

Cadmium and heat response of the fungus *Heliscus lugdunensis* isolated from highly polluted and unpolluted areas

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Summary. Induction of heat shock protein (Hsp) 70 and distinct metallothionein-like proteins (MTLPs) in response to Cd and heat treatment were studied in two strains of the aquatic hyphomycete *Heliscus lugdunensis*: HI-H4, isolated from a heavy metal polluted site, and HI-BB taken from an unpolluted area. Upon Cd-exposure, Hsp70 was actively synthesized in the strain HI-H4, and to a much lower degree in the strain HI-BB. The Hsp70-expression was time- and dose-dependent, reaching a maximum after 24 h incubation with 80 μ M Cd. Upon heat-stress, a similar response was observed: a strong Hsp70-expression in HI-H4, and only a marginal one in HI-BB. The strains reacted to Cd-exposure by a specific, environmentally related induction of MTLPs, as shown by the highly sensitive biamine derivatisation method of SH-rich proteins. In HI-H4, a strong expression of 11 kDa MTLP was registered, which followed strictly the induction pattern of Hsp70. This suggests interdependence of the induction mechanisms and roles of these stress proteins in metal resistance. On the contrary, in HI-BB a weak expression of MTLP of about 20 kDa was observed, exhibiting completely different induction pattern. The results suggest that the specific induction of Hsp70 and/or distinct MTLPs in the range of 11 kDa in *H. lugdunensis* strain HI-H4 are essential adaptive mechanisms to continuous heavy metal exposure.

Keywords: Cadmium – Heat stress – Metallothioneins – Aquatic hyphomycetes – *Heliscus lugdunensis* strains

Abbreviations: AA, acrylamide; AP, alkaline phosphatase; APS, ammonium peroxydisulphate; Bis-N,N'-methylene-bis-acrylamide; BCIP, 5-bromo-4-chloro-3-indolyl phosphate disodium salt; DTE, dithiothreitol; DTPA, bis-(2-aminoethyl)-amino-N,N',N'',N'''-pentaacetic acid; ECL, electrogenerated chemoluminescence; EDTA, ethylenediamine tetraacetic acid; NC, nitrocellulose; PMSF, phenylmethylsulphonylfluoride; PAGE, polyacrylamide gel electrophoresis; SDS, sodium dodecyl sulphate; PBS, phosphate buffered saline; Hsp, heat shock protein; MT, metallothionein; MTLP, metallothionein-like protein; NBT, p-nitro blue tetrazolium chloride; TCA, trichloroacetic acid; TCEP, tris-(2-carboxyethyl) phosphine chloride; Tris, tris(hydroxymethyl)aminomethane

Introduction

In recent years, there has been a renewed interest in the role of fungi as potential bioremediators of heavy metal

polluted environments (Gadd, 2001; Fomina et al., 2005). Fungi are capable of removing metals either by biosorption or bioaccumulation in the cytoplasm, binding them, e.g., by cysteine-rich peptides such as glutathione (GSH), metallothioneins (MTs) or phytochelatins (PCs) (Cobett and Goldsbrough, 2002; Jarosz-Wilkolazka et al., 2005). Special attention in this respect has been given to some aquatic fungi found in highly contaminated waters. The aquatic hyphomycete *Heliscus lugdunensis* was one of the most abundant species inhabiting heavy metal polluted sites (Krauss et al., 2005a, b; Sridhar et al., 2005). Heavy metal tolerance in filamentous fungi and their mechanisms of adaptation to polluted aquatic ecosystems are not well understood. In the filamentous fungus *Curvularia* sp. tolerance against Cd is acquired by biosorption on the mycelial surface (Vepachedu et al., 1997).

H. lugdunensis strains have been shown to respond to increased external Cd-concentrations by increasing the internal GSH-content, which is accompanied by rising levels of cysteine and γ -glutamylcysteine (Braha et al., 2007), as well as by the synthesis of some MTs and PCs (Jaeckel et al., 2005a, b). Yet, the detailed role of distinct MTs in the metal detoxification in fungi remains to be elucidated. Little is known about the participation of other stress proteins (e.g., heat shock proteins, Hsps) in these defense mechanisms. Hsps are ubiquitous tools of all living organisms to cope with stress-induced denaturation of other proteins (Wang et al., 2004).

