Hindawi Publishing Corporation BioMed Research International Volume 2015, Article ID 245649, 8 pages http://dx.doi.org/10.1155/2015/245649

Research Article

Purification and Characterization of a Polyextremophilic α -Amylase from an Obligate Halophilic Aspergillus penicillioides Isolate and Its Potential for Souse with Detergents

Imran Ali,^{1,2} Ali Akbar,³ Mohammad Anwar,² Sehanat Prasongsuk,¹ Pongtharin Lotrakul,¹ and Hunsa Punnapayak¹

Correspondence should be addressed to Imran Ali; imranalisheik@gmail.com and Hunsa Punnapayak; phunsa@chula.ac.th

Received 13 October 2014; Revised 25 December 2014; Accepted 25 December 2014

Academic Editor: Subash C. B. Gopinath

Copyright © 2015 Imran Ali et al. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

An extracellular α -amylase from the obligate halophilic *Aspergillus penicillioides* TISTR3639 strain was produced and enriched to apparent homogeneity by ammonium sulfate precipitation and Sephadex G100 gel filtration column chromatography. The mass of the purified amylase was estimated to be 42 kDa by SDS-PAGE. With soluble starch as the substrate it had a specific activity of $118.42~\mathrm{U\cdot mg^{-1}}$ and V_{max} and optimum activity at pH 9, 80°C, and V_{max} activity, while V_{max} and V_{max} and V_{max} and V_{max} and V_{max} and V_{max} and optimum activity at pH 9, 80°C, and V_{max} and $V_$

1. Introduction

Hypersaline environments are caused by the evaporation of water and they are also called thallasohaline environment. Due to the evaporation process the sodium chloride (NaCl) concentration rises above 300 psu. Many microbial communities have been found from the sesalterns. Halophilic microorganisms are extremophiles that are able to survive in and may require salt for their growth [1]. Fungi that are isolated from hypersaline environments with a salinity above $100~{\rm g}\cdot{\rm L}^{-1}$ and are able to grow *in vitro* at a 175 g·L⁻¹ salt concentration are categorized as halophilic fungi [2, 3]. Very few fungi have been reported yet to inhabit the hypersaline habitats. Their function in these environments is still not fully

understood [2]. Unlike other microbes, the fungi can grow independent of salt concentration in saline environments [1]. However, obligate halophilic fungi are those that are unable to grow in the absence of a salt concentration [4].

Extremophilic microorganisms adopt different strategies to survive in extreme available conditions. They harbor different metabolites such as enzymes that can work at extreme conditions [5] and so are of interest for diverse biotechnological applications. Halophiles are currently used in several fermentation processes [6], such as for the production of bioactive compounds [7], biorhodopsin, biosurfactants, food additives, and biocompatible solutes [8]. Extreme halophiles have been increasingly investigated for their hydrolytic enzymes since these have potential uses in several

¹Plant Biomass Utilization Research Unit, Department of Botany, Faculty of Science, Chulalongkorn University, Bangkok 10330, Thailand

²Institute of Biochemistry, University of Balochistan, Quetta 87300, Pakistan

³Department of Microbiology, University of Balochistan, Quetta 87300, Pakistan

industrial applications [9, 10]. However, the use of halophilic microorganisms and their metabolites has largely involved halophilic bacteria [11]. Despite the fact that halophilic fungi, especially the obligate strains, are better sources of extracellular enzymes, they have not been investigated very much for their potential in biotechnological applications [3, 12]. α -Amylase (1,4- α -D-glucan glucanohydrolase, EC 3.2.1.1) is a class of important industrial enzymes that are used in the food, textile, laundry, and pharmaceutical industries [13, 14] and currently form about 25% of the total enzyme market [15]. Although many microbial amylases have been used as an additive in laundry detergents [16], they do not perform well in hard or saline water, which then limits their use in such areas [3, 12].

Recently, the obligate halophilic *Aspergillus penicillioides* TISTR 3639 strain was isolated from an extreme hypersaline environment (a man-made solar saltern) in the Ban Laem district of Phetchaburi province, Thailand [1, 4]. The fungus was found to be positive for extracellular α -amylase activity [3] by plate screening method (Supplementary Figure S1; see Supplementary Material available online at http://dx.doi.org/10.1155/2015/245649). In this study, the purification and characterization of α -amylase from *A. penicillioides* TISTR 3639 were performed. Due to its polyextremophilic properties, the potential of using this enzyme as a laundry detergent additive was then investigated.

