species (52). Under the high Mn²⁺ conditions used in this study, one MnP isoenzyme, with a pI near 3.8 and showing a clear Mn²⁺-dependence of phenol red oxidizing in

activity stainings, was purified from J. separabilima (results not shown). Thus, MnP was expressed in all fungi cultivated under high $\mathrm{Mn^{2+}}$ level with malonate addition. In most fungal species this ligninolytic enzyme was found in multiple forms that were eluted as different hemeproteins (Figs. 3 –

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Fig. 3. FPLC elution profile of extracellular proteins separated by anion-exchange chromatography from 19-day culture of *Abortiporus biennis* supplemented with 50-fold Mn²⁺, 10 mM malonate, and 1 mM veratryl alcohol. —, absorbance at 280 nm; —, absorbance at 405 nm; P, protein peak; H, hemeprotein peak; ..., gradient NaCl; P2, laccases; H1-H6, MnPs.

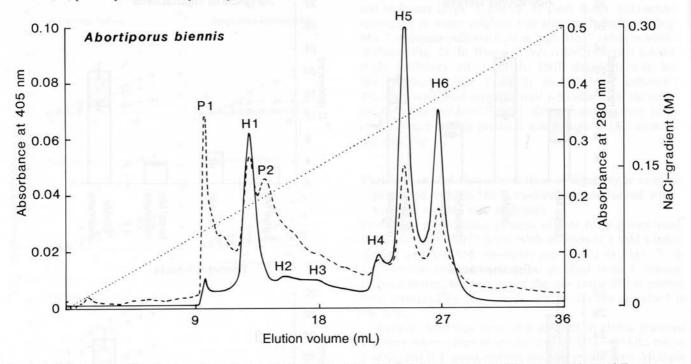
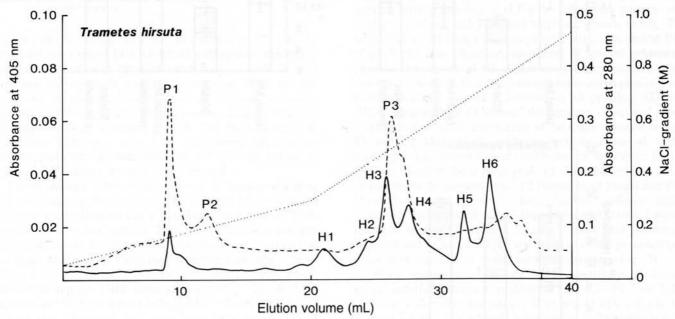


Fig. 4. FPLC elution profile of extracellular proteins separated by anion-exchange chromatography from 19-day culture of *Trametes hirsuta*. Culture conditions and symbols as indicated in Fig. 3. P2 and P3, laccases; H1-H6, MnPs.



6). In addition, some of these hemeproteins showed a high degree of heterogeneity when analysed by IEF, resolving into numerous MnP proteins (Fig. 7). In the fungi belonging to the genus *Trametes*, these MnP isoforms could be further subdivided into a group of MnPs with relatively high pI values (ca. from 4.4 to 7.0) and a group of more acidic MnP isoforms consisting of two major proteins with pIs of 3.9–4.0 and 3.7–3.9, respectively. *Abortiporus biennis* produced a large number of MnPs, which covered a pI range

from 3.9 to 6.5. Molecular masses of isolated MnPs were from 38 to 45 kDa.

Discussion

Until recently, basic research on lignin degradation, e.g., its mechanisms, physiology, enzymology, and molecular biology, was mainly carried out on the white-rot basidiomycete *Phanerochaete chrysosporium* (10, 12, 24–26, 38, 39, 49,

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Fig. 5. FPLC elution profile of extracellular proteins separated by anion-exchange chromatography from 19-day culture of *Trametes gibbosa*. Culture conditions and symbols as indicated in Fig. 3. P2, laccases; H1, LiP; H2, MnPs.

