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Screening of white-rot fungi manganese peroxidases: a comparison between the specific activities of the enzyme from different native producers

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Abstract

In this study manganese peroxidase (MnP) enzymes from selected white-rot fungi were isolated and compared for potential future recombinant production. White-rot fungi were cultivated in small-scale in liquid media and a simplified process was established for the purification of extracellular enzymes.

Five lignin degrading organisms were selected (*Bjerkandera sp., Phanerochaete (P.) chrysosporium, Physisporinus (P.) rivulosus, Phlebia (P.) radiata* and *Phlebia sp. Nf* b19) and studied for MnP production in small-scale. Extracellular MnP activity was followed and cultivations were harvested at proximity of the peak activity. The production of MnPs varied in different organisms but was clearly regulated by inducing liquid media components (Mn2+, veratryl alcohol and malonate). In total 8 different MnP isoforms were purified.

Results of this study reinforce the conception that MnPs from distinct organisms differ substantially in their properties. Production of the extracellular enzyme in general did not reach a substantial level. This further suggests that these native producers are not suitable for industrial scale production of the enzyme. The highest specific activities were observed with MnPs from *P. chrysosporium* (200 U mg-1), *Phlebia sp. Nf* b19 (55 U mg-1) and *P. rivulosus* (89 U mg-1) and these MnPs are considered as the most potential candidates for further studies. The molecular weight of the purified MnPs was estimated to be between 45–50 kDa.

Keywords: White-rot fungi, Manganese peroxidase, Lignin degradation, Lignocellulose, Enzyme purification

Introduction

At present lignocellulose is a major raw material for forestry, pulp and paper industry and the emerging second generation biofuel production. Among cellulose and hemicellulose, lignin is a major component of lignocellulosic biomass and largely responsible for its strength. Inside the Northern coniferous forest belt the importance of lignin utilization is stressed in wood-based biorefineries due to high amounts of lignin in softwoods (Li *et al.* 2009).

Lignin is a heterogeneous, branched and complex polymer consisting of phenylalanine-derived aromatic subunits (Whetten & Sederoff 1995). Because of its recalcitrance, lignin complicates the utilization of biomass

Out of these enzymes MnP is thought to play the most crucial role in lignin degradation, as it is found in all lignin degrading fungi white-rot fungi. This heme protein

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polysaccharides in biorefineries and increases the energy consumption in mechanical pulping (Jiang *et al.* 2008a). In nature one group of organisms, the *basidiomycetous* fungi, are able to effectively degrade lignin by employing a family of lignin degrading enzymes. These organisms can be divided into wood-colonizing white-rot fungi and soil litter-decomposing fungi. Fungal attack on lignin is attributed to certain secreted nonspecific oxidoreductases, which produce low molecular weight mediators able to intrude recalcitrant biopolymers. The family of extracellular ligninolytic enzymes typically includes lignin peroxidases (LiP, EC 1.11.1.14), laccases (EC 1.10.3.2), manganese peroxidases (MnP, EC 1.11.1.13), versatile peroxidases (VP, EC 1.11.1.16) and other accessory enzymes.

belongs to the commonly occurring class II peroxidase group in basidiomycetous fungi and has a highly specific Mn2+ -binding site. In the binding site of classical long MnPs there are three amino-acid residues (two Glu and one Asp) while several fungal Mn2+ -oxidizing enzymes with an additional tryptophan residue on the enzyme surface have been found. These enzymes are called VPs or hybrid MnPs, bearing resemblance to LiPs and able to perform oxidation though long-range electron transfer as well. The evolution of these class II peroxidases appears to be closely related to each other and even consistent with the sharp decline in coal accumulation rate during the Permo-Carboniferous period. Lignin is the main precursor for coal (Hofrichter et al. 2010; Floudas et al. 2012). MnP catalyses the oxidation of Mn2+ -ions to highly reactive Mn3+-ions. Chelated Mn3+ in turn act as low molecular weight mediators that are able to attack phenolic structures. MnP is able to cause substantial depolymerization if lignin in in vitro biomass treatments (Hofrichter 2002; Hofrichter et al. 2001; Maijala et al. 2008).

