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Enzyme characteristics of aminotransferase FumI of *Sphingopyxis* sp. MTA144 for deamination of hydrolyzed fumonisin B_1

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Abstract Fumonisins are carcinogenic mycotoxins that are frequently found as natural contaminants in maize from warm climate regions around the world. The aminotransferase FumI is encoded as part of a gene cluster of *Sphingopyxis* sp. MTA144, which enables this bacterial strain to degrade fumonisin B₁ and related fumonisins. FumI catalyzes the deamination of the first intermediate of the catabolic pathway, hydrolyzed fumonisin B₁. We used a preparation of purified, His-tagged FumI, produced recombinantly in *Escherichia coli* in soluble form, for enzyme characterization. The structure of the reaction product was studied by NMR and identified as 2-keto hydrolyzed

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Food Biotechnology Laboratory, Department of Food Sciences and Technology, University of Natural Resources and Life Sciences, Muthgasse 18, 1190 Vienna, Austria fumonisin B₁. Pyruvate was found to be the preferred co-substrate and amino group receptor ($K_{\rm M}$ =490 μM at 10 μ M hydrolyzed fumonisin B₁) of FumI, but other α -keto acids were also accepted as co-substrates. Addition of the co-enzyme pyridoxal phosphate to the enzyme preparation enhanced activity, and saturation was already reached at the lowest tested concentration of 10 µM. The enzyme showed activity in the range of pH 6 to 10 with an optimum at pH 8.5, and in the range of 6°C to 50°C with an optimum at 35°C. The aminotransferase worked best at low salt concentration. FumI activity could be recovered after preincubation at pH 4.0 or higher, but not lower. The aminotransferase was denatured after preincubation at 60°C for 1 h, and the residual activity was also reduced after preincubation at lower temperatures. At optimum conditions, the kinetic parameters $K_{\rm M}$ =1.1 μ M and $k_{\rm cat}$ =104/min were determined with 5 mM pyruvate as co-substrate. Based on the enzyme characteristics, a technological application of FumI, in combination with the fumonisin carboxylesterase FumD for hydrolysis of fumonisins, for deamination and detoxification of hydrolyzed fumonisins seems possible, if the enzyme properties are considered.

Keywords Mycotoxin · Detoxification · Degradation · Kinetics · Feed · Animal

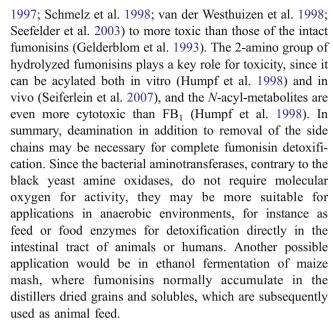
Introduction

Fumonisins are a group of structurally related mycotoxins produced by several species of *Fusarium* (Rheeder et al. 2002) as well as by *Alternaria alternata* (Chen et al. 1992), *Aspergillus niger* (Frisvad et al. 2007), and *Tolypocladium*



species (Mogensen et al. 2011). Fusarium verticillioides may be considered the most important fumonisin producer from the aspect of food and feed safety, as it is an important pathogen of maize in warm climate regions around the world. The fumonisins produced by this fungus were also the first to be isolated (Gelderblom et al. 1988; Bezuidenhout et al. 1988) and thoroughly studied (Marasas 2001). Fumonisin B₁ (FB₁), the most prevalent fumonisin of F. verticillioides, can cause equine leukoencephalomalacia (Marasas et al. 1988; Kellerman et al. 1990), porcine pulmonary edema (Harrison et al. 1990; Haschek et al. 2001), liver cancer in rats (Gelderblom et al. 1991), neural tube disorder in mice (Sadler et al. 2002; Gelineau-van Waes et al. 2005), alteration of the immune response in pigs (Taranu et al. 2005), and shows several other toxic effects in laboratory and domestic animals (Voss et al. 2007). Exposure to fumonisins is also associated with cancer and neural tube disorder in humans (Wild and Gong 2010). At the molecular level, fumonisins interfere with sphingolipid metabolism by inhibiting the enzyme ceramide synthase (Wang et al. 1991). The resulting imbalance of sphingolipids, which also have signaling functions as cellular messenger molecules, is responsible for toxic and carcinogenic effects of fumonisins (Merrill et al. 2001; Riley et al. 2001). However, other mechanisms of fumonisin toxicity and carcinogenicity were also proposed, including activation of mitogenactivated protein kinase (Wattenberg et al. 1996), lipid peroxidation (Abel and Gelderblom 1998), and alteration of the biosynthesis of other lipids in addition to the sphingolipids (Gelderblom et al. 2001).

