

Production of an alkaline protease by *Bacillus cereus* MCM B-326 and its application as a dehairing agent

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Abstract The present investigation describes microbial production of an alkaline protease and its use in dehairing of buffalo hide. *Bacillus cereus* produced extracellular protease when grown on a medium containing starch, wheat bran and soya flour (SWS). The ammonium sulphate precipitated (ASP) enzyme was applied for dehairing of buffalo hide. Microscopic observation of longitudinal section of buffalo hide revealed that the epidermis was completely removed and hair was uprooted leaving empty follicles in the hide. The ASP enzyme was stable for one month at ambient temperature between 25–35 °C. Enzymatic dehairing may be a promising shift towards an environment-friendly leather processing method.

Keywords *Bacillus cereus* · Alkaline protease · Dehairing · Buffalo hide · Leather processing

Introduction

Currently, over 30% of the world industrial enzyme market consists of alkaline proteases used in detergents, food processing and leather industry. Production of these enzymes using low cost substrates would reduce the cost of production. Oil seed cakes, which are a by product of oil extraction industry, are a potentially useful, low-cost substrate for the production of different enzymes (Kanekar et al. 1997; Tang et al. 2000; Alagarsamy et al. 2005). Keratinase, a type of bacterial protease, was produced by using cheap substrates such as hair, feathers and whey

milk—a dairy byproduct (Alessandro et al. 2003; Roberta et al. 2006; Philippe et al. 1999; Prakasham et al. 2006). Yang et al. (2000) have reported deproteinization of crustacean waste by protease of *B. subtilis*.

Leather industry waste leads to environmental pollution and health hazards (Malathi and Chakraborty 1991). Major pollutants from the leather industry that may have significant environmental impact include lime, sulphide and chromium (Alessandro et al. 2003). Huge amounts of lime sludge and total solids formation are the main drawbacks of lime (Thanikaivelan et al. 2003). Untreated sulphide can cause major problem in sewers. The severe alkaline conditions constitute a health hazard for the worker (Purushotham et al. 1996). Sulphide also reduces the strength of hair, which directly hampers the recovery of this value-added byproduct. The tannery effluent has high dissolved solids (TDS), high biological oxygen demand (BOD) and high chemical oxygen demand (COD) (Palanisamy et al. 2004). Enzymatic dehairing is being increasingly looked upon as a reliable alternative to the conventional lime-sulphide process, avoiding the problems created by sulphide. The advantages of enzymatic dehairing are (i) total elimination of lime and sulphide from the effluent, (ii) recovery of hair as a byproduct which may be used for production of synthetic fibres, biogas, foaming agent for fire extinguishers; while hydrolyzed hair is used as agricultural fertilizer, soil conditioner, compost, additive in chrome tanning or retanning processes, animal/poultry feed and also for production of cosmetics, pharmaceuticals and amino acid like cysteine (Palanisamy et al. 2004), (iii) elimination of the bating process required during deliming. However, these benefits remain unfulfilled, since enzymes are more expensive than the conventional chemical process, and require careful control (Kanagaraj et al. 2006; Qing et al. 2003). The latter point has been identified as the main obstacle in the wider application of enzymes. Pal et al.

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(1996) reported an enzyme-assisted dehairing with lime and protease from *R. oryzae*, where lime acts as a swelling agent. The potential for commercial use of enzymes in leather production is considerable, because of their properties of being highly efficient and substrate specific. Thus, the substitution of chemical dehairing agents in the leather industry by proteolytic enzymes produced by microorganisms will have an important economic and environmental impact.

Earlier studies using commercial enzymes for sheep skin dehairing revealed a high correlation between dehairing activity and proteolytic activity (Qing et al. 2003). Some microorganisms producing extracellular enzyme with dehairing activity have been described, e.g., *Streptomyces* sp. isolated from soil, degraded human hair, chicken feather, silk, wool, and unhaired goatskin (Mukhopadhyay and Chandra 1993). Among bacteria, strains of *B. subtilis* and *B. amyloliquefaciens* have been characterized for dehairing purposes (Varela et al. 1997; Nashy et al. 2005). However, many proteases are not suitable for dehairing, since they have collagen-degrading activity, which destroys the collagen structure of the hide. Therefore, it is essential to explore proteases with dehairing activity but no collagenolytic activity (Qing et al. 2003).

The present paper describes the production of an alkaline protease by *B. cereus* using inexpensive and easily available protein rich substrates, and its application in dehairing of buffalo hide.