2. Materials and Methods

- 2.1. Growth Conditions for Enzyme Production. The A. penicillioides TISTR3639 strain was grown in 150 mL flasks containing 100 mL of production medium (PM) at room temperature (25 \pm 2°C) at 150 rpm for 14 d. The PM was made according to Ali et al. [3] with a few modifications such that the composition was composed of 10 g·L $^{-1}$ soluble starch, 3.0 g·L $^{-1}$ mycological peptone, 100 g·L $^{-1}$ NaCl, 8.0 g·L $^{-1}$ CaCO $_3$, 6.6 g·L $^{-1}$ (NH $_4$) $_2$ SO $_4$, 3.5 g·L $^{-1}$ KH $_2$ PO $_4$, 0.15 g·L $^{-1}$ FeSO $_4$ ·7H $_2$ O, and 0.10 g·L $^{-1}$ MgSO $_4$ ·7H $_2$ O.
- 2.2. Amylase Purification. Enrichment to apparent homogeneity of the α -amylase was performed by ammonium sulfate precipitation and Sephadex G100 gel filtration chromatography as previously described [12, 17, 18]. The 14 d grown culture broth (100 mL) was centrifuged at 13,000 ×g at 4°C for 10 min and the supernatant was harvested. The amylase was then precipitated by bringing the supernatant to 90% saturation (NH₄)₂SO₄, storing overnight at 4°C, and harvesting the insoluble fraction by centrifugation at $12,000 \times g$ for 30 min. The pellet was then dissolved in 100 mM Tris-HCl buffer (pH 8) and dialyzed against the same buffer for 48 h. The dialyzate was then subjected to Sephadex G100 gel filtration using a 2.6 cm × 150 cm column, preequilibrated in and then eluted with 25 mM Tris-HCl buffer (pH 8) containing 5 mL·L⁻¹ Triton X-100 at a flow rate of 30 mL·h⁻¹. Fractions (5 mL) were collected and each was tested for α amylase activity and total protein content.

Determination of the purity and molecular weight of the enriched α -amylase was performed using sodium dodecyl

sulphate polyacrylamide gel electrophoresis (SDS-PAGE) resolution (150 g·L $^{-1}$ resolving gel) followed by coomassie blue staining, as reported by Hmidet et al. [18]. The purified amylase was mixed at a 1:5 volume ratio with the loading buffer (10 mM Tris-HCl, pH 8, 25 g·L $^{-1}$ SDS, 50 mL·L $^{-1}\beta$ -mercaptoethanol, 10 mL·L $^{-1}$ glycerol, and 0.002 g·L $^{-1}$ bromophenol blue). Prior to loading and electrophoresis, the sample was denatured and reduced by heating at 100°C for 5 min. Gels were stained with 2.5 g·L $^{-1}$ Coomassie Brilliant Blue R250 in 450 mL·L $^{-1}$ ethanol-100 mL·L $^{-1}$ acetic acid and destained with 50 mL·L $^{-1}$ ethanol-7.5 mL·L $^{-1}$ acetic acid. The molecular weight was determined in comparison to the Unstained Precision Plus Protein 161 molecular marker kit (Bio-Rad, USA).

- 2.3. α -Amylase Assay. α -Amylase activity was determined by the 3,5-dinitrosalicylic acid (DNS) method as described by Miller [19] using $10\,\mathrm{g\cdot L^{-1}}$ soluble starch as the substrate. The reaction mixture (0.1 mL enzyme solution, 0.5 mL 0.1 M phosphate buffer, and 5 mg soluble starch) was incubated at $40\,^\circ\mathrm{C}$ for $10\,\mathrm{min}$. The reaction was then stopped by the addition of 3 mL of DNS and heating in a boiling water bath for 5 min. After cooling, $10\,\mathrm{mL}$ of water was added and the absorbance of the reaction mixture was read at $540\,\mathrm{nm}$ (A₅₄₀). One unit (U) of enzyme activity was defined as the amount of enzyme that produced $1\,\mu\mathrm{mol}$ of glucose in 1 min.
- 2.4. Protein Estimation. The amount of protein was estimated by the method of Lowry [20], using bovine serum albumin (BSA) as the standard.
- 2.5. Characterization of the Enriched α -Amylase. The effect of the pH, temperature, and NaCl concentration on the enriched amylase enzyme activity was evaluated by sequential univariate analysis of the pH, temperature, and NaCl concentration, respectively, and monitoring the relative enzyme activity (the highest activity was referred to as 100%) as the selected parameter. For evaluation of the optimal pH, the reaction mixture was incubated in 0.1 M acetate buffer for pH 5-6 and 0.1 M phosphate buffer for pH 7–12, at a constant 30°C with no added NaCl. For evaluation of the optimal temperature, the reaction mixture was incubated at a temperature range of 40-100°C in 0.1 M phosphate buffer at the optimal pH (as determined above). Finally, for evaluation of the optimal salinity level, the sample mixture in 0.1 M phosphate buffer at the found optimal pH was supplemented with NaCl to a final concentration of 0-500 g·L⁻¹ and incubated at the optimal temperature.

The effect of various metal ions, or the enzyme inhibitors β -mercaptoethanol and EDTA, on the enzyme activity was investigated by separately adding BaCl₂, CaCl₂, FeCl₂, HgCl₂, MgCl₂, and ZnCl₂ to the reaction mixture at a final concentration of 2 mM. The relative amylase activity (%) was calculated in comparison to that without any additives.

2.6. Determination of the Kinetic Parameters of the Enriched α -Amylase. The kinetic parameters of the enzyme were determined by incubating the enriched α -amylase with 0.1–40 g·L⁻¹ soluble starch (substrate) under the previously found



FIGURE 1: SDS-PAGE analysis of the enriched α -amylase from A. penicillioides TISTR 3639. Lane 1: molecular mass ladder; lane 2 the enriched α -amylase, showing a single band at approximately 42 kDa.

optimum conditions. The $V_{\rm max}$ and K_m values were then calculated from the Lineweaver-Burk plot.