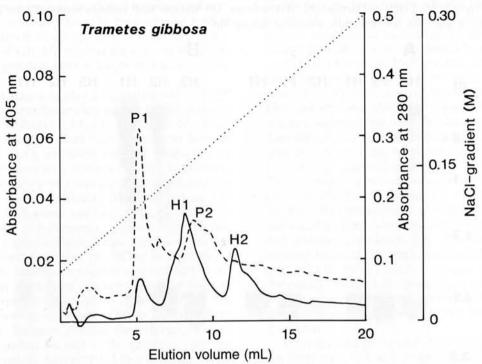
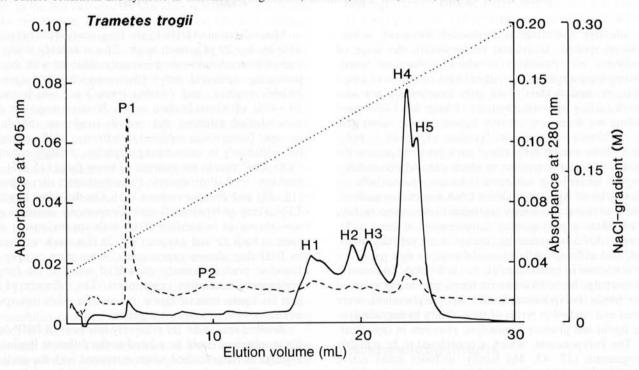


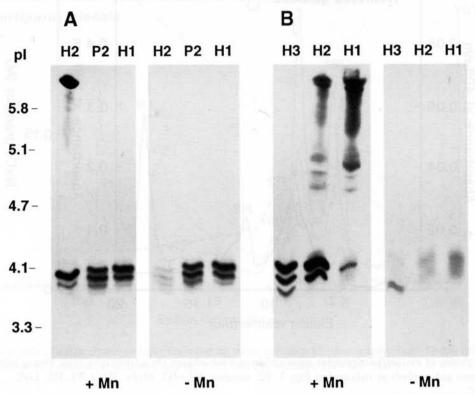
Fig. 6. FPLC elution profile of extracellular proteins separated by anion-exchange chromatography from 19-day culture of *Trametes trogii*. Culture conditions and symbols as indicated in Fig. 3. P2, laccases; H1-H3, MnPs; H4, LiP1; H5, LiP2.



50, 56) belonging to the family Corticiaceae (order Aphyllophorales) (1, 17). However, data on ligninolytic activity and lignin-modifying enzymes of other white-rot fungi are accumulating (5, 10, 14-16, 18-21, 29, 32-36, 40, 43, 44, 47, 51-55). These studies indicate that both the physiologi-

cal conditions for lignin degradation and enzyme systems expressed are fungal specific and could differ from those found in *Phanerochaete chrysosporium*. This diversity provokes a number of questions. Are such differences connected with the taxonomic position and (or) ecology of the fungi,

Fig. 7. Isoelectric focusing of ligninolytic enzymes purified from *Trametes gibbosa* and *Trametes trogii*. Protein (P) and hemeprotein (H) peaks are designated as in Figs. 5 and 6. (A) Comparative staining of *Trametes gibbosa* enzymes with phenol red in the presence and absence of Mn²⁺ reveals MnP (H2) and laccase (P2, H1) isoforms. (B) Multiple MnP isoforms separated from *Trametes trogii* are stained in the presence of Mn²⁺ but not seen under conditions lacking Mn²⁺.



e.g., substrate specialization (hardwood, softwood, or certain wood species, heartwood or sapwood), the stage of degradation, etc. Evolution of white-rot fungi as woodinhabiting microorganisms is linked to the evolution of forest ecosystems, and evidently not only morphology but also lignin-degrading enzymatic systems of fungi were developed following the formation of new lignocellulosic substrates. Thus, differences in enzymatic systems expressed in fungi could, to some extent, reflect their phylogeny. At present the taxonomy of Basidiomycetes, to which most white-rot fungi belong, is undergoing substantial changes, particularly on the family level. Applying modern DNA sequencing methods will help to revise previously segregated heterogeneous taxa and introduce a phylogenetic classification of fungi (17). However, this new taxonomic concept is not yet fully established, and although it is unsatisfactory, in this paper we refer to traditional systematics (1, 6, 13, 45, 46). To increase our knowledge base, 10 white-rot fungi, previously referred to the family Polyporaceae (order Aphyllophorales), were selected and studied in terms of their ability to degrade synthetic lignin and produce ligninolytic enzymes in liquid culture. The Polyporaceae, which is considered to be a highly heterogeneous (17, 45, 46) family, includes many active lignin degraders, e.g., Bjerkandera adusta, Ceriporiopsis subvermispora, Cerrena unicolor, Ischnoderma resinosum, Pycnoporus cinnabarinus, Trametes gibbosa, Trametes hirsuta, Trametes versicolor (3-6, 17, 45, 46). Ligninolytic properties and enzymes of B. adusta (20, 23), Ceriporiopsis subvermispora (5, 43, 44, 47) and Trametes versicolor (18, 19, 36, 41) were recently investigated.