The known fumonisin biodegradation pathways in black yeast strains (Blackwell et al. 1999) and bacteria (Duvick et al. 2003; Heinl et al. 2010) all start with the hydrolytic release of the two tricarballylic acid side chains, catalyzed by a fumonisin carboxylesterase. The second step is deamination of hydrolyzed fumonisin B₁ (HFB₁), either by an amine oxidase in black yeast strains (Blackwell et al. 1999) or an aminotransferase in bacterial strains (Heinl et al. 2010, 2011) (Fig. 1). Enzymatic detoxification of fumonisins may be a suitable approach to increase feed and food safety (Karlovsky 1999). However, it is unclear whether toxicity is sufficiently reduced by enzymatic cleavage of the two tricarballylic acid side chains. Hydrolyzed fumonisins were reported not to have cancerinitiating potency (Gelderblom et al. 1993), to be less hepatotoxic (Howard et al. 2002), less disruptive of sphingolipid metabolism, and not inducing neural tube disorder (Voss et al. 2009) in vivo. Hydrolysis of fumonisins by nixtamalization was reported to reduce, but not eliminate, toxicity (Hendrich et al. 1993; Voss et al. 1996). The in vitro effects of hydrolyzed fumonisins were reported to range from less (Flynn et al. 1997; Norred et al.



We have previously cloned the *fumI* gene of *Sphingo-pyxis* sp. MTA144, which catalyzes deamination of HFB₁ (Heinl et al. 2010), expressed the gene in *E. coli* and purified the enzyme (Hartinger et al. 2010). Here, we report the characterization of the enzyme together with its kinetic parameters as a starting point for evaluation of the application potential of the enzyme.

Materials and methods

Chemicals

FB₁ and ¹³C-FB₁ were obtained from Biopure Referenzsubstanzen GmbH (Tulln, Austria). HFB₁ was prepared by complete hydrolysis of FB₁ with purified recombinant carboxylesterase FumD (Heinl et al. 2010). All other chemicals were purchased from Sigma (St. Louis, MO, USA) or Sigma–Aldrich (Steinheim, Germany). Water was purified by reverse osmosis in an arium[®] RO 61316 system (Sartorius Stedim Biotech GmbH, Göttingen, Germany).

Recombinant enzyme production and purification

Construction of the expression vector pET-30a-AT144HIS, gene expression in *E. coli* ArcticExpress(DE3), and purification of the 6xHis-tagged recombinant aminotransferase FumI by immobilized nickel affinity chromatography were previously described (Hartinger et al. 2010). The recombinant enzyme FumI-HIS comprises the FumI sequence (GenBank ACS27061) with a C-terminal extension of ASSVD KLAAALEHHHHHH. The calculated molecular mass is 48,271.5 Da. Preparations of FumI-HIS were stored at -20°C after addition of glycerol to 25%.



Fig. 1 Initial steps of fumonisin B₁-degradation pathway of Sphingopyxis sp. MTA144: FB₁ (2-amino-12,16dimethyl-3,5,10-trihydroxy-14,15propan-1,2,3-tricarboxyicosane) is substrate of the fumonisin carboxylesterase FumD, which catalyzes hydrolytic cleavage of both tricarballylic acid (TCA) chains off the core chain to produce HFB₁ (2amino-12,16-dimethylicosane-3,5,10,14,15-pentol) and tricarballylic acid (1,2,3propanetricarboxylic acid). Aminotransferase FumI transfers the 2-amino group from HFB₁ to pyruvate, producing 2-keto-HFB₁ (3,5,10,14,15-pentahydroxy-12,16-dimethylicosane-2-one) and alanine

Enzyme activity assays

Possible co-substrates of the aminotransferase were tested in a reaction system containing 94 ng/ml FumI-HIS in 20 mM Tris–HCl pH 7.4, 50 mM NaCl, 2 mM CaCl₂, 20 μ M pyridoxal phosphate (PLP), and 0.1 mg/ml bovine serum albumin (BSA). HFB₁ as the substrate was employed at a concentration of 15 μ M, and no co-substrate or 15 μ M or 3 mM of the α -keto acids pyruvate, α -ketobutyrate, α -ketoglutarate, glyoxylate, or oxaloacetate, respectively, were added. Samples of 200 μ l were taken immediately after addition of the enzyme to the reaction mixture (t=0 h), as well as after 0.5, 1, and 3 h of incubation in a water bath

at 25°C. For correlation of enzyme activity with pyruvate and oxaloacetate concentration, the assay was performed as above except with 10 μ M HFB₁, and co-substrate concentrations were varied from 0.01 to 100 mM. Aminotransferase activity was determined by calculating the slope of the initial, linear reaction rates in Δ μ mol HFB₁ per liter per minute. Oxaloacetate was dissolved directly before use to avoid decarboxylation. To test the need for addition of PLP, the assay was performed as above with 15 μ M HFB₁ and PLP concentrations ranging from 0 to 400 μ M. The effect of salt on enzyme activity was tested by using 20 mM Tris–HCl pH 8.0 buffers with 1,000 mM, 100 mM, 10 mM NaCl, or without added NaCl, Teorell–Stenhagen buffer



pH 8.0 (Teorell and Stenhagen 1938), or 100 mM potassium phosphate buffer, all with 20 μ M PLP, 0.1 mg/ml BSA, 3 mM pyruvate, and 15 μ M HFB₁.