2.7. Compatibility of the Enriched α -Amylase with Commercial Detergents. The compatibility of the enriched α -amylase with three commercial detergents (a liquid detergent (A) and two powdered detergents (B and C) bought from Talad Thai market, Pathumthani, Thailand) was determined in terms of the respective enzyme activities in comparison with that of the enriched enzyme in the absence of the detergents.

The detergent solutions were prepared by dissolving the powdered or liquid detergent in distilled water to $7\,\mathrm{g}.\mathrm{L}^{-1}$ and heating at $100\,^\circ\mathrm{C}$ for 90 min to denature any enzyme activity present in the solutions. The cooled solutions were then mixed with the purified amylase at a 1:1 volume ratio and incubated for 1 h at $40\,^\circ\mathrm{C}$. The residual enzyme activity was calculated in comparison with the control (distilled water instead of the detergents solution), expressed as the relative activity (%) of the control experiment.

- 2.8. Performance of the Enriched α -Amylase with Commercial Detergents in Varying NaCl Concentrations. For the performance test, the control mixture was made using 0.1 mL of the enriched α -amylase from A. penicillioides TISTR3639 in 0.5 mL of 0.1 M phosphate buffer (pH 7) containing 10 g·L⁻¹ of soluble starch and 0–50 g·L⁻¹ NaCl, while the three detergents (A, B, and C) were assayed as above except for adding the respective detergent instead of the enriched amylase. A commercial α -amylase from A. oryzae (Sigma Aldrich, Germany) was also used for comparison.
- 2.9. Statistical Analysis. Each experiment with the required controls was performed in triplicate and the data are presented as the mean \pm one standard deviation (SD). Significance of the differences between means was tested for by analysis of variance (ANOVA) and Duncan's multiple means

tests (DMMT) on the parametric or arc-sine square root transformed data using the SPSS software, where a P value of less than 0.05 was considered as significant.

3. Results

3.1. Enrichment of the α -Amylase. The initial 90% saturation (NH₄)₂SO₄ cut gave a 2.5-fold increased enzyme specific activity for a 32% yield loss and a 76% total protein reduction (Table 1). Following Sephadex G100 gel filtration, the eluted fraction showing the highest amylase activity was then evaluated for its apparent purity and molecular weight by SDS-PAGE resolution and coomassie blue staining. Only a single protein band was visualized suggesting the apparent enrichment of the enzyme to homogeneity (Figure 1). Overall, the enzyme activity was increased 6.96-fold following the 90% saturation (NH₄)₂SO₄ precipitation and Sephadex G100 gel filtration for a loss of 56% activity. The specific activity of the enriched α -amylase was estimated to be 118.4 ± 1.2 U·mg⁻¹ (Table 1), with an estimated molecular weight of ~42 kDa by SDS-PAGE analysis (Figure 1).

3.2. Characterization of the Enriched α -Amylase. The effect of increasing the pH (range 5–12) on the enriched α -amylase activity, when assayed at 30°C, is shown in Figure 2(a), where the enzyme activity was higher (>80%) in a neutral to moderately alkaline pH (7–10). The enzyme activity declined above and below pH 9, but this decline was more marked with increasing pH above 9 (64% activity at pH 12) than with decreasing pH (~75 and ~73% at pH 6 and 5, resp.).

Increasing the temperature increased the activity of the enriched α -amylase, as assayed at the more optimal pH 9 (Figure 2(b)), to peak at 80°C (100±0.81%), and then declined at temperatures above 80°C, but it still retained more than 60% of its initial activity at 100°C. Thus, the thermophilic nature of the enzyme at temperatures higher than 60°C was clearly revealed.

With respect to the effect of the salinity, a gradual increase in the enriched α -amylase enzyme activity was observed with increasing NaCl concentrations up to $300~g\cdot L^{-1}$, as assayed at pH 9 and $80^{\circ}C$ (Figure 3). Although there was a sharp decrease in the enzyme activity above $300~g\cdot L^{-1}$ NaCl, the enzyme was still able to retain most (89 \pm 0.21%) of its initial activity at the nearly saturated salt concentration of $400~g\cdot L^{-1}$ NaCl.

Finally, the effect of various metal ions and enzyme inhibitors at a concentration of 2 mM on the activity of the enriched α -amylase is summarized in Table 2. Except for CaCl₂, which only weakly (but statistically significantly) increased the activity, none of the metal ions and enzyme inhibitors at this concentration were found to increase the enzyme activity. The slight decrease in the enzyme activity with the addition of 2 mM BaCl₂, HgCl₂, and *b*-mercaptoethanol was significant all the same, whilst the enzyme was moderately inhibited by the inclusion of FeCl₂ or EDTA and strongly inhibited by ZnCl₂.