Potential applications for MnP include biomechanical pulping, pulp bleaching, dye decolorization, bioremediation and production of high-value chemicals from residual lignin from biorefineries and pulp and paper side-streams (Maijala *et al.* 2008; Moreira *et al.* 2003; Moreira *et al.* 2001; Xu *et al.* 2010a and 2010b; Paice *et al.* 1993; Susla *et al.* 2008; Hofrichter *et al.* 1998; Sack *et al.* 1997). Applications of MnP are limited due to slow growth and low productivity of native enzyme producers and lack of an efficient recombinant production process (Hofrichter 2002; Jiang *et al.* 2008a).

The production of lignolytic enzymes and its regulation has been intensively studied in various lignin degrading fungi (Bonnarme & Jeffries 1990; Hakala et al. 2006; Jiménez-Tobon et al. 2003; Kamitsuji et al. 2004; Lankinen et al. 2005; Martínez et al. 1996; Moilanen et al. 1996; Nuske et al. 2002; Perie & Gold 1991; Palma et al. 2000; Petruccioli et al. 2009; Schneegab et al. 1997; Swamy & Ramsay 1999; Steffen et al. 2002; Susla et al. 2008; Silva et al. 2008; Sklenar et al. 2010; Singh et al. 2011; Taboada-Puig et al. 2011; Vares et al. 1995; Wang et al. 2001; Wang et al. 2008) and novel bacterial strains (Bharagava et al. 2009; Mishra & Thakur 2010; Yadav et al. 2011). Recombinant production has been studied in filamentous fungi (Conesa et al. 2000; Irie et al. 2001; Li et al. 2001; Mayfield et al. 1994; Stewart et al. 1996), yeasts (Jiang et al. 2008a), bacterial (Whitwam & Tien 1996) and insect (Johnson et al. 1992; Pease et al. 1991) hosts with successful production but modest yields of active enzyme. MnP from P. chrysosporium has been the target of most recombinant studies, which however suffer from unsuccessful posttranslational protein modification and the need for exogenous heme in high concentrations (Jiang et al. 2008a).

Furthermore, industrially robust enzymes need to have high stability in demanding process conditions which promotes the need for screening novel enzymes and enzyme modification. Multiple crystal structures and molecular models based on gene sequences have been published for MnP and VP (e.g. Sundaramoorthy et al. 1997; Sutherland et al. 1997 and Moreira et al. 2005). The effects of disulfide bonds and calcium-ions on the stability of MnP have been studied by Reading et al. (2001) and Sutherland et al. (1997), respectively. As a result of enzyme modification, a mutant MnP with one additional disulfide bond had increased stability in alkaline (pH 8) conditions (Reading et al. 2001). Another strategy is to screen for native MnPs with improved properties regarding thermostability, specific activity or pH optimum and stability (Petruccioli et al. 2009 and Urek & Pazarlioglu 2004). Increased specific activity (U mg-1 protein) of the selected protein would also increase the profitability of the recombinant process. Independent of the strategy the bottleneck is the establishment of an efficient recombinant production.

The aim of the present study was to screen MnPs from promising candidates of lignin degrading fungi for future recombinant production. Specific activities of MnPs were determined after cultivation and purification procedures. Isoforms of MnP from different organisms differ substantially and screening for novel MnP enzymes is beneficial for designing applications and developing a production process capable of producing the enzyme at an economically feasible cost. However high-throughput screening is challenging due to low enzyme yields and diverse regulation of MnP isoenzymes in different native hosts.

Materials and methods

Fungi and culture conditions

White-rot fungi strains *Bjerkandera sp.* BEL LLP4 (D-00810) and *P. radiata* Hatakka & Pirhonen strain 79 (ATCC 64658, D-84236) were ordered from the VTT Technical Research Centre of Finland culture collection, Finland. Strains *P. chrysosporium* (ATCC 24725, DSM 6909), *Phlebia sp. Nf* b19 (ATCC 201144, DSM 11239) and *P. rivulosus* T241i (DSM 14618) were ordered from the Leibniz institute DSMZ-German collection of microorganisms and cell cultures GmbH, Germany. *Phlebia sp. Nf* b19 was originally identified as *Nematoloma frowardii* species but reassigned to the family Corticiaceae and genus *Phlebia* based on a molecular level study by Hildén *et al.* (2008).