The optimum pH for enzyme activity was determined by setting Teorell–Stenhagen buffer to pH values ranging from pH 1.8 to 11.5 at 22°C (Teorell and Stenhagen 1938). The assay was performed using 67 ng/ml FumI-HIS, 10 μ M HFB₁, 20 μ M PLP, 0.1 mg/ml BSA, and 5 mM pyruvate at 25°C. Samples were taken before addition of enzyme and after 15, 30, 45, 60, 120, and 240 min. Enzyme activities were calculated from early time points, where reaction rates were linear, and are expressed as μ M 2-keto-HFB₁ formed per minute. The optimum temperature for enzyme activity was determined by using the same conditions as for the pH dependence, with the buffer set to pH 8.0 at 22°C, and incubation temperatures in water baths ranging from 6°C to 50°C. Reaction mixtures were temperature-equilibrated for 30 min before addition of enzyme.

To determine pH stability of FumI-HIS, the Teorell-Stenhagen buffer was set to pH values between pH 1.1 and 10.7, and 14 µg/ml of enzyme was incubated in these buffers at 35°C for 1 h. The residual enzyme activity was determined in the same buffer at pH 7.8 after dilution of FumI-HIS to 50 ng/ml, using 15 µM HFB₁, 20 µM PLP, 0.1 mg/ml BSA, and 5 mM pyruvate. Samples were taken, and reaction rates were determined as for the correlation of activity with pH and temperature. The thermal stability of FumI-HIS was determined by incubating 14 µg/ml of enzyme in Teorell-Stenhagen buffer, set to pH 8.0 at 22°C, for 1 h in water baths at temperatures ranging from 7°C to 60°C. Residual enzyme activity was measured at 35°C after dilution in the same buffer with final concentrations of FumI-HIS, HFB₁, PLP, BSA, and pyruvate as described for pH stability. Samples were taken, and reaction rates were calculated as for the determination of optimum pH and temperature.

Enzyme kinetics were determined by incubating 10 ng/ml FumI-HIS at 35°C in 20 mM Tris–HCl pH 8.0, 20 μ M PLP, 0.1 mg/ml BSA, and 5 mM pyruvate, while the concentration of HFB₁ was varied from 0.1 to 100 μ M. Samples were taken, and reaction rates were determined as above.

Liquid chromatography-mass spectrometry

Samples taken from enzyme assays were inactivated at 99°C for 10 min and stored frozen. After thawing, mixing, and centrifugation, aliquots of samples containing nominally 0.2 nmol HFB₁, but no more than 100 μ l, were dried at 50°C under a stream of nitrogen and dissolved in 200 μ l solvent with $^{13}\text{C-FB}_1$ as internal standard. Thus, the final nominal HFB₁ concentration for analysis was 1 μ M, or less if the original concentration was less than 2 μ M. All samples

were analyzed using the previously described instrumental setup and method (Heinl et al. 2010), to which 2-keto-HFB₁ has been integrated (Hartinger et al. 2010).

Preparative isolation of 2-keto-HFB₁ and structure determination by NMR

FumI-HIS was produced in E. coli ArcticExpress (DE3) as previously described (Hartinger et al. 2010). Biomass was resuspended in 1/100 of the culture volume, homogenized in a French Press, and 1.2 ml of the clarified lysate was added to 4.05 mg HFB₁ in 25 ml buffer (20 mM Tris-HCl pH 8.0, 6.6 mM pyruvate, 30 µM PLP, and 0.1 mg/ml BSA). After incubation at 30°C, complete deamination of HFB₁ was confirmed by liquid chromatography-mass spectrometry (LC-MS). The reaction mixture was centrifuged, lyophilized, and dissolved in 6 ml of 40% acetonitrile in water (v/v). After centrifugation, 600-µl portions were separated on a preparative Agilent 1100 HPLC system (Agilent, Waldbronn, Germany) consisting of a G1361A pump, a G2260A autosampler, an adjustable flow splitter (Analytical Scientific Instruments, CA, USA), an evaporative light-scattering detector (SEDEX LT-ELSD Model 85, Sedere, Alfortville, France), a G1365B multiwavelength detector, and a G1364B fraction collector. A Gemini-NX C18 column (150×21.2 mm, 5-µm particle size, 110-Å pore size; Phenomenex, Aschaffenburg, Germany) protected by a Gemini C18 pre-column (15×21.2 mm) was operated at a flow rate of 16 ml/min using gradient elution. Mobile phase A, consisting of acetonitrile/H₂O (10:90, v/v), was degassed by membrane filtration. Acetonitrile was used as mobile phase B. The gradient was as follows: 0-0.2 min: 0% B, 0.2-5 min: linear increase to 95% B, 5-6.4 min: isocratic elution at 95% B, 6.4-6.5 min: linear decrease to 0\% B, 6.5-9.5 min: re-equilibration at 0\% B. The column effluent was split 1:70, one part moving into the evaporative light-scattering detector and the main part passing through the multivariate wavelength detector to the fraction collector. The 2-keto-HFB₁ eluted between 4.85 and 5.7 min and fractions were collected in that time window. The combined fractions were evaporated to dryness on a rotary evaporator, dissolved in 7 ml of ethyl acetate, and transferred into an 8-ml screw cap vial; the ethyl acetate solution was evaporated to dryness under a stream of nitrogen at 50°C. The 2-keto-HFB₁, a viscous, clear, and colorless oily substance, was stored at 4°C until NMR analysis.

