FULL PAPER

Mizuho Kusuda · Mitsuhiro Ueda · Yasuhito Konishi Yoshihito Araki · Katsuji Yamanaka · Masami Nakazawa Kazutaka Miyatake · Takao Terashita

Detection of β -glucosidase as saprotrophic ability from an ectomycorrhizal mushroom, *Tricholoma matsutake*

Received: September 16, 2005 / Accepted: December 26, 2005

Abstract We investigated extracellular carbohydrase production in the medium of an ectomycorrhizal fungus, Tricholoma matsutake, to reveal its ability to utilize carbohydrates such as starch as a growth substrate and to survey the saprotrophic aspects. We found β -glucosidase activity in the static culture filtrate of this fungus. The β -glucosidase was purified and characterized. The purified enzyme was obtained from about 2.11 static culture filtrate, with 9.0% recovery, and showed a single protein band on SDS-PAGE. Molecular mass was about 160kDa. The enzyme was most active around 60°C and pH 5.0, and stable over a pH of 4.0– 8.0 for 30 min at 37°C. The purified enzyme was activated by the presence of Ca²⁺ and Mn²⁺ ions (about 2–3 times that of the control). The enzyme readily hydrolyzed oligosaccharides having a β-1,4-glucosidic linkage such as cellobiose and cellotriose. However, it did not hydrolyze polysaccharides such as avicel and CM-cellulose or oligosaccharides having an α-glucosidic linkage. Moreover, cellotriose was hydrolyzed by the enzyme for various durations, and the resultant products were analyzed by TLC. We concluded that the enzyme from T. matsutake seems to be a β -glucosidase because cellotriose with a β-1,4-glucosidic linkage decomposed to glucose during the enzyme reaction.

Key words Amylase \cdot β -Glucosidase \cdot Edible mushroom \cdot Mycorrhizal fungus \cdot *Tricholoma matsutake*

M. Kusuda \cdot M. Ueda \cdot M. Nakazawa \cdot K. Miyatake Laboratory of Biocycle Engineering, Graduate School of Life and Environmental Science, Osaka Prefecture University, Osaka, Japan

Y. Konishi · Y. Araki · T. Terashita (☒) Laboratory of Food Microbiological Science and Biotechnology, Faculty of Agriculture, Kinki University, 3327-204 Naka-machi, Nara 631-8505, Japan

K. Yamanaka Kyoto Mycological Institute, Kyoto, Japan

Introduction

The ectomycorrhizal mushroom *Tricholoma matsutake* (S. Ito et Imai) Singer is one of the most valuable edible mushrooms in Japan. However, artificial cultivation of the mushroom has not been established in a stable condition. This fungus can grow slowly on starch and inulin when a small amount of glucose is added as a starter (Norkrans 1950). The starch, which is used as a carbon source, is able to supply the conditions that allow a successful quantity of mycelial growth without raising osmotic pressure in the medium.

To reveal the ability to utilize starch and related substrates of T. matsutake, the production systems of extracellular amylase were studied using an artificial medium. As a result, it was shown that this fungus produced α -amylase and α -glucosidase in static culture filtrate. α -Amylase in these enzymes was already purified, and its enzymatic property was characterized from the aspect of substrate specificities. The purified α -amylase produced oligosaccharides from starch. However, α -glucosidase activity was only partially purified because the activity was very weak.

On the other hand, Vaario et al. (2002) reported that T. matsutake produced β -glucosidase when they used pine bark for cultivation of this fungus. In addition, they observed by transmission electron microscopy that T. matsutake decomposed sawdust cells of pine bark by the kind of strain and the fungus mycelia invaded the sawdust cell of pine bark (Vaario et al. 2003). We recently also detected a β -glucosidase having high activity from a static culture filtrate of this fungus. These results suggest that T. matsutake has saprotrophic abilities. However, no characterization of the highly purified enzyme from T. matsutake has been reported.