The fungi were maintained on solid agar plates. The agars used were 0.4% potato 2% dextrose agar (Difco, USA) for *P. radiata*, 3% malt extract-peptone agar (Merck, Germany) for *P. chrysosporium*, *Phlebia sp. Nf* b19 and *P. rivulosus* and 3% malt agar (Difco, USA) for *Bjerkandera sp.* The plates were incubated for 2–10 days

until covered with mycelium. Plates were stored at 4°C and replated monthly.

The inocula were prepared by suspending the mycelia from one plate to sterile 0.9% (w/v) Sodium chloride. One plate was used to inoculate two parallel fungicultivations.

The production of extracellular MnP in each organism was followed in two parallel cultivations in 1 L shake flasks containing 300 mL of liquid media with 90 rpm agitation. Cultivation temperatures were 37°C for P. chrysosporium, 25°C for Phlebia sp. Nf b19 and 28°C for all other whiterot fungi strains. P. radiata was grown in low-nitrogen ADMS medium pH 4.5 (Hatakka & Uusi-Rauva 1983) with 1% glucose, 0.05% (w/v) Tween 80 and 1 mM veratryl alcohol (VA, 3,4-dimethoxybenzyl alcohol). Supplements 180 µM Mn2+ and 10mM Sodium malonate were added on the 4th day. P. rivulosus was grown in low-nitrogen ADMS medium pH 4.5 (Hatakka & Uusi-Rauva 1983) with 1% glucose and 0.05% (w/v) Tween 20. Supplements 24 µM Mn2+ and 0.36 mM VA were added on the 4th day. P. chrysosporium was grown in Kirk's medium pH 4.5 (Urek & Pazarlioglu 2004; Tien & Kirk 1988 and Bonnarme & Jeffries 1990) with 1% glucose and 0.05% (w/v) Tween 80. Supplements 728 µM Mn2+ and 0.36 mM VA were added on the 4th day. Bjerkandera sp. was grown in Kirk's medium pH 4.5 (Palma et al. 2000) with 1% glucose and 0.05% (w/v) Tween 80, 235 μM Mn2+ and 0.36 mM VA. Phlebia sp. Nf b19 was grown in a glucose-yeast extract medium pH 4.5 (Nuske et al. 2002) with 0.5% glucose, 0.03% yeast extract and 200 µM Mn2+. Cultivations were continued until extracellular MnP activity was detected and it started to decline.

Sampling

1mL samples were taken every 24 h from each flask. Samples were centrifuged to remove cellmass (16000 \times g, 4°C). The supernatant was used for determination of MnP activity and glucose and total protein concentrations. All of the cultivations were performed in duplicate and the presented results are average values.

Enzyme assay and analytical procedures

Glucose concentration during cultivation was determined using the YSI 2700 SELECT $^{\text{\tiny TM}}$ (YSI limited, UK, Hampshire) biosensor.

MnP activity was determined spectrophotometrically at 270 nm by following the formation of Mn3+ –malonate complex at pH 4.5 in 50 mM sodium malonate buffer with 0.5 mM MnSO4. The reaction was initiated by adding H2O2 to the final concentration of 0.1 mM (Wariishi et al. 1992). The reaction was followed for 30 sec at room temperature. Δ Abs min-1 was converted to U L-1 using an extinction coefficient of 11590 M-1cm-1. Genesys

10UV spectrometer (Thermo Scientific, USA) was used in all the measurements with semi-micro 1.4 mL UV quartz cuvettes (Sigma-aldrich, Germany)

Total protein was determined with the Bio-Rad protein Assay (USA) using bovine serum albumin (BSA) as a standard. Measurements were made with Genesys 10UV spectrometer (Thermo Scientific, USA) at 595 nm wavelength.

Protein expression and purity was followed using sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) with silver nitrate staining. Ready-made Bio-Rad (USA) ready GelTM precast gels (12% Tris-HCl) were used with Bio-Rad (USA) prestained low range protein standards.

Purification of manganese peroxidases

Cellmass was separated from the culture fluids by centrifugation at $16000 \times g$, $4^{\circ}C$ (Avanti J series centrifuge, Beckman Coulter Inc, USA). Supernatant was then filtered through a Whatman grade 1 filter paper (11 μ m, Whatman, UK). Filtrate was then stored in $-20^{\circ}C$ until further processing.