¹H- and ¹³C-NMR spectra were obtained from CD₃CN solutions using an Avance DRX-400 FT-NMR spectrometer (Bruker BioSpin, Rheinstetten, Germany), operating at 400.13 MHz for ¹H and 100.62 MHz for ¹³C, at 295 K using a 5-mm inverse broadband *Z*-gradient probe head. Data were recorded and evaluated using TOPSPIN 1.3



software (Bruker Biospin). All pulse programs were taken from the Bruker software library. Chemical shifts were established on the basis of residual solvent resonances.

Results

Biochemical characterization of FumI-HIS

Clarified cell lysates of $E.\ coli$ overexpressing the fumI gene were previously used to obtain preliminary information, namely that HFB $_1$ is a substrate and that an α -keto acid such as pyruvate is required as co-substrate for the activity of the aminotransferase FumI (Heinl et al. 2010). The detailed biochemical characterization reported here was made using FumI-HIS, which was prepared and purified to electrophoretic homogeneity as described before (Hartinger et al. 2010). This enzyme preparation was found to be stable for several months when stored in 25% glycerol at -20° C. Clarified cell lysates of $E.\ coli$ ArcticExpress (DE3) producing FumI and FumI-HIS from the same expression vector pET-30a under identical conditions showed a very similar HFB $_1$ deamination activity, indicating that the Histag does not affect the enzymatic activity.

The α -keto acids pyruvate, α -ketobutyrate, α -ketoglutarate, glyoxylate, and oxaloacetate were tested as potential co-substrates for aminotransferase FumI (Fig. 2). No deamination of HFB $_1$ was detectable when no co-substrate was added or in the presence of α -ketoglutarate, while pyruvate, oxaloacetate, glyoxylate, and α -ketobutyrate, when included at 3 mM concentration, supported the reaction, with HFB $_1$ deamination efficiency decreasing in this order. When tested

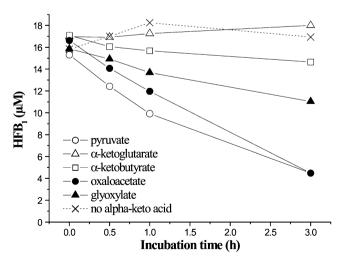


Fig. 2 Comparison of HFB $_1$ deamination activity of FumI-HIS with five different α -keto acids as co-substrate. The reactions were incubated in 20 mM Tris–HCl buffer (pH 7.4) at 25°C with 94 ng/ml FumI and 3 mM α -keto acid

with 10 μ M HFB₁, the enzyme showed maximal activity at 10 mM pyruvate, and the $K_{\rm M}$ for pyruvate was 490 μ M. Deamination activity was significantly reduced when the concentration of the co-substrate pyruvate was in the same range as that of HFB₁, and substrate inhibition was observed for pyruvate concentrations in the range of 50–100 mM (Fig. 3). With oxaloacetate as co-substrate, reaction rates were lower, and the $K_{\rm M}$ for oxaloacetate was 4,150 μ M (Fig. 3). Addition of PLP enhanced the aminotransferase activity of FumI-HIS, and saturation was reached already at the lowest concentration tested (10 μ M PLP) since a further increase in PLP concentration to 20, 40, 60, 80, 100, 120, 140, 160, 180, 200, 300, and 400 μ M in the assay mixture did not result in a further increase in the reaction rate. Without added PLP, FumI-HIS showed about 23% of the maximum activity.

Addition of NaCl to the 20 mM Tris-HCl (pH 8.0) reaction buffer reduced the HFB₁ deamination rate (Fig. 4). When using 100 mM potassium phosphate or Teorell–Stenhagen buffer, enzyme activity was also lower than in 20-mM Tris-HCl buffer of the same pH. Nevertheless, Teorell–Stenhagen buffer was used to determine the optimum pH for enzyme activity, since this buffer can be adjusted to any pH between 2.0 and 12.0 (Teorell and Stenhagen 1938). Aminotransferase FumI-HIS was active in the range of pH 6.5 to 9.7 and showed a rather sharp optimum at pH 8.0 and 8.5 (Fig. 5a). The enzyme was active over a broad temperature range of 6°C to 45°C with an optimum for the 30 min assay at 35°C (Fig. 5b). Significant and rapid thermal inactivation was observed at 50°C, where traces of 2-keto-HFB₁ were formed only at the