In this work, we compared the Cd-stress and heat response of *Heliscus lugdunensis* strain H4-2-4, isolated from a highly polluted area with that of HI-BB isolated from an unpolluted site. We focused our attention on

involvement of Hsp70 and distinct MTs in fungal defence and adaptation against heavy metal toxicity.

Materials and methods

Chemicals and Hsp70 antibody

Acrylamide; N,N'-methylene-bis-acrylamide-Solution (Rotiphorese gel 30, Roth, Karlsruhe); sodium dodecyl sulphate (Roth); Tricine (Roth); Tris (Roth); Temed- (N,N,N',N'-tetramethylethylenediamine, Serva, Heidelberg), ammonium peroxydisulphate (Merck, Darmstadt); NBT, p-nitro blue tetrazolium chloride (USB Corp., Cleveland Ohio); BCIP, 5-bromo-4-chloro-e-indolyl phosphate (USB Corp., Cleveland Ohio); DTPA, bis-(2-aminoethyl)-amino-N,N,N',N',N''-pentaacetic acid (Sigma-Aldrich, München); monobromobimane (Fluka, Darmstadt); polyclonal antibody against cytosolic Hsp70 of tomato (*Lycopersicon peruvianum*) was a gift of Neumann et al. (1987); rabbit anti-Hsp70 alkaline phosphatase conjugate (Biomol, Hamburg); Western-Light chemoluminescence-test (CDP-Star, Tropix, Invitrogen, Carlsbad CA, Cat-No. SO-W008).

Fungal strains and cultivation

Strains of *Heliscus lugdunensis* Sacc. et Therry were isolated from single conidia from following sites: strain HI-BB, from a non-polluted stream (Boss Brook near Amherst, Nova Scotia, by Prof. F. Bärlocher, Mount-Allison-University Sackville, Canada (Bärlocher, 1987)); strain H4-2-4, from a polluted location (Hergisdorf near Eisleben, Saxony-Anhalt, Germany, by Krauss et al. (1998, 2001)). The strains are available from DSMZ Braunschweig (HL-BB: DSM 18483) or Dr. G. Krauss (H4-2-4, Department Groundwater Microbiology, UFZ, Centre for Environmental Research, Leipzig-Halle, Germany), abbreviated to HI-H4.

All cultures were maintained on malt extract agar (0.5% malt extract, 0.1% peptone, 1.5% agar). To examine the effects of heavy metals on biomass production, the fungi were grown in broth cultures (0.5% malt extract, 0.1% peptone; 75 ml medium in 200 ml Erlenmeyer flasks). The cultures were inoculated with homogenized agar slices overgrown with a 14 days old culture; the total fungal biomass introduced per flask was 1–1.3 mg. The flasks were incubated on an orbital shaker ($14 \pm 0.7^\circ\text{C}$; 120 rpm). Cd^{2+} was added as chloride salts (Sigma) in the first variant, at beginning of cultivation, and in the second variant, at 4th day of cultivation. Standard solutions were filter sterilized (0.2 μm membrane filters) before adding to achieve final metal concentrations of 40, 80 and 120 μM in the cultures. The distinct procedure of time and temperature arrangements in induction experiments of Hsp70 and thiol proteins (metallothioneins or metallothionein-like proteins) are given in the corresponding legends of figures.

Determination of dry mass

To determine growth, mycelia incubated for 12 d in three parallels without or with Cd at a final concentration of 80 μM were used (log phase) and then harvested by filtration (Merck, filter 0869) after 2, 4, 6, 8, 10 and 12 days. Mycelia were rinsed twice with distilled water, blotted dry between filter papers, weighed (fresh weight) and used for analysis. To determine wet to dry weight ratio, pre-weighed fresh mycelia were exposed to 80°C for at least 2 h, and weighed again. The standard deviation of the conversion factor was 15–18%. The influence of Cd on growth of the *Heliscus* strains HI-BB and HI-H4 is shown in Fig. 2 by average values and standard deviations.

Protein determination

Hundred microliters of the supernatant were mixed with 1 ml of 5% TCA and left overnight at 4°C to precipitate the proteins. The precipitate was

washed 1-time with 1% TCA and 3-times with water, and the amount of protein determined according to Lowry et al. (1951).

Isolation of cytosolic protein to identify Hsp70

The mycelia were harvested by filtration and the wet mass was homogenized under liquid N_2 by stirring using mortar and pestle. The material was extracted with 50 mM Tris-HCl (pH 8) containing 3 mM EDTA, 10 mM DTE and 0.2% PMSF. The homogenate was centrifuged at 8500 g for 10 min and aliquots were used for protein determination and sample preparation.