Concentration of the culture fluids from the screening cultivations was done by using Vivaspin 20 centrifugal concentrators with 10 kDa cut-off membranes (GE Healthcare, USA and Sartorius Stedim Biotech, Germany). Crude culture liquids were concentrated approximately tenfold by volume. Megafuge 1.0R (Heraeus, England) was used for centrifugation at $3400 \times g$, 4°C. Diafiltration was performed in Vivaspin tubes by filtering 20 mM Bis-TRIS-propane buffer (pH 6.2) through the membrane (three times the volume of the concentrate).

ÄKTAavantTM liquid chromatography system (GE Healthcare, USA) equipped with a strong anion exchange column (HiScreenTM CaptoTM Q, GE Healthcare, USA) was used for protein separation. The outflow was monitored at two wavelengths: 280 nm for protein and 405 nm for hemeprotein detection. The column was first equilibrated with 20 mM Bis-TRIS-propane buffer (pH 6.2) and proteins were eluted with a linear gradient of 0–1 M NaCl in the same buffer. 2 mL fractions were collected during the elution phase. The fractions were assayed for MnP activity and total protein. Peaks were pooled separately and stored at –20°C.

Results

MnP production in selected white-rot fungi

The activity (U l-1) of MnP enzyme produced and the time period it took to reach the maximal activity varied greatly between the selected white-rot fungi. A unifying trend of accelerated glucose consumption just before and during MnP production was observed in these batch cultivations (Figure 1A & 1B). The MnP peak timing and maximal activity varied between 7–21 days and

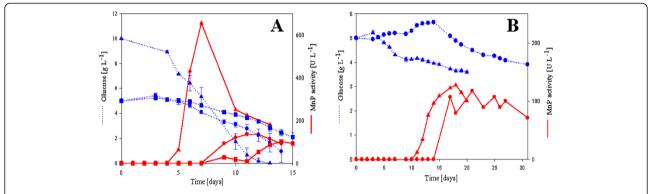


Figure 1 Glucose consumption (dotted line) and MnP activity (solid line) by white-rot fungi during 300 mL batch cultivations. Each strain was cultivated in specific conditions and media described in the materials section. Values are calculated averages of two parallel cultivations. Standard deviation of MnP activity in parallel cultures is on average ±40 UL⁻¹. *Bjerkandera sp.* (**A**, blue and red triangles), *Phanerochaete chrysosporium* (**A**, blue and red circles), *Physisporinus rivulosus* (**A**, blue and red squares), *Phlebia radiata* (**B**, blue and red triangles), *Phlebia sp.* Nf b19 (**B**, blue and red circles). Harvesting point is designated by the end of sampling.

100–800 U l-1, respectively. Fastest growing white-rot fungus in these experiments was *Bjerkandera sp.*, which also produced the highest activities. However, the MnP activity of *Bjerkandera sp.* increased and subsequently declined very sharply. These cultures were harvested 6 days after the MnP activity peak, which may have resulted in inactivated protein and lowered specific activity.

Purification of MnP isoenzymes from crude culture filtrates

Crude culture filtrates were concentrated and purified in a rapid purification scheme described in the materials section. Based on the SDS-PAGE gels (Figure 2A-E) MnP was the predominant protein produced in most of the cultures and highly purified in the pooled anion exchange fractions (Figures 3, 4, 5, 6, 7). In the Phlebia sp. Nf b19 culture supernatant and concentrate, a larger enzyme (around 70 kDa) was the predominant one. Based on the size, this is probably a laccase. Loss of MnP activity was high during the purification process. MnP activity was lost especially in the concentration step, but this is not relevant for the determination of specific activity. The role of enzyme inactivation in the purification procedure is minimized due to low kept temperatures (4°C) during each step. After the strong anion exchange (Mono Q column) one MnP isoenzyme for Bjerkandera sp. (Figure 3), Phlebia sp. Nf b19 (Figure 7) and P. rivulosus (Figure 5); two MnP isoenzymes for P. chrysosporium (Figure 4) and possibly three isoenzymes for P. radiata (Figure 6) were detected. All of the MnP active fractions showed absorbance peaks in the 280 nm wavelength for protein and in the 405 nm wavelength for the heme containing protein. Wavelenght 405 nm could easily be used for selecting fractions likely to show MnP activity. MnP proteins were eluted between 175 and 320 mM NaCl concentrations.