3.3. Enzyme Kinetics. From the Lineweaver-Burk plot (Figure 4), the enriched amylase had a K_m of 5.41 mg·mL⁻¹ and

Properties	Cell-free supernatant	(NH ₄) ₂ SO ₄ precipitation	Gel filtration chromatography
Total protein (mg)	2301.1 ± 1.0	785.4 ± 1.1	219.4 ± 1.3
Total activity (U)	$39142 \pm 1.5 (100\%)$	$34015.8 \pm 1.0 \ (68\%)$	$25891.9 \pm 1.1 (44\%)$
Specific activity (U·mg ⁻¹)	17 ± 0.0	43.3 ± 0.1	118.4 ± 1.2
Purification fold	1.0	2.5	6.9

TABLE 1: Stepwise summary of the enrichment of the α -amylase from A. penicillioides TISTR 3639.

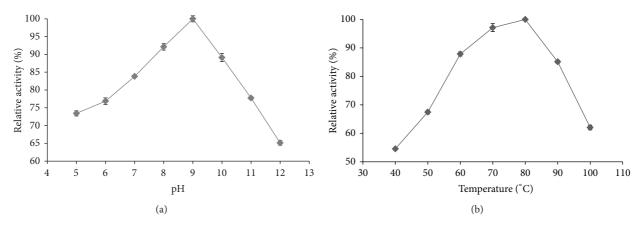


FIGURE 2: Effect of the (a) pH (at 30°C) and (b) temperature (at pH 9.0) on the activity of the enriched α -amylase from *A. penicillioides* TISTR 3639. Data are shown as the mean relative activity (%) ±1 SD (error bars), derived from three repeats. Means with a different lowercase superscript letter (a, b, and c) are significantly different (ANOVA and DMRT of the transformed data, P < 0.05).

Table 2: Effect of various metal ions and additives on the activity of the enriched α -amylase from A. penicillioides TISTR 3639.

Additives (2 mM)	Relative activity (%)
None	100*f
$BaCl_2$	97.8 ± 0.8^{e}
CaCl ₂	104.2 ± 1.7^{g}
$FeCl_2$	$78.1 \pm 1.5^{\circ}$
$HgCl_2$	97.1 ± 1.4^{e}
MgCl_2	$99.1 \pm 2.2^{\text{ef}}$
$ZnCl_2$	44.3 ± 0.9^{a}
β -Mercaptoethanol	95.1 ± 1.5^{d}
EDTA	73.4 ± 0.4^{b}

^{*} One hundred percent activity corresponded to the activity of the amylase without additive. Data are shown as the mean relative value \pm 1SD, derived from three repeats. Means followed by a different lowercase superscript letter (a, b, c, d, e, f, and g) are significantly different (ANOVA and DMRT of the transformed data, P < 0.05).

 $V_{\rm max}$ of 1.05 $\mu{\rm mol\cdot min}^{-1}{\cdot}{\rm mg}^{-1}$ with soluble starch as the substrate.

3.4. Compatibility of the Enriched α -Amylase with the Selected Three Commercial Detergents. The compatibility results of the enriched enzyme with three commercial detergents (A, B, and C) as well as the commercial amylase enzyme are summarized in Table 3. The residual activity taken in percentage, as compared with control, showed that the enzyme was seen to retain more than 80% of the activity without any added

Table 3: Effect of various detergents on the residual activity of the enriched α -amylase from *A. penicillioides* TISTR 3639.

Additive	Residual activity (%)
Distilled water (control)	100*d
Detergent A	81 ± 1.5^{a}
Detergent B	87 ± 0.1^{b}
Detergent C	94 ± 1.5^{c}

^{*}One hundred percent activity corresponded to the activity of the amylase without any added detergent. Data are shown as the mean relative value \pm ISD, derived from three repeats. Means followed by a different lowercase superscript letter (a, b, c, and d) are significantly different (ANOVA and DMRT of the transformed data, P < 0.05).

detergent. The powdered B and C detergents were found to be more compatible with the enriched amylase, by giving the residual enzyme activity of 87% and 94%, respectively, than was liquid detergent A, which was observed to provide 81% amylase residual activity.

3.5. Performance Test with Commercial Detergents. The performance test results of this enriched α -amylase in the presence of the respective commercial detergents are summarized in Figure 5, where the relative activity (%) was found to decrease with increasing salt concentration. However, in similar salt conditions the commercial amylase and the detergents had an inferior activity in comparison to this enriched α -amylase from A. penicillioides. At 50 g·L $^{-1}$ NaCl, only 60% of the A. penicillioides amylase activity was detected in the commercial amylase and 60–70% in the three detergents.

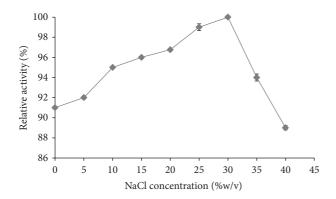


FIGURE 3: Effect of the NaCl concentration (g·L⁻¹) on the activity of the enriched α -amylase from *A. penicillioides* TISTR 3639 at pH 9 and 80°C. Data are shown as the mean relative activity (%) ±1 SD (error bars), derived from three repeats. Means with a different lowercase superscript letter (a, b, and c) are significantly different (ANOVA and DMRT of the transformed data, P < 0.05).