In this article, we purified and characterized β -glucosidase from this ectomycorrhizal mushroom and discuss possible roles of this extracellular enzyme of *T. matsutake*.

Material and methods

Organism

Tricholoma matsutake Z-1 strain was used in this experiment. This strain was isolated from the fruit-body of *T. matsutake* in 1985 by Inaba et al. (1995). The fungus was stored on partially modified matsutake agar medium (PMMA) (Terashita et al. 2000a) at a low temperature (4°C).

Medium compositions and culture conditions

A mycelial block ($5 \times 5\,\text{mm}$) was cut from a plate culture that had grown on PMMA (1.5% agar powder) medium (Terashita et al. 2000a) for 40 days at 24°C in a petri dish (diameter, $90\,\text{mm}$). It was inoculated in an Erlenmeyer flask ($100\,\text{ml}$) containing $20\,\text{ml}$ PMM liquid (PMML) medium after sterilization at 121°C for $5\,\text{min}$. The composition of the PMML medium was $22.7\,\text{g}$ glucose, $5.0\,\text{g}$ yeast extract, $77.0\,\text{g}$ potato extract, and $5.0\,\text{g}$ Sunpearl-CP (commercial name; sulfite pulp waste from softwood) per $1000\,\text{ml}$ distilled water, with an initial pH of 5.1. After inoculation, it was static cultured at 24°C for 60 days in the light (about $200\,\text{lux}$).

Extraction of crude enzyme

For purification of the extracellular β -glucosidase, 60 days after incubation the culture medium was filtered through no. 2 filter paper (Advantec, Osaka, Japan) for mycelial removal. The culture filtrate was used as the crude enzyme for starting materials for purification.

Measurement of α - and β -glucosidase activity

α-Glucosidase and β-glucosidase activity was assayed using p-nitrophenyl- α - and β-glucopyranoside (p-NαG and p-NβG) as the substrates. The enzyme reaction was carried out at 37°C for 30 min. One unit of α - and β-glucosidase activity was defined as the activity that forms 1 μmol p-nitrophenol in 1.0 ml reaction mixture at 37°C in 1 min.

Purification procedures of extracellular β -glucosidase

Ammonium sulfate was added to make a 60% saturation of the culture filtrate (2096 ml). The precipitate by ammonium sulfate was then separated by centrifuge (9000 g, 20 min). The precipitate was dissolved in 20 mM Tris-HCl buffer containing 10 mM CaCl₂ (pH 7.0) and then was used as the starting material for the enzyme purification.

All purification steps were carried out at 4°C. The crude enzyme solution (520 ml) was dialyzed (in 20 mM Tris-HCl buffer containing 10 mM CaCl₂, pH 7.0, overnight, 4°C) and put on a Toyopearl-DEAE 650 M column (Tosoh, Tokyo, Japan) that was previously equilibrated with 20 mM Tris-HCl buffer containing 10 mM CaCl₂ (pH 7.0). The

enzyme was eluted with a linear gradient of Tris-HCl buffer (pH 7.0) from 20 to $500\,\text{mM}$ at a flow rate of $0.5\,\text{ml/min}$. The β -glucosidase activity of each fraction was measured.

The fractions containing β -glucosidase (fractions 25–50; total volume, 151.5 ml) from the Toyopearl-DEAE column were concentrated at 4°C under reduced pressure to about 200 μ l. The concentrated solution was charged on a column of Superdex 200 equilibrated with 20 mM Tris-HCl buffer (pH 7.0) containing 20 mM CaCl₂, pH 7.0. Elution was performed with the same buffer at a flow rate of 250 μ l/min, and 250- μ l fractions were collected.