The molecular weight of the MnPs was between 45 and 50 kDa (Table 1). Specific activities of the MnPs were calculated on the basis of the activity measurements and measured total protein in the pooled active fractions. *P. chrysosporium* MnP1 had the highest specific activity (200 U mg-1). The second highest specific activity was observed for *P. rivulosus* (89 U mg-1). *Phlebia sp. Nf* b19 MnP had the specific activity of 55 U mg-1. Rest of the MnP isoenzymes had a specific activity below 40 U mg-1 (Table 1).

Preliminary results of *in vitro* enzymatic processing of milled pinewood sawdust with crude MnP concentrates showed up to 17% (data not shown) decrease in Klason lignin content and benefitted from the use of co-oxidants (Tween 80 and glutathione) in some cases.

Discussion

These results show that a simplified purification process is capable of producing comparable specific activity results for the screening of novel MnP enzymes. Culture supernatant concentration and anion exchange chromatography using a Mono Q column was enough to purify the enzyme to a high degree. Although the initial enzyme amounts were low and the losses during concentration were quite high, these results provide a basis for enzyme ranking. Based on these results there are significant differences in the specific activities of MnP enzymes from different white-rot fungi. Out of the white-rot fungi used in this study P. chrysosporium, P. rivulosus and Phlebia sp. Nf b19 MnPs were the most promising. For P. rivulosus and Phlebia sp. Nf b19 these results are also consistent with the MnP activity and protein absorbance (280 nm) ratio in anion exchange chromatography (Table 1). The MnP activity (U l-1) of Bjerkandera sp. had come down over 70% at the point of harvesting. If the lost activity is accounted as inactivated MnP in the total protein

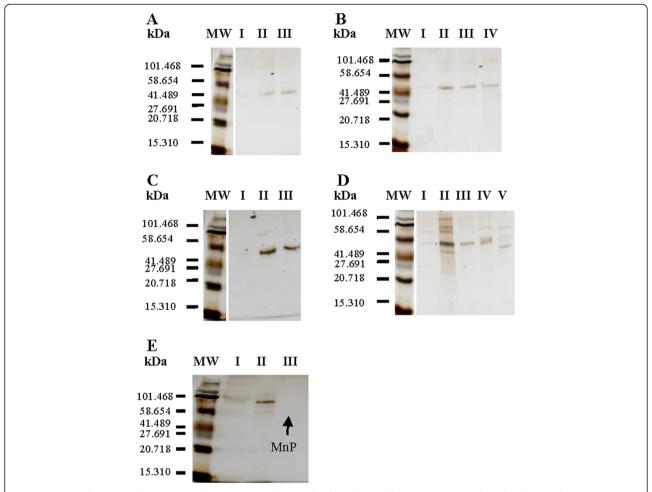


Figure 2 (A-E) Silver stained SDS-PAGE gels in corresponding order from the purification process with molecular weight protein standards (MW), crude extracellular fluids (I), concentrated samples (II) and MnP isoenzymes from MonoQ anion exchange column (III-V).

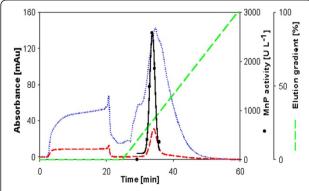


Figure 3 Anion exchange chromatogram from the purification of MnP from *Bjerkandera sp.* Dotted line is the absorbance at wavelength 208 nm; Short dash line is the absorbance at wavelength 405 nm; Long dash line is the concentration of the eluent in percentage.

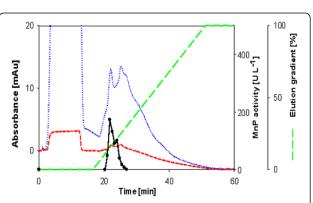


Figure 4 Anion exchange chromatogram from the purification of MnP from *Phanerochaete chrysosporium*. Symbols as in Figure 2.