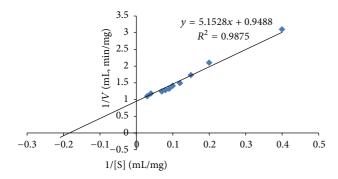


FIGURE 4: Lineweaver-Burk plot for the determination of the $V_{\rm max}$ and K_m values of the enriched α -amylase from A. penicillioides TISTR 3639, at optimum conditions (pH 9 and 80°C), in the presence of different concentrations of soluble starch.

4. Discussion

Many attempts have been made to find suitable fungus strains for the production of amylases with desirable properties [21]. Mesophilic fungi are reported to be the principal amylase producers [16] and especially members of the Aspergillus and Penicillium genera that appear to be the dominant producing species [22]. Fungal amylases are preferred for use in various industries, including the food and pharmaceutical industries, due to their nontoxic characteristics [3, 12, 23]. Consequentially, Aspergillus species, such as Aspergillus niger and Aspergillus oryzae, are frequently used in the industrial production of amylases [24], but there are few reports on the purification and detailed characterization of α -amylases from halophilic fungi [12]. In addition, to the best of our knowledge, this is the first report of the determination of the potential of an amylase from any obligate halophilic fungus to be used as an additive in laundry detergent.

The molecular mass of amylases from halophilic fungi is mostly reported in the range of 50–75 kDa [25]. In this study for *A. penicillioides* TISTR 3639, it was found slightly smaller with an approximate mass of 42 kDa (Figure 1). Moreover, the

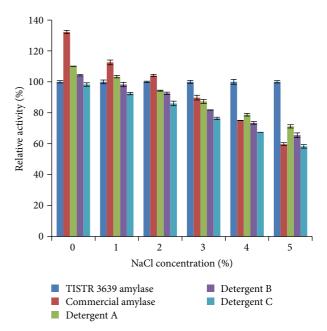


FIGURE 5: Performance comparison of a commercial amylase, detergents A, B, and C, and the enriched α -amylase (control) from *A. penicillioides* TISTR 3639 at pH 7 and 30°C in the presence of different concentrations of NaCl. The results are expressed as the relative activity (%) to that of the enriched α -amylase from *A. penicillioides* TISTR 3639 without NaCl and are shown as the mean relative activity (%) ±1 SD (error bars), derived from three repeats. Means with a different lowercase superscript letter (a, b, and c) are significantly different (ANOVA and DMRT of the transformed data, P < 0.05).

specific activity of the amylase was in the moderate range [17], whilst the K_m value (5.41 mg·mL $^{-1}$) of the enzyme falls in the middle of the range (0.35–11.66 mg·mL $^{-1}$) reported for amylases from halophilic fungi [26]. The K_m and $V_{\rm max}$ values of the enriched α -amylase of this study (with soluble starch as the substrate) suggested that it has a moderately high affinity for soluble starch and requires a relatively low concentration of this substrate to achieve $V_{\rm max}$.

The optimal pH profile of this enriched amylase mirrors the pH-dependent growth profile of its producing *A. penicillioides*, where it was previously reported that *A. penicillioides* grew relatively well at a neutral to alkaline pH [1]. This also matches the pH found in most of the hypersaline environments [6]. The enzyme in this study retained almost 80% of its relative activity at an extreme pH (pH 11), but at pH 5 it was slightly lower at less than 75% relative activity, showing that it is potentially more tolerant of alkaline pH. An optimal pH of 9-10 has been reported for many enzymes from halophilic organisms and an alkalophilic property has been considered as the nature of halophilic enzymes [27, 28].

The alkaline amylases from microorganisms have been found to have tremendous applications in detergent industries [15]. However, most halophilic enzymes are denatured and lose their activity at temperatures over 50°C [29], whereas this halophilic and alkalophilic amylase from *A. penicillioides* TISTR 3639 showed an optimal activity at 80°C and was

still relatively stable at 90°C or lower with more than 80% of its initial activity remaining after a 1 h incubation at this temperature (data not shown). Thus, it is extremely thermophilic [30, 31]. Thermophilic amylases are mostly applied in various starch industries [29].

Salinity is a crucial factor in the normal functioning of most enzymes from obligate halophilic microorganisms. Previously, it was found that *A. penicillioides* grew best at a salinity of $100\,\mathrm{g}\cdot\mathrm{L}^{-1}$ NaCl [1], a salinity level that has been widely reported to favor amylase production in most halophilic microbes [12, 32, 33]. However, the α -amylase from *A. penicillioides* TISTR 3639 still had a high catalytic activity even at extreme salt concentrations (300–400 g/mL⁻¹ NaCl), much higher than those of the amylases reported from other extreme halophilic Archaea and Bacteria [10, 33, 34], adding a novelty to this enzyme. This extremophile property of the enzyme suggests the potential to be used in saline waste water management, in bioremediation processes in saline areas [12, 35, 36], and for biofuel production, where halophilic enzymes are reported to work better than normal enzymes [37].