Enzyme homogeneity and estimation of molecular mass by SDS-PAGE

Protein homogeneity of the purified enzyme was measured by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) by the method detailed by Laemmli (1970), using a mixture of 0.1% SDS and 100mM Trisglycine buffer, pH 8.3, at 15mA for 5h. The apparent molecular mass was calculated from the mobility versus molecular mass plots of the marker proteins (myosin, 200.0kDa; β -galactosidase, 116.2kDa; bovine serum albumin, 66.3kDa; aldolase, 42.4kDa; carbonic anhydrase, 30.0kDa; myoglobin, 20.1kDa). Protein bands on the gel were stained with Coomassie brilliant blue after SDS-PAGE.

Effect of pH and temperature

The temperature dependence of β -glucosidase activity was measured in a 0.1 M McIlvaine buffer, pH 5.0, for 30 min. The thermal stability of this enzyme was investigated by incubating it in a 0.1 M McIlvaine buffer, pH 5.0, for 30 min at various temperatures. The effect of pH on enzyme activity was measured using a 20 mM p-N β G solution as the substrate at 37°C for 30 min. The effect of pH on the stability of β -glucosidase activity was investigated by measurement of the remaining activity after incubation for 30 min at 37°C in a buffer solution with pH values from 3.0 to 8.0.

Michaelis constant

p-Nitrophenyl- β -D-glucopyranoside and cellobiose were incubated at various concentrations with purified β -glucosidase; then, the initial rate of p-nitrophenol and glucose formation was measured.

Effect of metal ions

The β -glucosidase activity was assayed with various metal ions and ethylene diamnetetraacetic acid (EDTA) by the *p*-nitrophenol method (Kusuda et al. 2003).

Table 1. Purification of extracellular β-glucosidase from *Tricholoma matsutake* Z-1

Purification steps	Total protein (mg)	Total activity (U)	Specific activity (U/mg)	Purification (-fold)	Yield (%)
Culture filtrate 60% (NH ₄) ₂ SO ₄ -sat. ppt.	23 456	133	0.006	1	100
	1 505	152	0.101	18	114
Toyopearl-DEAE	14.3	36	2.52	442	27
Superdex 200	0.2	12	63.89	11 200	9

Substrate specificities

The substrate specificity of β -glucosidase was tested using synthetic substrates (p-N β G, p-N α G, and p-nitrophenyl β -D-cellobioside) and natural substrates (disaccharides; different degrees of oligosaccharides and polysaccharides: maltose, cellobiose, isomaltose, sucrose, lactose, trehalose, cellotriose, isomalto oligosaccharide, dextrin, CM-cellulose, avicel, curdlun). The enzyme activity was measured by the p-nitrophenol method for synthetic substrates and the F-kit D-glucose method (Kusuda et al. 2004) for natural substrates. Moreover, cellotriose was hydrolyzed by β -glucosidase at various durations, and the resultant products were analyzed by thin-layer chromatography (TLC).

Results

Purification procedure

The extracellular β -glucosidase from T. matsutake that shows activity at about pH 5.0 was purified using the crude enzyme solution prepared from a static culture filtrate (about 2100 ml, for 60 days cultivation) as the starting material. In the final purification step, using a Superdex 200 column chromatography (Fig. 1), β -glucosidase was recovered as a single and symmetrical protein peak (Fr. no. 45–52). However, a small peak (Fr. no. 53–55) having α -glucosidase activity appeared. The overall process for the purification of β -glucosidase is summarized in Table 1. The purified extracellular β -glucosidase represents about 11 200-fold purification over the original culture filtrate with about 9% recovery. The homogeneity of the purified enzyme was ascertained by SDS-PAGE. The purified β -glucosidase gave a single sharp protein band on SDS-PAGE (Fig. 2).

Molecular mass

The molecular mass of purified amylase was about 160kDa by SDS-PAGE.