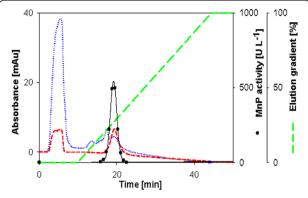


Figure 5 Anion exchange chromatogram from the purification of MnP from *Physisporinus rivulosus*. Symbols as in Figure 2.

measurement, the specific activity could be at least three-fold higher. In addition to specific activity the stability of the purified MnP enzymes towards temperature, pH and inactivating compounds needs to be studied before candidates for recombinant production are selected. The reported specific activities are significantly lower than those previously reported for *P. chrysosporium* by Urek *et al.* (2004), *Bjerkandera sp.* by Palma *et al.* (2000) and Taboada-Puig *et al.* (2011) and for *Phlebia sp. Nf* b19 by Schneegaβ *et al.* (1997)

The production of MnP by native producers, as also observed in this study, is limited due to relatively low maximal enzyme activities, slow growth and the sensitivity of white-rot fungi towards shear forces, difficult induction strategies and low adaptivity to submerged fermentations. In this study MnP was in most cases induced with several supplements (Mn2+, VA, Tween, Sodium malonate). However, as reported by Hakala *et al.* (2006), the regulation of different MnP isoforms can be largely dependent on the inducing compounds and nutrient (nitrogen and carbon) sources and amounts. This suggests that novel isoforms would be found by changing the culture conditions. Mn2+-ions are typically necessary inducers for

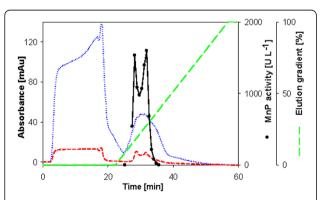


Figure 6 Anion exchange chromatogram from the purification of MnP from *Phlebia radiata.* Symbols as in Figure 2.

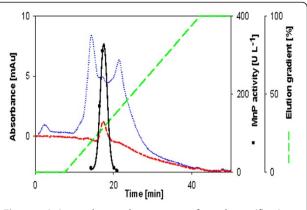


Figure 7 Anion exchange chromatogram from the purification of MnP from *Phlebia sp. Nf* b19. Symbols as in Figure 2.

MnP production although repressive effects of Mn2+-ions have also been reported (Martinez *et al.* 1996). Although high activity level was not the focus in this study, the culture conditions and inducing strategies (see the materials section) for this study were selected on previous literature to maximize enzyme production. Specified effects of individual inducing components cannot be separated for any strain, but clearly no MnP was produced before inducing components were added.

In recombinant MnP processes with Pichia pastoris (Jiang et al. 2008b), Aspergillus oryzae (Stewart et al. 1996), Aspergillus niger (Conesa et al. 2000) and Echerichia coli (Whitwam et al. 1995; Whitwam & Tien 1996) production of high amounts of active protein has been a problem. These processes suffer from incorrect protein folding, insufficient heme synthesis and heme incorporation. However, MnP yield might be significantly increased with the right combination of recombinant enzyme, production host, promoter system, protein secretion system and optimized process conditions. Jiang et al. (2008b) also suspected that P. pastoris lacks the proper heme escorts and reseptors to transport exogenous heme to the site of MnP synthesis. This problem might be relieved by inserting the genes for such heme transport from e.g. Shigella (S.) dysenteriae (Mills & Payne 1995). S. dysenteriae is a known pathogen causing shigellosis, which would make the GMO rating of the recombinant strain difficult. On the other hand *P. pastoris* synthesizes large amounts of a homologous heme-containing catalase protein under alchohol oxidase (AOX) promoted recombinant expression. Thus gene expression under AOX promoter might be beneficial for MnP production due to the added need for heme synthesis inside the cells. Coprinus cinereus produces a class II excreted fungal peroxidase (CiP) that readily oxidizes phenols, but is unable to oxidize veratryl alcohol or Mn2+-ions (Hofrichter et al. 2010). This enzyme is successfully produced in a recombinant process using P. pastoris (Kim et al. 2009a) with high

Table 1 Purification of MnP isoenzymes from *Bjerkandera sp.* (A), *Phanerochaete chrysosporium* (B), *Physisporinus rivulosus* (C), *Phlebia radiata* (D) and *Phlebia sp.* Nf b19 (E)