Sample preparation to identify Hsp70

The proteins were precipitated by a pre-chilled acetone at -20°C overnight, centrifuged at 5000 g for 30 min and dissolved as follows: For SDS-PAGE, 100 μg of protein were dissolved in sample-buffer (50 mM Tris-HCl, pH 6.8, 10% (w/v) glycerol, 2% (w/v) SDS, 5% (v/v) β -mercaptoethanol) to obtain a final concentration of 2 $\mu\text{g}/\mu\text{l}$. Before electrophoresis, to the samples 2 μl of 50% bromophenol blue were added.

Sample preparation to identify sulfur-rich proteins (metallothioneins)

The mycelia were harvested and homogenized as above. The frozen material (15–30 mg fresh mass) was filled in pre-chilled (-20°C) septum-vials, 100 μl of 500 μM TCEP were added and the proteins were reduced for 1 h at room temperature in darkness. Then, to each sample 0.4 ml of 5.8 mM monobromobimane in 50 mM Hepes-NAOH-buffer at pH 8, containing 5 mM DTPA and 50% acetonitrile, was added. The derivatization and sample preparation for SDS-PAGE followed the procedure of Fan et al. (2004).

For SDS-PAGE, 100 μg of protein were dissolved in sample-buffer (50 mM Tris-HCl, pH 6.8, 10% (w/v) glycerol, 4% (w/v) SDS) to obtain a final concentration of 2 $\mu\text{g}/\mu\text{l}$. Before electrophoresis, to the samples 2 μl of 50% bromophenol blue were added.

SDS-PAGE

For Hsp70 analysis, one-dimensional SDS-polyacrylamide gel electrophoresis was carried out in 5% (w/v) stacking gel and 12.5% (w/v) separating gel, or in a 12–17.5% gradient gel, according to Laemmli (1970). The proteins were stained with Coomassie Brilliant Blue R-250 (Serva, Heidelberg) according to Weber and Osborne (1969) or silver, according to Blum et al. (1987). To analyze sulfur-rich-proteins (MT), the proteins were separated by one-dimensional PAGE on a modified Tricine-urea gel, according to Schägger and Jagow (1987): The stacking gel contained 4% AA; 0.44% Bis; 0.31 M Tris-HCl, pH 8.5; 0.4% SDS, the separation gel 12–20% AA, 1.9% Bis, 4.6 M urea, 0.76 M Tricine-Tris, pH 8.5; 0.1% SDS. To 7.5 ml separation gel mixture 2 μl Temed and 20 μl 10% APS were added and the mixture left to polymerize overnight at 8°C . For relative molecular weight calibration, recombinant proteins prestained (Fermentas SMO671, 11, 17, 24, 30, 40, 55, 70, 100, 130, 170 kDa or SM1811, 10, 15, 27, 35, 55, 70, 100, 130, 250 kDa) were used.

The gels were run in a Minigel-tool (Biometa, Göttingen) for about 1 h at 80 V, 15–18 mA and about 1.5–2 h at 100 V, 28–50 mA. After finishing of separation, the gels were stained using silver-reagent or coomassie as indicated above, and prepared for Hsp70 analysis by Western-blotting. In the case of monobromobimane-conjugated proteins, the gels were analyzed by UV-light (UV-Gel-Yet-Imager, INTAS) and quantified by 1-D-Image-System (Biotec-Fischer).

Western blot

Proteins were transblotted from gels onto nitrocellulose membranes (0.2 or 0.45 μm , Schleicher and Schüll, Germany) for 2 h at 300 mA, and 1 h at

350 mA using 0.02 M NaH_2PO_4 , pH 6.4 in a tank-blot tool. The immunoblotting procedure was carried out as described by Neumann et al. (1987) using a polyclonal anti-rabbit Hsp70-antibody. After blotting, the proteins on the membrane were stained by Ponceau S, washed thrice with water, once with PBS-solution and blocked for 1 h at 37 °C with 1% non-fat-milk (Bio-Rad, Germany) in PBS-solution, pH 7.5 and 0.1% Tween 20. Thereafter, the membrane was washed for 1 h with PBS and probed for 1 h with a polyclonal antibody against cytosolic Hsp70 (*Lycopersicon peruvianum*; 1:1000) as indicated. After washing again with PBS for 1 h, the membrane was incubated for 1 h with AP-conjugated secondary goat anti-rabbit antibody (1:2000), and after washing for 1 h with PBS either analyzed by the colour-test (substrates of alkaline phosphates, NBT and BCIP), or a highly sensitive luminescence technique (ECL, CDP-star, Tropix) was applied to detect immunoreactive spots on membranes.