Effect of pH and temperature

The results are shown in Table 2. The optimal temperature of the enzyme activity toward p-N β G was 60°C; the thermal stability of this enzyme was at 40°C. The enzyme exhibited

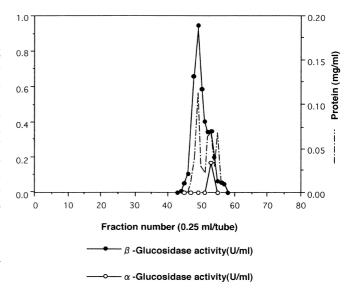


Fig. 1. Gel filtration profiles of β-glucosidase from *Tricholoma matsutake* Z-1 by Superdex 200 column chromatography. Column, Superdex 200 column (Pharmacia, 10×300 mm); flow rate, 0.25 ml/min; fraction size, 0.25 ml/tube; elution, 20 mM Tris-HCl buffer (pH 7.0) containing 20 mM CaCl₂

maximum activity at pH 5.0 and was stabilized at pH 4.0–8.0 after incubation at 37°C for 30 min.

Michaelis constant

A Lineweaver–Burke plot of β -glucosidase activity with p-N β G and cellobiose as substrates was obtained. As shown in Table 2, the $K_{\rm m}$ values for p-N β G and cellobiose were calculated to be 0.22 mM and 0.99 mM for this β -glucosidase. Conversely, the $V_{\rm max}$ values for both were 2.89 mM/min (p-N β G) and 3.59 mM/min (cellobiose), respectively.

Effect of metal ions

The effects of metal ions are shown in Table 3. β -Glucosidase activity was remarkably activated by the presence of Ca²⁺ (287%) and Mn²⁺ (257%). Among the metal ions tested, Ba²⁺, Mg²⁺, Zn²⁺, Pb²⁺, Na⁺, Cu²⁺, Li⁺, Co²⁺, Fe²⁺, Al³⁺, and Hg²⁺ and EDTA were in some degree activators for β -glucosidase. In contrast, K⁺ and Ag⁺ weakly inhibited the activity.

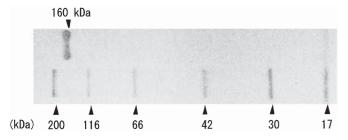


Fig. 2. Sodium dodecyl sulfate-polyacrylamide gelelectrophoresis (SDS-PAGE) of purified extracellular β-glucosidase from *T. matsutake* Z-1. Molecular weight markers: 200kDa, myosin; 116kDa, β-galactosidase; 66kDa, bovine serum albumm (BSA); 42kDa, ardolase; 30kDa, carbonic anhydrase; 17kDa, myoglobin; *arrow*, protein band of β-glucosidase

Table 2. Some properties of extracellular β -glucosidase from *T. matsutake* Z-1

Molecular weight (SDS-PAGE) pH stability (37°C, 30 min) Optimal pH (37°C) Thermal stability (pH 5.0, 30 min) Optimal temperature (pH 5.0) $K_{\rm m}^{\ \ a}$ $V_{\rm max}^{\ \ a}$	160 kDa 4–8 5.0 ~40°C 60°C 0.22 mM (<i>p</i> -NβG) 0.99 mM (cellobiose) 2.89 mM/min (<i>p</i> -NβG) 3.59 mM/min (cellobiose)
Final resolution for substrate	Glucose Cellobiose

p-NβG, p-nitrophenyl-β-D-glucopyranoside

Table 3. Effect of metal ions and chemical agents on β -glucosidase activity from T. matsutake

Chemical reagent	Relative activity (%)		
Standard	100		
CaCl ₂ · 2H ₂ O	287		
$MnCl_2 \cdot 4H_2O$	257		
BaCl ₂ · 2H ₂ O	186		
$MgCl_2 \cdot 6H_2O$	180		
ZnCl ₂	177		
PbCl ₂	163		
NaCl	163		
$CuSO_4 \cdot 5H_2O$	160		
LiCl	154		
CoCl ₂	151		
$FeCl_2 \cdot 4H_2O$	149		
$AlCl_3 \cdot 6H_2O$	143		
$HgCl_2$	129		
KCl	94		
$AgNO_3$	71		
Ethylenediaminetetraacetic acid (EDTA)	214		