	Activity [U]	Protein [mg]	Specific activity [U/mg]	Activity protein absorbance ratio [UL ⁻¹ absorbance at 280nm ⁻¹]	Yield [%]	Purification fold	Size of MnP [kDa]
A					,		
Culture fluid	38.0	2.9	13.3		100	1.0	
Ultrafiltration	19.8	2.6	7.6		52.2	0.6	
Mono Q BSMnP1	15.3	0.5	32.3	18	40.2	2.4	45
В							
Culture fluid	38.7	1.447	26.7		100	1.0	
Ultrafiltration	6.9	0.403	17.0		17.7	0.6	
Mono Q PCMnP1	1.1	0.005	200.5	13	2.8	7.5	45
Mono Q PCMnP2	0.4	0.013	33.0	8	1.1	1.2	45
С							
Culture fluid	36.6	2.685	13.6		100	1.0	
Ultrafiltration	7.7	0.278	27.6		20.9	2.0	
Mono Q PRMnP1	2.3	0.026	88.6	122	6.2	6.5	50
D							
Culture fluid	52.0	4.669	11.2		100	1.0	
Ultrafiltration	24.6	2.713	9.1		47.2	0.8	
Mono QPradMnP1	5.8	0.245	23.7	33	11.1	2.1	50
Mono Q PradMnP2	4.0	0.199	19.9	21	7.6	1.8	50
Mono Q PradMnP3	7.2	0.234	30.8	33	13.8	2.8	50
E							
Culture fluid	7.1	0.671	10.6		100	1.0	
Ultrafiltration	3.4	0.511	6.6		46.9	0.6	
Mono Q NFMnP1	0.7	0.012	55.2	67	9.3	5.2	49

productivities (peroxidase activity over 1200 U ml-1 and total protein over 1.6 g l-1) and is also sold as a commercial enzyme by Novozymes (Baylase®). The productivity of this heme-containing peroxidase was optimized by host and expression promoter selection (Kim *et al.* 2009b). Highest productivities were obtained by using the AOX promoter with a fast methanol utilization strain (Mut+) of *P. pastoris*. This supports the theory, that inducing the methanol utilization pathway in a production host can promote the recombinant production of heme-containing peroxidases. Commercially available MnP (from *Phlebia sp. Nf* b19) and VP (from *Bjerkandera adusta*) from Jena Bioscience GmbH are native enzymes and overly expensive for any kind of industrial use.

In this study several well known MnP enzymes were compared. MnP and laccase and their regulation in *P. rivulosus* has been well characterized by Hakala *et al.* (2005 & 2006). The Differential regulation of MnP isoforms in *P. rivulosus* was also noted in these articles. In this study only one MnP from *P. rivulosus* is characterized, but this may very well be a group of MnP isoenzymes with approximately the same isoelectric points

and molecular sizes. In characterization studies by Hildén et al. (2008), the MnP2 enzyme from Phlebia sp. Nf b19 showed a 96% amino acid identity to the MnP2 enzyme of *P. radiata* in the primary structure. In this study the specific activities and sizes of the MnP enzymes produced by these related fungal strains were in the same range. Even in growth and MnP production these strains showed similarities (extremely slow growth and late onset of MnP production). Hildén et al. (2005) describe two MnP enzymes from P. radiata that are different in their primary structure, intron amount, length and crystal structure. The other isoenzyme being structurally related to LiP, but having an alanine residue instead of tryptophan while still having a conserved Mn2+-binding site. In this study three MnPs from P. radiata were separated, but they seemed to be highly similar in size and specific activity for Mn2+ oxidation.

Many technical applications for MnP have been reported with promising results. The utilization of the enzyme is still dependent on a cost-effective recombinant production process and possibly the discovery of more robust novel isoenzymes or modifications of the currently known ones. Furthermore, optimization of enzymatic treatment processes for various technical lignins (Vishtal & Kraslawski 2011), paper pulps (Maijala et al. 2008; Xu et al. 2010) and organopollutants (Sack et al. 1997) with proper process conditions and co-oxidants will probably increase the interest in MnP. Delignification of pinewood sawdust using a MnP treatment in this study was relatively inefficient. The use of several co-oxidants and other enzymes involved in biological ligninolysis may help to achieve more thorough enzymatic delignification demonstrated by many MnP producing and LiP-negative white-rot fungi (Hammel & Cullen 2008). In previous laboratory experiments by Hofrichter et al. (1998) and Kapich et al. (1999) Mineralization and solubilization of synthetic (14C-labeled) large molecular weight lignin by MnP has been reported. These and various other studies suggest that isolated MnP enzymes can be used to delignify biomasses. However, for now the utilization of class II peroxidases to degrade polluting substances in soils is technically more appealing.

Competing interests

The authors declare that they have no competing interests.

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