Statistical analysis

Determination of standard deviation (SD) for dry weight (growth of *Heliscus*) and protein spots (Hsp70, metallothioneins) was performed with MS Excel 6.0. Significant differences of these data were calculated using Student's *t*-test. *P* values <0.05 were considered significant (Timischl, 2000).

Results

Comparison of aquatic hyphomycete *Heliscus lugdunensis* strains HI-BB and HI-H4 showed significant differences in their growth and the induction of Hsp70 and metallothionein-like proteins by cadmium.

At first, the strains were studied in liquid culture for their response to Cd or heat stress, using Hsp70-induction as a bioindicator. Seven days old cultures were incubated

for 4 or 24 h at 14 °C in the presence of 40, 80 or 120 μM CdCl_2 . Thereafter, cytosolic proteins were isolated and analyzed by SDS-PAGE and immunoblotting, using a polyclonal Hsp70-antibody. The results are presented in Fig. 1. In HI-BB, a very low level of Hsp70 under the conditions tested was observed. In HI-H4, however, concentrations in the range of 40–120 μM Cd produced a clear expression of Hsp70. The highest Hsp70 expression was registered with 80 μM Cd after 24 h incubation, yet a 4 h treatment with 120 μM Cd also resulted in an elevated Hsp70 synthesis. A prolonged treatment with 120 μM Cd caused no Hsp70-induction, probably because of toxic effects.

This remarkable difference in Cd-response of both strains was illustrated by a growth-inhibitory test in the presence of 80 μM Cd. Upon 12 days incubation, the growth of strain HI-BB (as evaluated by the dry mass of mycelia) decreased to 40% compared to control culture (without Cd), while the growth of strain HI-H4 reached 86% under the same conditions (Fig. 2).

Both strains responded to heat stress in a similar way, as with Cd. 7 d-Cultures of HI-BB and HI-H4 were incubated for 2–4 h at 40° and the cytosolic proteins were analyzed as above. In HI-BB, only a marginal presence of Hsp70 was registered. In contrast, in HI-H4 a pronounced expression of the stress protein was evident, especially after 2 h of heat stress (Fig. 3). Upon prolonged heating (4 h), the amount of Hsp70 decreased, probably because of damaging effects on mycelia.

When subjecting the strains to a double stress, first to Cd, and then to heating, the response remained the same: a marginal induction in HI-BB, and a definite expression

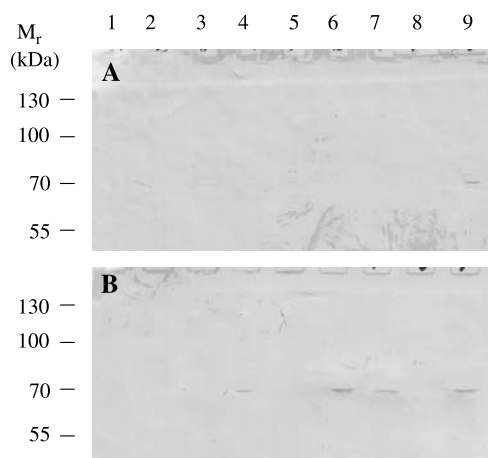


Fig. 1. Induction of Hsp70 in *H. lugdunensis* strains HI-BB and HI-H4 by cadmium. The mycelia of HI-BB (A) and HI-H4 (B) were cultivated 7 d at 14 °C (control, 2), then incubated for 4 h (3) or 24 h (4) at 40 μM Cd, for 4 h (5) or 24 h (6) at 80 μM Cd, and for 4 h (7) or 24 h (8) at 120 μM Cd. 9 Illustrates the serological result with 2 μg of *Cucurbita* Hsp70. Buffer soluble proteins were extracted, 20 μg were subjected to each lane, separated by 10% Laemmli-SDS-PAGE, transferred to NC-membrane and incubated with Hsp70-antibody. 1 Fermentas marker proteins, 11–170 kDa. Electrophoregrams after analysis by phosphatase-colour-test are depicted