Materials and methods

Microorganism

The bacterium used in this study was *B. cereus* MCM B-326, a buffalo hide isolate, deposited in MACS Collection for Microorganisms (MCM). The stock culture was maintained in 15% glycerol at -20°C .

Substrates

Different industrial deoiled cakes like soya cake, saffola cake, rape seed cake, groundnut cake, wheat bran and flours like soya flour, bengal gram flour were purchased from local market. Soybean meal was procured from Hi-Media Laboratories, Mumbai.

Production of enzyme

The enzyme was produced in 250 mL capacity Erlenmeyer flask using 100 mL medium, containing 1% starch and 0.3% CaCO_3 along with one of the following different cheap nitrogen sources: deoiled soyabean cake, saffola cake, rape seed cake, groundnut cake, soya flour, wheat bran, bengal

gram flour, bengal gram flour combined with wheat bran, bengal gram flour combined with deoiled groundnut cake, bengal gram flour combined with soybean meal, wheat bran combined with deoiled ground nut cake, wheat bran combined with soya flour, and soybean meal, in the medium at 1% level. The media at pH 9.0 were inoculated with 1% of 21 h old inoculum (2.8×10^8 cells mL^{-1}) and incubated at 30°C for 36 h under shake culture conditions. The broth was centrifuged at 13,000g for 10 min to obtain cell free supernatant (CFS). The CFS was then partially purified by 60% saturation of ammonium sulphate. Ammonium sulphate precipitated (ASP) enzyme was applied for dehairing of buffalo hide.

Partial purification and zymogram

The proteins from CFS were precipitated with ammonium sulphate (60% saturation) followed by membrane dialysis. The enzyme pattern of precipitated proteins was obtained by zymogram with 1% casein substrate and detected using comassive brilliant blue R250. The molecular weight of the protease was determined by comparing with mobility of standard molecular weight marker proteins (bovine albumin 66 kDa, chicken ovalbumin 45 kDa, glyceroldehyde-3-phosphate dehydrogenase 36 kDa, trypsinogen 24 kDa, cytochrome C 12.4 kDa).

Protease assay

The protease activity was determined by caseinolytic assay (Nilegaonkar et al. 2000). One mL of the cell free supernatant was mixed with 4 mL of casein (0.625% w/v) and incubated at 37°C for 30 min. The reaction was stopped by addition of 5 mL of 5% trichloroacetic acid (TCA). Enzymatically hydrolyzed casein was measured by modified Folin–Ciocalteu method (Jayaraman 2003) against casein treated with inactive enzyme as blank. The TCA precipitated reaction mixture was centrifuged and 0.1 mL of supernatant was mixed with 0.9 mL of distilled water and 0.5 mL of alkaline copper tartarate solution, and incubated for 10 min. Two mL of 1N Folin–Ciocalteu reagent was added and the absorbance read at 660 nm after 20 min of incubation. A standard graph was generated using standard tyrosine solutions of 5–50 $\mu\text{g mL}^{-1}$. All the incubations were carried out at room temperature. One unit of protease activity was defined as the amount of enzyme, which liberated 1 μg tyrosine per min at 37°C .

Protein estimation

The protein content of the cell free supernatant was estimated using Biuret method, with bovine serum albumin as a standard (Jayaraman 2003).

Enzyme storage stability

The storage stability of the crude enzyme was assessed by storing the enzyme at room temperature (28 ± 2 °C) and estimating the activity at different time intervals up to 90 days. This was done by conducting caseinolytic assay.

Dehairing activity

The salted buffalo hide pieces of size 2.5 cm² or weight 3 g were soaked overnight in tap water (400% v/w of salted hide) with 0.3% calcium carbonate. The soaked hide pieces were incubated with the enzyme (1% w/w of hide) in tap water having pH 7.0 at room temperature (28 ± 2 °C) along with control i.e., hide piece incubated without enzyme, under the same conditions. For comparison, chemical dehairing was also carried out with 10% lime and 2% sodium sulphide. After 21 h, pieces were dehaired by mechanical means and visually analyzed for color, smoothness and silkiness of the pelt.

Histological study

The pelts were prepared for histopathological analysis by fixing in 10% (w/v) formaldehyde. The pelt was dehydrated with 80, 95 and 100% (v/v) of alcohol gradients followed by xylene treatment, and then embedded in paraffin. Longitudinal sections (L.S.) of hide embedded in paraffin wax were obtained using a microtome. The sections were fixed on slides using starch paste containing thymol, which acts as a preservative. The sections were stained with Harris's haematoxylin stain followed by 0.5% (v/v) HCl and dilute ammonia (John and Merrilline 2002). The slides were observed microscopically (Labophot 2 microscope, Japan) for (a) epidermis, (b) hair root follicles, and (c) corium (collagen layer).