Mineralization of DHP by the fungi studied did not exceed 14% by day 29 of growth in air. This is notably lower than rate of mineralization we previously obtained with the LiPproducing corticioid fungi Phanerochaete chrysosporium, Phlebia radiata, and Phlebia tremellosa, which showed 35-40\% of mineralization within 30 days using the same radiolabelled substrate and growth conditions (54) but in oxygen. Oxygen was reported to influence DHP mineralization differently in various fungal species, strongly enhancing ¹⁴CO₂ evolution in the cultures of many fungi (15, 25), particularly in LiP-producers Phanerochaete chrysosporium (15, 25) and Phlebia radiata (15). On the other hand, the LiP-lacking polyporous fungus Ceriporiosis subvermispora was shown to mineralize DHP with approximately equal rates in both air and oxygen (44). In this work we used air in DHP-degradation experiments, since pure oxygen was found to prevent primary growth of some of the fungi in preliminary cultivation experiments. The influence of oxygen on lignin mineralization by various white-rot species evidently needs more investigation.

Another reason for the relatively low rates of DHP degradation obtained could be related to the different ligninolytic systems of fungi studied when compared with the corticioid species *Phanerochaete chrysosporium*, *Phlebia radiata*, and *Phlebia tremellosa*, as was shown by the earlier reported results on DHP mineralization by *Phanerochaete chrysosporium* versus *Ceriporiopsis subvermispora* in liquid cultures and in wood (47). These observations indicated that the DHP assay may not reflect the true ligninolytic activity of many white-rot fungi growing under natural conditions,

e.g., in wood substrates, that possess ligninolytic enzyme systems different from that of *Phanerochaete chrysosporium*. The DHP method was initially developed for liquid cultures of *Phanerochaete chrysosporium* (25), producing high titers of LiP under the growth conditions used (49, 50). Nevertheless, a great deal of valuable information e.g., on the physiology of fungal lignin degradation, has been obtained in this manner (10, 24).

Manganese was shown to play a regulative role in DHP degradation by Phanerochaete chrysosporium (38, 39) and some other white-rot fungi (7, 14, 21, 32, 40, 44, 54). High Mn2+ level in the presense of manganese chelators leads to the repression of ¹⁴CO₂ evolution and LiP production in Phanerochaete chrysosporium cultures (38, 39). In contrast, higher Mn²⁺ concentrations stimulate DHP mineralization in several fungal species producing MnP but lacking LiP activity, such as Dichomitus squalens (40), Ceriporiopsis subvermispora (44), and Pleurotus ostreatus (21). In this study manganese caused variable fungal specific effects on DHP mineralization. Evolution of 14CO2 in cultures of Trametes trogii and Trametes versicolor was inhibited by the 50-fold Mn²⁺ level. Both fungi produce rather high titers of LiP, as has been found for Trametes trogii here and reported previously for Trametes versicolor (18). In the other LiPproducing species, Trametes gibbosa, the influence of manganese remained unclear because of the generally very low DHP-degrading capacity demonstrated by this fungus under the growth conditions used. Stimulation of DHP mineralization by the Mn²⁺ could indicate the participation of MnP in the case of A. biennis, J. separabilima, and Trametes hirsuta. In the last two species stimulation by Mn²⁺ was found to be the highest in the presence of the chelator (sodium malonate). However, the addition of malonate triggered diverse responses in the fungi, showing clear inhibitory effects on DHP mineralization in the Mn2+-low cultures of A. biennis and Trametes trogii. These results indicate that malonate could be a self-regulatory factor in lignin degradation, irrespective of Mn2+, as was shown in the cultures of Phlebia radiata grown in air (32), in which the addition of malonate decreased 14CO2 evolution both under high- and low-manganese conditions. Despite the observation that in some cases the addition of malonate inhibited ¹⁴CO₂ evolution from the DHP-degrading cultures of the white-rot fungi, it did not notably reduce (or it even increased) the amount of water-soluble products derived from DHP by the fungi. A similar effect was previously reported for liquid cultures of Pleurotus ostreatus (21). A decrease of ¹⁴CO₂ evolution in the presence of high malonate concentration could be due to the known inhibitory effect of malonate on the tricarboxylic acid (TCA) cycle (48). Although malonate cannot penetrate the fungal cell membrane to a significant extent (9, 11), its permeability could be increased by forming molecular complexes of the ionophore type (11). Monitoring the fate of malonate during DHP-degradation experiments would be necessary to explain its physiological effects, including the possible metabolism of malonate by various fungi.