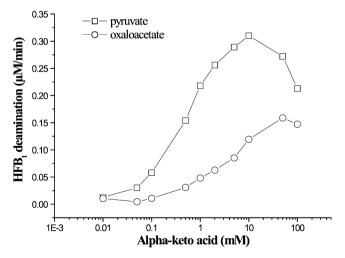


Fig. 3 Correlation between pyruvate concentration or oxaloacetate concentration and HFB $_1$ deamination activity. The reactions were incubated with 94 ng/ml FumI-HIS and 10 μM HFB $_1$ in 20 mM Tris–HCl (pH 7.4) with 0.1 mg/ml BSA, 20 μM PLP, and the indicated concentration of pyruvate or oxaloacetate at 25°C. Samples were taken and analyzed, and reaction rates were calculated as described in Materials and Methods. The reaction mixtures without co-substrates (negative controls), which are not shown in the semi-logarithmic plot, gave HFB $_1$ deamination rates of zero



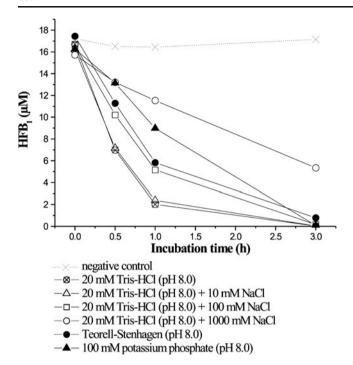
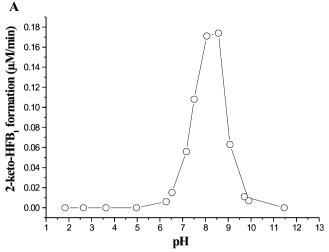


Fig. 4 Effect of different buffers and salt concentrations on the HFB₁-deamination rate of aminotransferase FumI-HIS. Negative control: 20 mM Tris-HCl (pH 8.0) with buffer instead of enzyme

beginning of the incubation period (data not shown). Chromatograms of samples from the HFB₁ deamination reaction of this experiment at 35°C are shown in Fig. 6. Furthermore, FumI-HIS was incubated with intact FB₁ instead of hydrolyzed FB₁ under identical conditions, but FB₁ was not deaminated. Reaction with a hydrolyzed total fumonisin extract from F. verticillioides culture material performed under the same conditions showed that, in addition to HFB₁, HFB₂, and HFB₃ were substrates of FumI-HIS as well. When FumI-HIS was preincubated at an acidic pH and 35°C for 1 h before HFB₁ deamination was measured, the activity was reduced after incubation at pH 6.5 or less, and completely abolished after incubation at pH below 4.0 (Fig. 7a). Incubation at pH 8.0 for 1 h at various temperatures resulted in a steady decrease of FumI activity with increasing temperature, and the enzyme was completely inactivated after incubation at 60°C. Even after incubation at 7°C, a 15% loss of enzyme activity compared to non-preincubated FumI was determined (Fig. 7b).

The apparent steady-state kinetic parameters of FumI-HIS catalyzed HFB₁ deamination were determined for a saturating co-substrate concentration of 5 mM pyruvate in 20 mM Tris–HCl buffer (pH 8.0) at 35°C, and the Michaelis–Menten plot is shown in Fig. 8. The apparent $V_{\rm max}$ value of 0.022 μ M/min was reached at 5 μ M HFB₁ when 10 ng/ml of FumI-HIS was present in the reaction mixture. A further increase in substrate (HFB₁) concentration resulted in a significant reduction of the reaction rate,



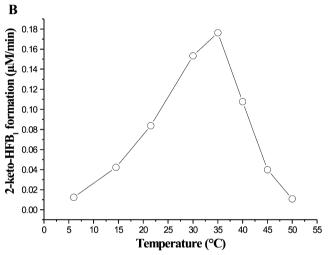


Fig. 5 Determination of optimum pH (a) and temperature (b) for FumI-HIS activity. The reactions were incubated with 67 ng/ml FumI-HIS and 10 μ M HFB₁ in Teorell–Stenhagen buffer, with supplements as described in Materials and Methods, set to the indicated pH values (a) or to pH 8.0 (b), at 25°C (a) or the indicated temperatures (b)

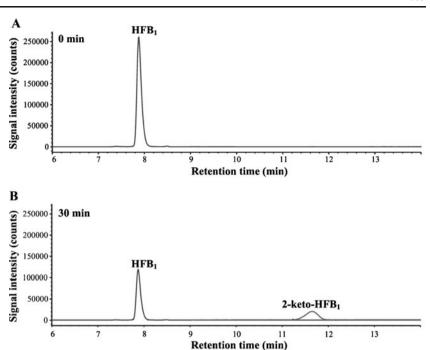
indicating substrate inhibition. The $k_{\rm cat}$ value was calculated to be $104~{\rm min}^{-1}$ using the theoretical molecular mass of FumI-HIS, and the Michaelis constant $K_{\rm M}$ for HFB₁ was determined to be $1.1~{\rm \mu M}$. The specific HFB₁ deamination activity was $2.2~{\rm \mu mol/min/mg}$.