Enzyme activity was assayed in the presence of metal ions or chemical agents in the final concentration of 1 mM under standard conditions Enzyme activity as a standard was measured in the absence of $CaCl_2 \cdot 2H_2O$ β -Glucosidase activity was assayed with various metal ions and EDTA by the *p*-nitrophenol method

Table 4. Substrate specificity of purified extracellular β -glucosidase from *T. matsutake* Z-1

Substrate	Relative activity (%)
Synthetic substrate ^a	
<i>p</i> -Nitrophenyl-β-D-glucopyranoside	100
<i>p</i> -Nitrophenyl-α-D-glucopyranoside	0
<i>p</i> -Nitrophenyl-β-D-cellobioside	0
Natural substrate ^b	
Cellobiose $(\beta 1 \rightarrow 4)$	100
Cellobiose $(\beta 1 \rightarrow 4)$	63
Maltose $(\alpha 1 \rightarrow 4)$	0
Isomaltose $(\alpha 1 \rightarrow 6)$	0
Lactose $(\beta 1 \rightarrow 4)$	0
Treharose $(\alpha 1 \rightarrow 1)$	0
Sucrose $(\alpha \hat{1} \rightarrow 2)$	0
Isomaltooligosaccharride ($\alpha 1 \rightarrow 6$)	0
Dextrin $(\alpha 1 \rightarrow 4, \alpha 1 \rightarrow 6)$	0
CM-cellulose $(\beta 1 \rightarrow 4)$	0
Avicell $(\beta 1 \rightarrow 4)$	0
Curdlun $(\beta 1 \rightarrow 3)$	0

^aReaction mixture (200 μl) containing 3 μl purified enzyme solution, $100 \mu l$ each synthetic substrate (20 mM), and $20 \, \text{mM}$ CaCl₂ containing 0.1 M Mcllvaine buffer (97 μl) was incubated at pH 5.0 and 37°C for 30 min; reactant product (yield) was calculated from the decomposition of each substrate, and enzyme activity was expressed in values relative to p-nitrophenyl-β-D-glucopyranoside, 100, as standard; enzyme activity was measured by the method of p-nitrophenol

^bReaction mixture (200 μl) containing 5 μl enzyme, 100 μl each natural substrate (oligosaccharide, 20 mM; polysaccharide, 0.2%), and 20 mM CaCl₂ containing 0.1 M Mcllvaine buffer (95 μl) was incubated at pH 5.0 and 37°C for 120 min, and enzyme activity was expressed in values relative to cellobiose, 100, as standard; enzyme activity was measured by the method of F-kit D-glucose

Substrate specificities toward synthetic substrates and natural substrates

The substrate specificity of β -glucosidase was tested using different glycosidic bonds of saccharides for synthetic substrates (Table 4). The glucosidase readily activated p-N β G [relative activity (R.a.), 100%]. However, the glucosidase did not hydrolyze other synthetic substrates. Also, the substrate specificity of β -glucosidase was tested using carbohydrates with different glycosidic bonds of saccharides for natural substrates (Table 4). The β -glucosidase activity was measured by the F-kit D-glucose method. The glucosidase readily hydrolyzed cellobiose (R.a., 100%) and cellotriose (R.a., 62%) but did not hydrolyze other saccharides. As a result, it was indicated that β -glucosidase hydrolyzed only oligosaccharides having the β -1,4-glycosidic bond.

Also, the natural substrate cellotriose was hydrolyzed by β -glucosidase. The resultant products were analyzed by TLC (Fig. 3). It was observed that the cellotriose was decomposed to cellobiose and glucose. In the long-term reaction (24h), the substrates were decomposed completely.