The amylases from halophilic microorganisms have frequently been reported to have polyextremophilic characteristics [10, 12, 33]. They are mostly reported to be thermotolerant with haloalkalophilic properties [25], where the enzyme must have the capability to withstand extreme conditions for several industrial processes [38]. The amylase from *A. penicillioides* TISTR 3639 has the same trend of being polyextremophile, which makes it potentially versatile for use in different industrial operations, where harsh conditions are available.

The detergent industries are one of the primary consumers and users of enzymes that enhance the performance of detergents or allow the product to be more environmentally friendly [21]. Approximately 90% of commercial liquid detergents contain a mixture of enzymes that include amylases [39], since starch is considered as an attractant for various soil particles [18]. Amylases are also used in the detergent industries to remove starchy food stains, such as chocolate, custard, gravy, and potato amongst others, which are found on kitchen utensils as well as on clothes [16]. The addition of any enzyme in the detergent requires its compatibility and ability to perform in the presence of the detergents for inclusion [40]. Currently, there have been only a few reports of suitable and stable amylases that can be added into laundry detergents [41], and these need to work well at an alkaline pH [21]. Thus, it is of interest that the enriched amylase from A. penicillioides TISTR 3639 was found to be relatively stable in the presence of different detergents as well as alkaline conditions. Note that the compatibility test was performed under normal conditions (30°C, pH 7) that are suboptimal for this enzyme and so its activity may be improved at higher temperatures, alkalinity, and salinity levels, although of course its compatibility with these detergents would need to be evaluated at these conditions.

The rapidly changing world provides some tough challenges to humans. The limiting water resources in many parts of the world compel the use of saline or hard untreated water for daily domestic uses [42], including untreated underground hard water [43], and these decrease the cleaning

efficiency of detergents [44]. In this study, the enriched α -amylase from A. penicillioides TISTR 3639, in comparison with three tested detergents and a commercial amylase, was found to work well in a low water activity, which means that the inclusion of this amylase could help solve the problem of a low cleaning capacity of detergents in a high saline environment. Moreover, when such underground hard water is not suitable for drinking it can be used for laundry purposes, saving the clean and drinkable water for consumption in areas where drinking water resources is limited.

5. Conclusions

Increasing demands from biotechnology, climate change, and decreasing water resources form the need to find amylases that can withstand high temperatures, salt concentrations (low water activity), and alkalinity levels. The polyextremophilic behavior of this enriched α -amylase obtained from *A. penicillioides* TISTR 3639 appears to be a promising candidate for fulfilling the current needs of many industrial processes requiring amylases, especially for laundry detergent industries. Of interest was the fact that the α -amylase from the obligate halophilic *A. penicillioides* TISTR 3639 was more extremophilic than the fungus itself. This provides an opportunity to exploit more interesting biotechnological applications from obligate halophilic fungi.

Conflict of Interests

The authors declare that they have no conflict of interests.

Acknowledgments

The authors would like to thank all who helped in their research. This work was financially supported by a Post-Doctoral Research Grant Allocation from the Ratchadaphiseksomphot Fund, the Ratchadaphiseksomphot Endowment Fund 2014 of Chulalongkorn University (CU-57-043-EN) and Research Center for Thermotolerant Microbial Resources (RCTMR) in Yamaguchi University.

References

- [1] I. Ali, L. Kanhayuwa, S. Rachdawong, and S. K. Rakshit, "Identification, phylogenetic analysis and characterization of obligate halophilic fungi isolated from a man-made solar saltern in Phetchaburi province, Thailand," *Annals of Microbiology*, vol. 63, no. 3, pp. 887–895, 2013.
- [2] N. Gunde-Cimerman, J. C. Frisvad, and P. Zalaretal, "Halotolerant and halophilic fungi," in *Biodiversity of Fungi-Their Role in Human Life*, S. K. Deshmukh and M. K. Rai, Eds., pp. 96–128, Oxford & IBH, New Delhi, India, 2005.
- [3] I. Ali, N. Siwarungson, H. Punnapayak et al., "Screening of potential biotechnological applications from obligate halophilic fungi, isolated from a man-made solar saltern located in Phetchaburi province, Thailand," *Pakistan Journal of Botany*, vol. 46, pp. 983–988, 2014.
- [4] I. Ali, S. K. Rakshit, A. Akbar et al., "Identification and phylogenetic analysis of halophilic fungus isolated from a man-made