Statistical analysis

Analysis of variance with repeated measures was carried out using GLM command of software SPSS (SPSS version 10, Windows 98). Activity means and standard deviation were calculated (Table 1). Univariate analysis of variance (ANOVA) was applied to the data on supplementation of different cheap nitrogen sources and protease activity.

Results and discussion

Protease production

Enzyme, as a biological catalyst, depends upon various factors for its activity. Most of the enzymes used for the

Table 1 Protease production by *B. cereus* MCM B-326 using different cheap nitrogen sources supplements

Cheap sources*	Enzyme activity**(U mL ⁻¹)
Deoiled soybean cake	95.85 ± 1.36
Deoiled saffola cake	72.09 ± 4.07
Deoiled rape seed cake	71.28 ± 3.80
Deoiled groundnut cake	119.54 ± 2.32
Wheat bran	75.54 ± 2.05
Soya flour	81.32 ± 2.73
Bengal gram flour	38.21 ± 2.25
Bengal gram flour + wheat bran	160.80 ± 6.11
Bengal gram flour + groundnut cake	118.41 ± 5.00
Bengal gram flour + soybean meal	90.23 ± 4.06
Wheat bran + groundnut cake	142.89 ± 3.07
Wheat bran + soya flour	176.05 ± 4.24
Soybean meal (Himedia)	140.73 ± 4.83

* Cheap sources with 1% starch and 0.3% CaCO₃, pH 9.0 at 30 °C, shaking culture condition

** The results given are mean values of six observations ± standard deviation (SD)

Data was significant ($P < 0.01$)

dehairing process have been obtained from microorganisms isolated from soil adjacent to tanneries, and decomposing skins (George et al. 1995; Dayanandan et al. 2003).

Earlier, *B. cereus* MCM B-326 was grown in starch-soybean meal and calcium carbonate medium revealing an alkaline metalloprotease, non-collagenolytic and non-keratinolytic in nature. Ammonium sulphate precipitated enzyme showed two bands, viz., Cereus 1 and Cereus 2, with approximate molecular weights 45-kDa and 36-kD on non-denaturing PAGE (Fig. 1). It is reported that *B. cereus* has two proteases with molecular masses of approximately 38-kDa and 36-kDa (Kim et al. 2001). A calcium dependent protease from *B. cereus* BG1 was reported to have molecular weight of 34-kDa (Ghorbel et al. 2005). According to Qing et al. (2003) the molecular weight of the purified dehairing protease from *B. pumilus* was 32-kDa. Thus molecular weight of Cereus 2 is similar to the molecular weight reported for other proteases from *B. cereus*, while molecular weight of Cereus 1 is different (Nilegaonkar et al. 2007).

The same organism is studied for production of protease with inexpensive substrates and its application in dehairing of buffalo hide. *Bacillus cereus* MCM B-326 has maximum protease production of 176.05 ± 4.24 U mL⁻¹ in medium containing starch, wheat bran, soya flour (1% each) and 0.3% CaCO₃, at pH 9.0 within 36 h, under shake culture condition (Table 1). Wheat bran can be used in combination with other substrates like bengal gram flour or soya flour or soybean meal or groundnut cake for the production