Different patterns of ligninolytic enzymes produced by the white-rot fungi grown under similar conditions in liquid cultures were observed even in taxonomically close species, e.g., among the fungi from the genus *Trametes*. Some of the investigated white-rot species (*Trametes pubescens*, *Trichap*tum abietinum, and G. applanatum) did not produce any notable ligninolytic activity and did not degrade synthetic lignin under the conditions studied, although in natural conditions they are known to be efficient lignin-degrading fungi (4, 6, 45, 46). Laccase was the most easily detectable enzyme produced in high levels by several species. Manganese peroxidase was also generally present in the fungi studied, and numerous MnP isoforms, similar to the pattern found in Ceriporiopsis subvermispora (43) and Trametes versicolor (18), were resolved in four species. Expression of MnP isoforms was obviously stimulated in high-manganese cultures supplemented with the chelator. Lignin peroxidase does not seem to be either widespread or easily produced in detectable titers in the fungi belonging to the Polyporaceae (14, 34), as demonstrated earlier in the corticioid fungi Phanerochaete chrysosporium (49, 50), Phlebia radiata (33), and Phlebia tremellosa (15, 54). The probable reason for this is the non-optimized growth conditions used. LiP and LiP encoding genes were characterized in B. adusta (20, 23) and Trametes versicolor (18, 19). In the latter fungus, multiple forms of LiP were described (19). In this work we purified LiP from Trametes gibbosa and Trametes trogii. Previously three LiP isoenzymes were isolated from J. separabilima (52), and now one MnP isoenzyme was detected in this fungus when grown under high Mn2+ concentration.

In conclusion, the results of this study showed that whiterot fungi are different in respect to their enzyme production patterns, DHP-degrading capacities, and regulation by Mn²⁺ or malonate. Extracellular peroxidases (LiP and MnP) were detected in DHP-degrading cultures of the fungi studied, although the role of these enzymes is still unclear. One group of fungi possesses LiP-dominating ligninolytic systems, the other group MnP-dominating ligninolytic systems, which involve either LiP or MnP in DHP mineralization. Fungi in the former group (Trametes versicolor and Trametes trogii) produced rather high levels of LiP, with the highest DHP mineralization occurring in the low-manganese medium, whereas DHP mineralization by the other fungi (A. biennis, J. separabilima, and Trametes hirsuta) was stimulated by Mn²⁺. However, although no definitive conclusions about the connection between ligninolytic systems of the white-rot fungi and their taxonomy or ecology can be drawn from this study, a recent investigation of more white-rot species (51) indicates that evolutionary LiP might be older compared with MnP. Evolution of fungal ligninolytic peroxidases needs further research using genetic methods. The more advanced and younger groups of fungi might utilize the MnP enzyme system for lignin degradation, but they might also have LiPencoding genes that are not expressed. Thus, biochemical data, such as characterization of ligninolytic enzymes, might contribute to the further development of fungal phylogenetic classification.

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