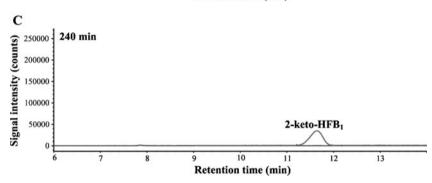
Structure determination of the reaction product by NMR

From the 1D ¹H- to ¹³C-APT-NMR spectra, it was obvious that the sample did not contain a single species, but three different forms. In order to analyze the molecular structures of these compounds, ¹H¹H COSY, ¹H¹³C HSQC, and ¹H¹³C HMBC spectra were recorded. Using these two-dimensional methods, all of the structures could be elucidated unambiguously, and complete assignments of ¹H and ¹³C signals were established (Table 1). Thus, the structures were



Fig. 6 LC–MS chromatograms of samples from a HFB $_1$ transamination reaction. Samples of 10 μ M HFB $_1$ in Teorell–Stenhagen buffer (pH 8.0) at 35°C were taken before (a), 30 min (b), or 240 min (c) after addition of purified aminotransferase Fuml-HIS to a final concentration of 67 ng/ml. HFB $_1$ and 2-keto-HFB $_1$ were separated on a C8 reversed phase column in a formate–acetonitrile gradient and detected in SIM mode





identified as 2-keto-HFB₁ and two cyclic hemiketals formed by ring closure between the 2-keto moiety and the 5-OH. While the hemiketals have already been described (Blackwell et al. 1999), the open-chain keto form has not yet been characterized.

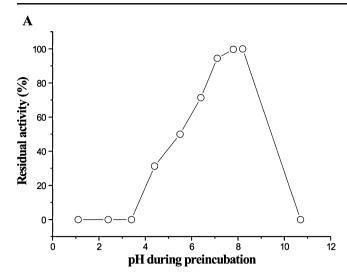
Discussion

Gastrointestinal detoxification of fumonisins by specific enzymes is a promising concept to ameliorate the effects that fumonisins, which are frequently found as natural contaminants of maize in large parts of the world, have on the health and performance of domestic animals. It is important to know the characteristics of an enzymecatalyzed reaction before a technological application can be considered. We determined some key properties of the aminotransferase FumI of *Sphingopyxis* sp. MTA144 with respect to its reaction with fumonisins. The enzyme is active with hydrolyzed FB₁, the first intermediate in all known pathways of FB₁ catabolism, and not with FB₁ itself

and catalyzes the transfer of the 2-amino group onto a suitable acceptor (Heinl et al. 2010). Since the corresponding amino group of structurally similar sphinganine is acylated by ceramide synthase, which is the molecular target that is inhibited by fumonisins (Wang et al. 1991), the deamination reaction can be considered as important for detoxification of fumonisins. To our knowledge, no reports on the toxicity of 2-keto-HFB₁ have been published, and we have not studied its toxicity yet. However, the amino group is thought to play a key role for the toxicity of FB₁ (Gelderblom et al. 1993; Lu et al. 1997; Norred et al. 2001; Fernandez-Surumay et al. 2004), and obviously, 2-keto-HFB₁ cannot be acylated in a similar way as HFB₁ (Humpf et al. 1998; Seiferlein et al. 2007). Therefore, it seems reasonable to speculate that 2-keto-HFB₁ may be non-toxic.

Numerous aminotransferases have been identified and studied since the first description of this class of enzymes (Needham 1930), and several are used in industrial applications for the production of a range of natural and non-natural amino acids and amines (Taylor et al. 1998;





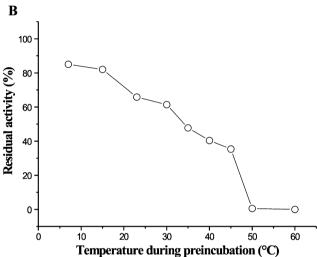


Fig. 7 Stability of FumI-HIS at various pH values (a) and temperatures (b). Fourteen micrograms per milliliter FumI-HIS was incubated in Teorell–Stenhagen buffer at the indicated pH at 35°C (a) or in the same buffer at pH 8.0 at the indicated temperature (b) for 1 h. Residual enzyme activity was determined as described in Materials and Methods. The measured activities after incubation at pH 8.0 (a) or without preincubation (b) were set to 100%

Hwang et al. 2005; Zhu and Hua 2009). However, this is the first characterization of an aminotransferase that catalyzes deamination of a fumonisin. Our work may lead to the further expansion of the range of aminotransferase applications to mycotoxin detoxification. Since FB₁ is not a substrate for FumI, a possible technological application of the enzyme will require its combined use together with a fumonisin carboxylesterase such as FumD, so that HFB₁, the substrate of FumI, is generated by hydrolytic cleavage of the two tricarballylic acid side chains of FB₁. In this respect, the substrate specificity of FumI is quite remarkable, since the direct vicinity of the 2-amino group is identical in FB₁ and HFB₃ (Fig. 1). Since HFB₂ and HFB₃