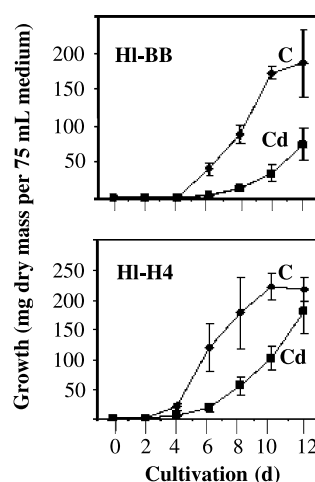


Fig. 2. Growth of *H. lugdunensis* strains HI-BB and HI-H4 without (C, control) or in the presence of cadmium (Cd, 80 μM) in the time range of 12 days. Cd was added at beginning of the experiment

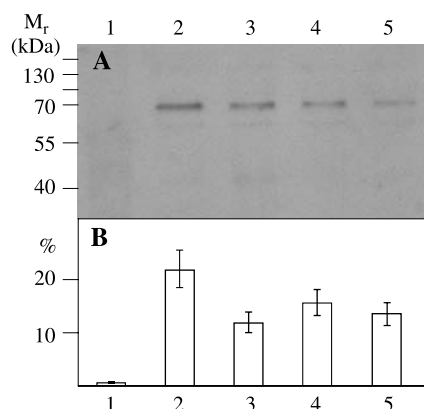


Fig. 3. Induction of Hsp70 in *H. lugdunensis* strains HI-BB and HI-H4 upon heat stress. **A** Electrophoresis after analysis by ECL-test. **B** Semi-quantitative calculation of spot intensity. The mycelia of two strains were cultivated for 7 d at 14 °C without (control) or with 80 μ M Cd, added at 4th day. Then, control- and Cd-treated mycelia were incubated for 2 or 4 h at 40 °C. Buffer soluble proteins were extracted, 20 μ g were subjected to each lane and separated by 10–17% Laemmli-SDS-PAGE: Strain HI-BB upon 2 or 4 h at 40 °C (1). Note that similar results were obtained upon heating of Cd-treated HI-BB; strain HI-H4 upon 2 or 4 h at 40 °C (2 and 4); Cd-treated HI-H4 upon 2 or 4 h at 40 °C (3 and 5). Control strains exhibited similar Hsp70-expression, as that in 1

of Hsp70 in HI-H4 (Fig. 3). It should be mentioned that the Hsp70 levels in this case were lower compared to those upon heat stress only, due to the toxic effect of the combined treatment.

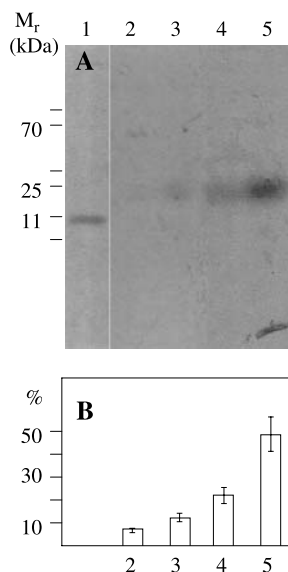


Fig. 4. Induction of sulphur-rich proteins in *H. lugdunensis* strain HI-BB by cadmium. **A** Electrophoresis after analysis by UV-light. **B** Semi-quantitative calculation of spot intensity. The strain was cultivated for 7 d at 14 °C by shaking and then incubated for 24 h without or with cadmium. Proteins were extracted and each of 20 μ g protein was separated on 12–20% Schagger-gradient gel: Monobromobimane-derivative of 5 μ g MT-1 (rabbit, see Materials and methods) (1); control of shaking culture (2); 40, 80 and 120 μ M Cd (3–5)

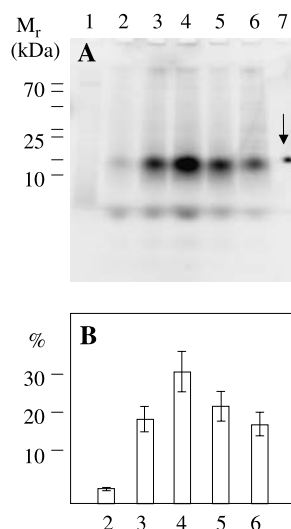


Fig. 5. Induction of sulphur-rich proteins in *H. lugdunensis* strain HI-H4 by cadmium. **A** Electrophoresis after analysis by UV-light. **B** Semi-quantitative calculation of spot intensity. The material was cultivated for 7 d at 14 °C by shaking and then incubated for 24 h without or with cadmium. Proteins were extracted and each of 20 μ g protein was separated on 12–20% Schagger-gradient gel: Fermenta-marker SM1811, 9 recombinant proteins 10–250 kDa, 5 μ l (1); control of shaking culture (2); 40, 80 and 120 μ M Cd (3–5); control of a fixed culture (6); monobromobimane-derivative of 2 μ g MT-1 (rabbit) at 11 kDa (arrow) (7)