- solar saltern in Thailand," *Lasbella University Journal of Science and Technology*, vol. 2, pp. 47–52, 2013.
- [5] G. Antranikian, "Extreme environments as a resource for microorganisms and novel biocatalysts," *Marine Biotechnology*, vol. 96, pp. 219–262, 2005.
- [6] C. Gostinčar, M. Lenassi, N. Gunde-Cimerman, and A. Plemenitaš, "Fungal adaptation to extremely high salt concentrations," *Advances in Applied Microbiology*, vol. 77, pp. 71–96, 2011.
- [7] A. Oren, "Industrial and environmental applications of halophilic microorganisms," *Environmental Technology*, vol. 31, no. 8-9, pp. 825–834, 2010.
- [8] S. DasSarma and P. DasSarma, *Halophiles*, Encyclopedia of Life Sciences, Wiley, London, UK, 2002.
- [9] M. Delgado-García, B. Valdivia-Urdiales, C. N. Aguilar-González, J. C. Contreras-Esquivel, and R. Rodríguez-Herrera, "Halophilic hydrolases as a new tool for the biotechnological industries," *Journal of the Science of Food and Agriculture*, vol. 92, no. 13, pp. 2575–2580, 2012.
- [10] M. Moshfegh, A. R. Shahverdi, G. Zarrini, and M. A. Faramarzi, "Biochemical characterization of an extracellular polyextremophilic α-amylase from the halophilic archaeon *Halorubrum* xinjiangense," Extremophiles, vol. 17, no. 4, pp. 677–687, 2013.
- [11] H. Dalbøge, "Expression cloning of fungal enzyme genes: a novel approach for efficient isolation of enzyme genes of industrial relevance," FEMS Microbiology Reviews, vol. 21, no. 1, pp. 29–42, 1997.
- [12] I. Ali, A. Akbar, B. Yanwisetpakdee, S. Prasongsuk, P. Lotrakul, and H. Punnapayak, "Purification, characterization, and potential of saline waste water remediation of a polyextremophilic α-amylase from an obligate halophilic Aspergillus gracilis," BioMed Research International, vol. 2014, Article ID 106937, 7 pages, 2014.
- [13] X. D. Liu and Y. Xu, "A novel raw starch digesting α -amylase from a newly isolated *Bacillus* sp. YX-1: purification and characterization," *Bioresource Technology*, vol. 99, no. 10, pp. 4315–4320, 2008.
- [14] M. Shafiei, A.-A. Ziaee, and M. A. Amoozegar, "Purification and biochemical characterization of a novel SDS and surfactant stable, raw starch digesting, and halophilic α-amylase from a moderately halophilic bacterium, *Nesterenkonia* sp. strain F," *Process Biochemistry*, vol. 45, no. 5, pp. 694–699, 2010.
- [15] K. K. Kiran and T. S. Chandra, "Production of surfactant and detergent-stable, halophilic, and alkalitolerant alpha-amylase by a moderately halophilic *Bacillus* sp. Strain TSCVKK," *Applied Microbiology and Biotechnology*, vol. 77, no. 5, pp. 1023–1031, 2008.
- [16] R. Gupta, P. Gigras, H. Mohapatra, V. K. Goswami, and B. Chauhan, "Microbial α-amylases: a biotechnological perspective," *Process Biochemistry*, vol. 38, no. 11, pp. 1599–1616, 2003.
- [17] S. Chakraborty, A. Khopade, C. Kokare, K. Mahadik, and B. Chopade, "Isolation and characterization of novel α-amylase from marine *Streptomyces* sp. D1," *Journal of Molecular Catalysis B: Enzymatic*, vol. 58, no. 1–4, pp. 17–23, 2009.
- [18] N. Hmidet, A. Bayoudh, J. G. Berrin, S. Kanoun, N. Juge, and M. Nasri, "Purification and biochemical characterization of a novel α-amylase from *Bacillus licheniformis* NH1. Cloning, nucleotide sequence and expression of *amy* N gene in *Escherichia coli*," *Process Biochemistry*, vol. 43, no. 5, pp. 499–510, 2008.
- [19] G. L. Miller, "Use of dinitrosalicylic acid reagent for determination of reducing sugar," *Analytical Chemistry*, vol. 31, no. 3, pp. 426–428, 1959.