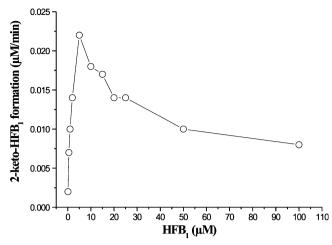


Fig. 8 Initial HFB $_1$ deamination reaction rate as a function of substrate concentration. The reactions were performed with 10 ng/ml FumI-HIS and 5 mM pyruvate, a saturating co-substrate concentration, at 35°C in 20 mM Tris–HCl pH 8.0, 20 μM PLP, 0.1 mg/ml BSA. HFB $_1$ concentrations ranged from 0.1 to 100 μM

are also deaminated by the enzyme, the 10-hydroxyl and 5-hydroxyl groups of HFB₁, which are lacking in HFB₂ and HFB3, respectively, are apparently not essential for substrate recognition. The ability of FumI to use several different α-keto acids as amino group acceptors is also noteworthy. Even though pyruvate was the preferred co-substrate when various α -keto acids were tested at the identical molar concentrations in our study, it is tempting to speculate that the relaxed co-substrate specificity of FumI may have biological significance for its function in Sphingopyxis sp. MTA144. Depending on the available α -keto acids, FumI could thus produce alanine from pyruvate, aspartate from oxaloacetate, or glycine from glyoxylate in vivo. The affinity of FumI for its co-substrate (Fig. 3), which is much lower than the affinity for HFB₁ (Fig. 8) when judged by the $K_{\rm M}$ values, may also be important for the enzyme in vivo, since the compatible α -keto acids are also intermediates in energy metabolism. In this context, it is interesting to note that the fum gene cluster of Sphingopyxis sp. MTA144 encodes three genes, the tricarballylate proton symport pump FumG, the tricarballylate dehydrogenase FumE, and the citrate utilization protein B FumF, which supposedly allow utilization of the two tricarballylic acid side chains of FB₁ and their conversion to citric acid (Heinl et al. 2010). Oxaloacetate and glyoxylate, which both accept the amino group during the FumI-catalyzed reaction, can both be generated from citric acid in the Krebs cycle or the glyoxylate cycle. For a technological application of FumI, this co-substrate requirement together with the comparatively low $K_{\rm M}$ value implies that pyruvate or one of the other suitable α -keto acids will have to be present or will have to be provided together with the enzyme at



Table 1 ¹H and ¹³C NMR assignment of 2-keto-HFB₁ and its hemiketal forms

	2-Keto-HFB ₁		α-Hemiketal ^a		β-Hemiketal ^a	
	¹ H	¹³ C	¹ H	¹³ C	¹ H	¹³ C
1	2.14	25.8	1.32	22.8	1.29	25.0
2	_	212.4	_	106.9	_	102.2
3	4.28	75.7	3.87	78.2	3.80	76.8
4	1.72, 1.50	41.4	2.40, 1.35	40.5	2.23, 1.52	38.9
5	3.70	68.7	3.97	77.0	3.80	76.1
6	1.50-1.40	38.8	1.58, 1.44	37.4	1.45-1.40	38.5
7	1.45-1.25	≈27	1.45-1.25	≈27	1.45-1.25	≈27
8	1.45-1.25	≈27	1.45-1.25	≈27	1.45-1.25	≈27
9	1.40-1.35	39.4	1.40-1.35	39.4	1.40-1.35	39.4
10	3.59	69.8	3.59	69.8	3.59	69.8
11	1.48, 1.06	44.4	1.48, 1.06	44.4	1.48, 1.06	44.4
12	1.87	26.9	1.87	26.9	1.87	26.9
12-Me	0.94	21.7	0.94	21.7	0.94	21.7
13	1.40-1.25	40.1	1.40-1.25	40.1	1.40-1.25	40.1
14	3.61	70.2	3.61	70.2	3.61	70.2
15	3.16	80.2	3.16	80.2	3.16	80.2
16	1.55	35.6	1.55	35.6	1.55	35.6
16-Me	0.84	16.4	0.84	16.4	0.84	16.4
17	1.61, 1.20	32.2	1.61, 1.20	32.2	1.61, 1.20	32.2
18	1.36, 1.20	30.0	1.36, 1.20	30.0	1.36, 1.20	30.0
19	1.25-1.35	23.9	1.25-1.35	23.9	1.25-1.35	23.9
20	0.90	14.5	0.90	14.5	0.90	14.5

^a α/β according to Blackwell et al. (1999)

millimolar concentration to enable efficient and fast HFB_1 deamination. However, if the enzyme is to be used for gastrointestinal HFB_1 deamination, this co-substrate requirement might be covered to some extent by the presence of α -keto acids in the chymus. It cannot be excluded that other amino group acceptors that we have not yet identified may enable higher reaction rates even at low concentration.