Discussion

When the fungi forms fruit bodies, large amounts of mycelia may be needed. However, it is very difficult to cultivate large amounts of mycelia using monosaccharides in a pure

 $^{^{\}rm a}$ β -Glucosidase activity was assayed with various concentrations of p-N β G and cellobiose by the p-nitrophenol method and F-kit D-glucose method, respectively

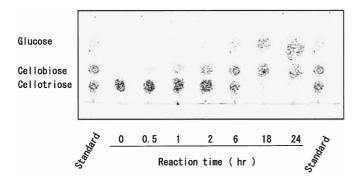


Fig. 3. Resultant products of cellotriose by β-glucosidase from *T. matsutake* Z-1. Reaction mixture was composed of $5\,\mu$ l enzyme solution, $100\,\mu$ l cellotriose (20 mM), $95\,\mu$ l 0.1 M McIlvaine buffer, pH 5.0, in a final volume of $200\,\mu$ l. The degradation pattern of the cellotriose was analyzed by thin-layer chromatography (TLC). Reactions were done at 37°C for 0–24h. Developing solution, acetic acid:chloroform:distilled water, 7:5:1; developing time, about 4h; chromoagent, 20% sulfalic acid-methanol

culture because of the osmotic pressure in the medium (Ohta 1994). Kawai (1973) indicated that *T. matsutake* had poor amylase activity among the many mushroom fungi. Ohta (1997) reported that several strains of *T. matsutake* had the ability to utilize starch as a carbon source.

For the artificial cultivation of this fungus, the ability of starch utilization is very important because the fungus had previously not been known to use other polysaccharides except starch. From these facts, it is important to investigate the ability of amylase production in this fungus. Lee et al. (1998) reported that *T. matsutake* had relatively high amylase activity among its other extracellular enzymatic activities. Terashita et al. (2000b) have reported that the mycelial growth was remarkably promoted, and amylase production in the culture medium obviously increased, when starch from yam and potato was added to the medium as a growth substrate.

In our previous report, we studied the production of amylases and purified extracellular $\alpha\text{-amylase}$ from a static culture filtrate. Enzymatic properties of the highly purified $\alpha\text{-amylase}$ had already been reported from the aspect of starch utilization as a growth substrate (Kusuda et al. 2003). This $\alpha\text{-amylase}$ readily hydrolyzed the $\alpha\text{-1,4-glucosidic}$ bond in soluble starch and amylose but did not hydrolyze the $\alpha\text{-1,4-glucosidic}$ bond, and cyclic polysaccharides, the $\alpha\text{-1,6-glucosidic}$ bond, and cyclic polysaccharides such as $\alpha\text{-}$ and $\beta\text{-cyclodextrin}$.

In this article, we have reported that T. matsutake produce a β -glucosidase having high activity from static culture filtrate in the research of carbohydrase production to elucidate the utilization of carbohydrates as the growth substrate in ectomycorrhizal mushrooms. This finding suggests the saprotrophic abilities of this fungus.

Then, we tried purification procedures and characterization of β -glucosidase from *T. matsutake*. Until quite recently, it was reported that *T. matsutake* does not use substrates having a β -glucosidic bond such as cellooligosaccharides and cellulose (Ogawa 1978). Conversely,

Nakazawa et al. (1974) reported that *T. matsutake* cellulolytic enzymes produced a very weak activity. Enokibara et al. (1993) reported the production of cellulolytic enzymes in mycorrhizal mushrooms and showed that *Hebeloma vinosophyllum* Hongo and *Laccaria bicolor* Maire P.D. Orton produced very weak activities of the subneutral cellulases active at pH 5.5–6.8. Also, *Lepista nuda* (Bull.: Fr.) Cooke produced strong activities of the acid cellulase active at pH 3.0–5.5. Furthermore, Terashita et al. (1995) also showed the production of exoglucosidases of mycorrhizal mushrooms such as *T. matsutake* and *Lyophyllum shimeji* (Kawam.) Hongo on potato dextrose liquid medium. As a result, they detected that *T. matsutake* produced weak activities of CM-cellulase and avicelase in all test strains.