- [20] O. H. Lowry, N. J. Rosebrough, A. L. Farr, and R. J. Randall, "Protein measurement with the Folin phenol reagent," *The Journal of Biology and Chemistry*, vol. 193, no. 1, pp. 265–275, 1951.
- [21] P. M. de Souza and P. D. O. Magalhães, "Application of microbial α-amylase in industry—a review," *Brazilian Journal of Microbiology*, vol. 41, no. 4, pp. 850–861, 2010.
- [22] K. Kathiresan and S. Manivannan, "α-Amylase production by *Penicillium fellutanum* isolated from mangrove rhizosphere soil," *African Journal of Biotechnology*, vol. 5, no. 10, pp. 829–832, 2006.
- [23] F. Niknejad, M. Moshfegh, M. J. Najafzadeh et al., "Halotolerant ability and α-amylase activity of some saltwater fungal isolates," *Iranian Journal of Pharmaceutical Research*, vol. 12, pp. 113–119, 2013
- [24] R. Kammoun, B. Naili, and S. Bejar, "Application of a statistical design to the optimization of parameters and culture medium for α -amylase production by *Aspergillus oryzae* CBS 819.72 grown on gruel (wheat grinding by-product)," *Bioresource Technology*, vol. 99, no. 13, pp. 5602–5609, 2008.
- [25] M. E. Setati, "Diversity and industrial potential of hydrolaseproducing halophilic/halotolerant eubacteria," *African Journal* of *Biotechnology*, vol. 9, no. 11, pp. 1555–1560, 2010.
- [26] M. F. Najafi and A. Kembhavi, "One step purification and characterization of an extracellular α -amylase from marine *Vibrio* sp.," *Enzyme and Microbial Technology*, vol. 36, no. 4, pp. 535–539, 2005.
- [27] B. Prakash, M. Vidyasagar, M. S. Madhukumar, G. Muralikrishna, and K. Sreeramulu, "Production, purification, and characterization of two extremely halotolerant, thermostable, and alkali-stable α -amylases from *Chromohalobacter* sp. TVSP 101," *Process Biochemistry*, vol. 44, no. 2, pp. 210–215, 2009.
- [28] K. N. Varalakshmi, B. S. Kumudini, B. N. Nandini et al., "Production and characterization of α-amylase from Aspergillus niger JGI 24 isolated in Bangalore," Polish Journal of Microbiology, vol. 58, no. 1, pp. 29–36, 2009.
- [29] O. Prakash and N. Jaiswal, "α-Amylase: an ideal representative of thermostable enzymes," *Applied Biochemistry and Biotechnol*ogy, vol. 160, no. 8, pp. 2401–2414, 2010.
- [30] S. A. Fincan and B. Enez, "Production, purification, and characterization of thermostable a-amylase from thermophilic *Geobacillus stearothermophilus*," *Starch/Staerke*, vol. 66, no. 1-2, pp. 182–189, 2014.
- [31] I. Gomes, J. Gomes, and W. Steiner, "Highly thermostable amylase and pullulanase of the extreme thermophilic eubacterium *Rhodothermus marinus*: production and partial characterization," *Bioresource Technology*, vol. 90, no. 2, pp. 207–214, 2003.
- [32] M. A. Amoozegar, F. Malekzadeh, and K. A. Malik, "Production of amylase by newly isolated moderate halophile, *Halobacillus* sp. strain MA-2," *Journal of Microbiological Methods*, vol. 52, no. 3, pp. 353–359, 2003.
- [33] K. K. Kondepudi and T. S. Chandra, "Production of surfactant and detergent-stable, halophilic, and alkalitolerant alpha-amylase by a moderately halophilic *Bacillus* sp. strain TSCVKK," *Applied Microbiology and Biotechnology*, vol. 77, no. 5, pp. 1023–1031, 2008.
- [34] M. Shafiei, A.-A. Ziaee, and M. A. Amoozegar, "Purification and characterization of a halophilic α-amylase with increased activity in the presence of organic solvents from the moderately halophilic *Nesterenkonia* sp. strain F," *Extremophiles*, vol. 16, no. 4, pp. 627–635, 2012.

- [35] J. Chung, S. Shin, and J. Oh, "Characterization of a microbial community capable of reducing perchlorate and nitrate in high salinity," *Biotechnology Letters*, vol. 31, no. 7, pp. 959–966, 2009.
- [36] R. Margesin and F. Schinner, "Biodegradation and bioremediation of hydrocarbons in extreme environments," Applied Microbiology and Biotechnology, vol. 56, no. 5-6, pp. 650–663, 2001.
- [37] T. Zhang, S. Datta, J. Eichler et al., "Identification of a haloalkaliphilic and thermostable cellulase with improved ionic liquid tolerance," *Green Chemistry*, vol. 13, no. 8, pp. 2083–2090, 2011.
- [38] W. Asad, M. Asif, and S. A. Rasool, "Extracellular enzyme production by indigenous thermophilic bacteria: partial purification and characterization of α-amylase by *Bacillus* sp. WA21," *Pakistan Journal of Botany*, vol. 43, no. 2, pp. 1045–1052, 2011.
- [39] S. Mitidieri, A. H. S. Martinelli, A. Schrank, and M. H. Vainstein, "Enzymatic detergent formulation containing amylase from Aspergillus niger: a comparative study with commercial detergent formulations," Bioresource Technology, vol. 97, no. 10, pp. 1217–1224, 2006.
- [40] A. K. Mukherjee, H. Adhikari, and S. K. Rai, "Production of alkaline protease by a thermophilic *Bacillus subtilis* under solidstate fermentation (SSF) condition using *Imperata cylindrica* grass and potato peel as low-cost medium: characterization and application of enzyme in detergent formulation," *Biochemical Engineering Journal*, vol. 39, no. 2, pp. 353–361, 2008.
- [41] A. K. Mukherjee, M. Borah, and S. K. Rai, "To study the influence of different components of fermentable substrates on induction of extracellular α-amylase synthesis by *Bacillus subtilis* DM-03 in solid-state fermentation and exploration of feasibility for inclusion of α-amylase in laundry detergent formulations," *Biochemical Engineering Journal*, vol. 43, no. 2, pp. 149– 156, 2009.
- [42] I. Ali, A. Akbar, H. Punnapayak et al., "Seven big challenges for Pakistan—and the lessons they could teach," *The Futurist*, vol. 48, pp. 22–26, 2014.
- [43] S. Ghizellaoui and A. Chibani, "Use of nanofiltration for partial softening of very hard water," *Desalination*, vol. 179, no. 1–3, pp. 315–322, 2005.
- [44] J.-S. Park, J.-H. Song, K.-H. Yeon, and S.-H. Moon, "Removal of hardness ions from tap water using electromembrane processes," *Desalination*, vol. 202, no. 1–3, pp. 1–8, 2007.