Secondly, protein fractions were isolated, thiol groups of proteins reacted with monobromobimane and then, the labelled proteins were separated by SDS-PAGE. A high fluorescence of bimane-derivatives of thiol-rich proteins was only detected (Figs. 4, 5). *Heliscus* cells of these two strains responded to cadmium over 24 hours treatment by a different concentration-dependent induction of sulfur-rich metallothioneins or metallothionein-like proteins. Thus, with HI-BB a maximum MTLP-induction was obtained at 120 μ M Cd reaching about 600% of that of controls (being 100%) (Fig. 4A, B, lane 5 compared to lane 2, control), whereas with HI-H4 a maximal protein induction was produced at 80 μ M Cd reaching about 800% of control levels (Fig. 5A, B, lane 4 compared to lane 2, control). Surprisingly, the relative molecular masses of MTLPs analyzed by SDS-PAGE were 20 and 11 kDa in the strains HI-BB and HI-H4, respectively (Figs. 4A, 5A). Further unknown fluorescent spots at about 5 kDa, and very small ones at about 2 kDa were detected (Fig. 5). No significant increase of MTLPs was observed by heating of 7 days old mycelia for 4, 8 or 24 h at 40 °C (not shown).

Discussion

The mechanisms of heavy metal adaptation and tolerance in filamentous fungi are not well understood. Current

Table 1. Induction of Hsp70 in *H. lugdunensis* strains from polluted and unpolluted areas

Induction agent	Concentration (μM)	Time of treatment (h)	HI-BB	HI-H4
Heat (40°C)		2	±	++
Cd	40	1	–	–
		2	–	–
		4	–	–
		24	–	±
	80	1	–	–
		4	–	–
		24	toxic	++
	120	1	–	+
		4	toxic	+
		24	toxic	toxic
	IC ₅₀ ^a	120	45 ^b	100 ^c

++ Strong positive

+ Positive

± Low induction

– Negative

^a Inhibitory concentration reducing the growth of fungi by 50%^b Miersch et al. (1997)^c Jaeckel et al. (2005a)

evidence suggests that stress proteins may play an important role in this process, yet their detailed functions remain to be elucidated. Special attention in this respect is paid to metallothionein-like proteins (MTLPs) and heat-shock proteins (Hsp). In this work, we were concerned with the induction of Hsp70 and some MTLPs in *H. lugdunensis* strains HI-H4 and HI-BB, growing on differently polluted natural habitats, in order to correlate their expression with heavy metal resistance of the strains. In vitro studies have revealed a significantly higher resistance of HI-H4 towards Cd (IC₅₀ of 100 μM) (Jaeckel et al., 2005a) as compared to HI-BB (IC₅₀ of 45 μM) (Miersch et al., 1997). An elevated resistance of HI-H4 to Cd is also shown in this study (Fig. 2). We demonstrate that upon Cd-stress, Hsp70 are actively synthesized in the strain HI-H4 isolated from a highly polluted site, and to a much lower degree (trace) in HI-BB growing on a non-contaminated area. The Hsp70-expression, as summarized in Table 1, was time and dose-dependent, reaching a maximum after 24 h at 80 μM Cd. This acquired Hsp70 response was further evidenced by subjecting both strains to heat (40°C). Again, there was a strong Hsp70-induction in HI-H4, and only a marginal one in HI-BB. This remarkable feature of HI-H4 merits special attention. Transcription of Hsp70 gene, known to be activated by heat stress, is also induced by a variety of metals such as Zn, Cd, Cu and Hg (Williams and Morimoto, 1990; Murata et al., 1999). It has been shown that the induction of Hsp70