Guerin-Laguette et al. (2001) recently reported the saprotrophic growth of T. matsutake in soil. They showed mycelial growth stimulation and increase in β -glucosidase excretion from the hyphae of T. matsutake using surfactants such as Tween 80 and Tween 40. The addition of the surfactants seemed to greatly improve the saprotrophic potential of this fungus on pine bark. Moreover, Vaario et al. (2003) reported T. matsutake produced β -glucosidase when pine bark was used as the substrate for cultivation of this fungus. They indicated that hyphae of T. matsutake were able to degrade and invade the xylem cell walls of sawdust (from a pine tree) by transmission electron microscope observations. However, no characterization of the highly purified β -glucosidase from a mycorrhizal mushroom has been reported up to the present.

In this report, we showed good purification procedures and the enzymatic properties of highly purified β -glucosidase from T. matsutake for the first time. The molecular mass of the β-glucosidase (160kDa) was large compared to enzymes from other glucosidases of mushroom origin, such as β-glucosidase (110kDa) from Termitomyces clypeatus Heim (Sengupta et al. 1991), CMCase IIIa (24.3 kDa) from Polyporus arcularius (Fr.) Ames, 69B (ATCC 24461) (Ishihara et al. 2005), and α -glucosidase (51 kDa) from Lentinula edodes (Berk.) Sing. (Yamazaki and Suzuki 1978). The β -glucosidase from *T. matsutake* has a wide range of pH stability (pH 4.0-8.0) as compared with βglucosidases from other fungi. This enzyme activity showed optimum temperature at 60°C. We studied the properties of extracellular α -amylase from *T. matsutake* (Kusuda et al. 2003), and the optimum temperature was about 60°C.

Among the metal ions tested for β -glucosidase activity, the enzyme from *T. matsutake* was activated remarkably by the presence of Ca²⁺ (2.87 times that of the control), Mn²⁺ ion (2.57 times), and EDTA (2.14 times). Ag⁺ weakly inhibited glucosidase activity.

In the present study, a β -glucosidase from T. matsutake hydrolyzed disaccharides (cellobiose and p-N β G) and cellotriose having the β -1,4-glucosidic bond. However, the enzyme did not act upon disaccharides and polysaccharides having α -1,4-, α -1,1-, and α -1,6 bonds. Moreover, we examined the activity of the enzyme toward cellotriose by TLC analysis of hydrolyzates from the enzyme. From this result, it was decided that the enzyme β -glucosidase acted on oligosaccharides such as cellobiose. This β -glucosidase seem to

be similar to CMCase IIIa from a saprophytic mushroom, *Polyporus arcularius*, which had been reported by Ishihara et al. (2005) as to substrate specificity. The β -glucosidase from *T. matsutake* can be considered as acting on oligosaccharides in the range di–hepta. It is thought that the high concentration of the substrate or feedback inhibition is a cause for the results in our experiments. More detailed information is needed to reveal the substrate specificities of the β -glucosidase from *T. matsutake*.

To reveal the ability to utilize starch as a growth substrate and investigate the saprotrophic aspects of this fungus, we purified and characterized the α-amylase (Kusuda et al. 2003). Moreover, we identified a β-glucosidase from this mushroom. This enzyme hydrolyzed the β -1,4glucosidic bond in oligosaccharides having a β-1,4 bond. Our finding suggests that T. matsutake is able to utilize oligosaccharides released from cellulose and its related compounds in nature such as "shiro soil." However, more detailed information is needed about amylase and β-glucosidase production from T. matsutake and other microbes, including litter inhabited by microbes, plants, and animals in "shiro soil." Ectomycorrhizal fungi are believed to obtain much of the carbon necessary for growth from the host plants through ectomycorrhizae. Recently, Hatakeyama and Ohmasa (2004) have showed that many strains of the genus Suillus and Boletinus can grow well at relatively high glucose concentrations (33.3-100 g/l). Conversely, Vaario et al. (2002) discussed that carbon transfer may be low via the mycorrhizae compared to their experimental results. From these results, quantification of carbon source flow from the host plant to the mycobiont *T. matsutake* is also important.

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