Blackwell et al. have noted before that 2-keto-HFB₁ can undergo cyclization between the 5-hydroxyl group and C-2 of 2-keto-HFB₁ to form two isomers of a hemiketal, and that this reaction is favored by the presence of methanol in the solvent (Blackwell et al. 1999). We made the same observation and adapted our procedure for purification and NMR characterization of 2-keto-HFB₁ to work without methanol. Nevertheless, the NMR characterization showed that our preparation of 2-keto-HFB₁ was not pure, and the signals indicative of the hemiketals were still present. As the C-1-C-5 part of 2-keto-HFB₁ resembles a simplified ketose structure, an equilibrium between the open-chain and cyclized forms may be the explanation for this behavior since in carbohydrate chemistry, such equilibria are ubiquitous. This may also be one reason that the 2-keto-HFB₁ peaks that eluted from a C8 HPLC column were broader than the HFB₁ peaks.

Overexpression of FumI in E. coli and the subsequent purification procedure (Hartinger et al. 2010) result in enzyme preparations that apparently are not saturated with the prosthetic group PLP, as can be concluded from the finding that addition of PLP to such an enzyme preparation enhances activity. Aminotransferases were reported to occur as mixtures of their apo- and holo-forms also when isolated from their natural environment (Moss 1976). Increasing the PLP concentrations above 10 µm in the reconstitution experiments of recombinant FumI-HIS showed no further increase of aminotransferase activity, which is consistent with the tight binding of PLP that has been reported for other aminotransferases (Ford et al. 1980; Eliot and Kirsch 2004). If batches of enzyme need to be prepared for technological application in the future, they can be saturated with low concentrations of PLP before their application to ensure that FumI is predominantly available as a holoenzyme.

FumI-HIS displayed HFB₁ deamination activity in all buffers we tested, and these covered a wide range of salt concentrations (Fig. 4). This indicates that the proposed application of FumI is unlikely to be limited by buffer salt requirements. The lack of enzyme activity at acidic pH (Fig. 5a) rules out that FumI is already active in the stomach when provided as an animal feed supplement.



However, FumI can be expected to function in sections of the upper intestine where the pH is close to neutral or even slightly higher. The temperature optimum of 35°C and good activity at 40°C found for FumI-HIS (Fig. 5b) imply that the actual body temperature of animals, which e.g. was reported to be 38.8°C for piglets (Ingram and Legge 1970), should be well suited for high enzyme activity. Based on our determination of pH stability (Fig. 7a), it will be necessary to provide FumI in a formulation that enables shuttling through the stomach without inactivation at low pH. Simultaneously, the enzyme could thus be protected from proteolytic degradation. Technologies for microencapsulation and targeted release in the intestine have previously been developed (Champagne and Fustier 2007).

We were surprised to see a reduction of HFB_1 deamination activity after preincubation of FumI-HIS at 7°C, because we had observed that a preparation of enzyme that was stored in the fridge apparently retained full activity over several weeks. Maybe handling of the enzyme at low concentration contributed to the observed loss of activity. Likewise, the maximal specific reaction rate measured in the experiment shown in Fig. 8, using 10 ng/ml FumI-HIS, was lower than in the experiment shown in Fig. 3, when 94 ng/ml FumI-HIS was used.

The $K_{\rm M}$ value we determined for HFB₁ deamination, 1.1 µM, corresponds to 0.794 mg/kg FB₁. Since the fumonisin concentrations frequently found in maize (Binder et al. 2007; Gonzalez Pereyra et al. 2008; Gong et al. 2009; Monbaliu et al. 2010) and known to have toxic effects on animals (Ross et al. 1991; Taranu et al. 2005) are higher, FumI should have sufficient affinity for the substrate HFB₁ to enable detoxification in the range of fumonisin concentrations that are relevant in agriculture. The decrease of FumI activity observed at HFB₁ concentrations higher than 10 μM, which is equivalent to 7.22 mg/l FB₁, can be interpreted as substrate inhibition. Since the guidance value for the maximum acceptable concentration of FB₁ and FB₂ in e.g. pig feed is 5 mg/kg according to the European Commission Recommendation 2006/576/EC, this observed inhibition of activity at higher HFB₁ concentrations may not be relevant from a technological point of view. In case enzymatic detoxification by a combination of FumD and FumI of a highly contaminated commodity is required, substrate inhibition of FumI may be avoided by adapting FumD activity to that of FumI, so that the HFB₁ is continuously formed in sub-inhibitory concentrations.

Feed enzymes may be useful for the degradation of antinutritive substances that are naturally contained in agricultural commodities, or to enhance the nutritional value of feed. The use of phytase as a feed enzyme to release phosphate from phytate that is naturally contained in plant material is well established (Lei and Porres 2003;

Rao et al. 2009). Based on the enzyme characteristics reported here, FumI may also be suitable for future applications as a feed enzyme to enhance the salubriousness of maize that is naturally contaminated with fumonisins, if the specific properties and requirements of FumI are considered in this application.

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Conflict of interest The authors declare no conflict of interest.

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