by heavy metals, as well as by heat shock, is mediated by metal-activated HSF1 (heat shock factor 1) (Mosser et al., 1988; Williams and Morimoto, 1990; Murata et al., 1999). We assume that under continuous heavy metal exposure, HI-H4 has acquired a higher steady-state activation ability of HSF1 compared to HI-BB, thus ensuring a better survival. To our knowledge, this is the first report to show the involvement of Hsp70 in the adaptive response of aquatic hyphomycetes to heavy metal toxicity. The majority of studies on Hsp70 in fungi have been performed on ascomycetes such as *Saccharomyces cerevisiae* (Boorstein et al., 1994), and some other fungal phyla including zygomycete *Rhizopus nigricans*, basidiomycetes *Boletus edulis* and *Cryptococcus curvatus*, and chytridiomycete *Blastocladiella emersonii* (Cernila et al., 2003). The authors report elevated Hsp70-mRNA content in *R. nigricans* upon heat or Cu-stress (Cernila et al., 2000), or elevated Hsp70 concentrations in *B. edulis* after Cd-exposure (Collin-Hansen et al., 2005), the effects being in all cases considered as immediate stress and/or defence response (but no adaptation) against metal toxicity.

HI-H4 and HI-BB exhibit specific, environmentally related induction of sulphur-rich proteins, MTs or MTLPs, upon Cd-treatment, differing in molecular mass and induction pattern, as unambiguously shown by the highly sensitive biamine derivatisation method of SH-containing proteins (Fan et al., 2004). HI-H4 responded by a strong expression of 11 kDa SH-rich protein (most likely metallothionein). The cadmium-inducibility of 11 kDa MT in HI-H4 only suggests its involvement in metal tolerance and detoxification. Remarkable is the similarity of the induction pattern of this metallothionein to that of Hsp70 (both stress proteins reached maximum expression at 80 μM Cd) (compare Figs. 1 and 5), which suggests interdependence of their induction mechanisms and roles in metal resistance. Recently, a cross-talk of heat shock and heavy metal stress pathways in human cell has been suggested (Uenishi et al., 2006). On the contrary, HI-BB reacted to Cd-stress by a weak induction of SH-rich protein of about 20 kDa (not expressed in HI-H4). The induction pattern of this MT (Fig. 4) differed completely from that of 11 kDa MT or Hsp70, pointing rather to an immediate stress bioindication than detoxification in HI-BB.

MTs are ubiquitous, functionally diverse metal-binding proteins of relatively low molecular mass that play major roles in metal homeostasis and/or detoxification (Kägi and Schaffer, 1988; Cobett and Goldsbrough, 2002; Roosens et al., 2005). Fungi contain mainly small MTs (3–6 kDa). In aquatic fungi, only a few of them have been identified so far to be involved in metal detoxification.

The content of thiol-rich proteins was reported to increase significantly in the aquatic hyphomycetes *Fontanospora fusiramosa* and *Flagellospora curta*, exposed to Cd (Guimaraes-Soares et al., 2007). Recently, 2.3 kDa MT has been identified and characterized in *H. lugdunensis* strain H8-2-1, isolated from a moderately polluted site (Jaekel et al., 2005b). In other phyla, MT of 53 amino acid residues has been found in certain mutants of *S. cerevisiae* to produce copper or cadmium resistance (Tohoyama et al., 1995). Small copper binding MTs were described for the ectomycorrhizal fungi *Laccaria laccata* and *Paxillus involutus* (Howe et al., 1997). In *P. involutus*, Cd has been found to induce 3 kDa MT-related proteins, thought to be involved in metal tolerance (Courbot et al., 2004). Class II MTs were induced by Cu- and Cd-exposure in the heavy metal resistant ascomycete *Beauveria bassiana* (Kameo et al., 2000). Our article is the first to describe MT of about 11 kDa involved in adaptation of aquatic hyphomycetes to continuous heavy metal exposure, thus expanding the knowledge on the function of diverse MTs. Interestingly, 11 kDa Cd-binding metallothionein-like protein from the unicellular green algae *Chlorella sorokiana* has been reported to confer an extremely high resistance of this species to heavy metals (Yoshida et al., 2006). Other sulphur-rich proteins between 70 and 140 amino acids, such as hydrophobins (Temple and Horgen, 2000) were not found in aquatic fungi.

In conclusion, the specific induction of Hsp70 and/or distinct MTs in the range of 11 kDa in *Heliscus* strain H1-H4 appear to be essential adaptive mechanisms to continuous heavy metal exposure.

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