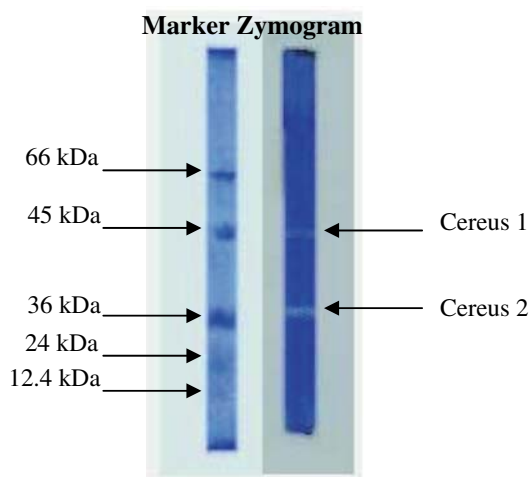
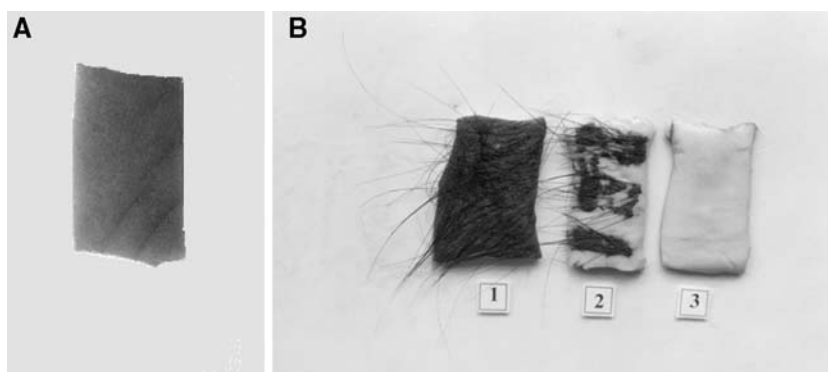


Fig. 1 Zymogram of protease from *Bacillus cereus* MCM B-326

of protease. In our previous study, it was observed that *B. cereus* shows protease activity of $126.87 \pm 1.32 \text{ U mL}^{-1}$ in starch, soybean meal and calcium carbonate medium under identical experimental conditions (Nilegaonkar et al. 2007). Earlier, protease with $3541.66 \text{ U L}^{-1} \text{ h}^{-1}$ productivity using nug-meal—a byproduct of oil extraction from seeds of *Guizotia abyssinica*, using *Bacillus* sp. AR-009 was reported (Amare 1997). Sweet potato residue and peanut meal residue were also reported as substrates for the production of protease (17.5–26.7 U) at 200–250 h using *Streptomyces rimosus* TM-55 in submerged and solid state cultivations method (Yang and Wang 1999). As reported here, the protease from *B. cereus* MCM B-326 has similar activity to that reported on nug meal substrates, and higher activity than that produced using sweet potato residue and peanut residue. *B. licheniformis*, when grown at 50 °C on oil seed cakes of linseed, castor and groundnut, yielded low levels of enzyme activities, against a combination of soybean meal and wheat bran (Sinha and Satyanarayana 1991). *Bacillus* sp. JB 99 when grown in solid state on wheat bran and rice bran exhibited enzyme activities of 7836 and 7540 U mL^{-1} respectively (Johnvesly et al. 2002).

Fig. 2 (A) Chemically dehaired buffalo hide. (B) Enzymatically dehaired buffalo hide (1) control-hide piece incubated without enzyme, in tap water, (2) enzymatically dehaired pelt after 12 h, (3) enzymatically dehaired pelt after 21 h



Storage stability

The ASP enzyme retained 80, 70 and 60% activity after one, two and three months respectively, when stored at room temperature. Our laboratory data (Nilegaonkar et al. 2007) suggests that Ca^{+2} is not necessary for the activity/stability of the enzyme, but enzymes reported and used in leather processing were stabilized by addition of calcium or co-polymers (Jaquess et al. 1999). In *Streptomyces moderatus* NRRL 3150, the enzyme activity of multiple proteinase concentrate was found to be stable with calcium chloride and sodium azide up to six months at 4 °C (Chandrasekaran and Dhar 1985).

Enzymatic dehairing of buffalo hide

Enzyme precipitated with ammonium sulphate (60% saturation) was used for dehairing of buffalo hide. In our previous studies, the enzyme was shown to be an alkaline metalloprotease, non-collagenolytic and non-keratinolytic in nature (Nilegaonkar et al. 2007). The presence of lime or sulphide was not required for this process. The enzyme-treated hide showed visible dehairing activity after 12 h of incubation and complete dehairing at 21 h. In the control sample, hair loosening was not observed, even by mechanical means such as plucking by forceps (Fig. 2).

In the chemically treated pelt, epidermis was removed (Fig. 3a). Microscopic examination of the pelt treated with enzyme showed complete removal of epidermis and presence of empty hair follicles (Fig. 3b) suggesting removal of hair from hair root follicle. Collagen layer was not damaged or modified in both chemical and enzyme-treated pelts. Microscopic examination of the pelt after lime-sulphide dehairing showed presence of parts of disintegrated shaft and entire bulb in the follicles. Similar results were observed by using multiple proteinase concentrate in goat skin (Chandrasekaran and Dhar 1985). In our studies, it was also observed that protease treated pelt swells moderately with adequate opening up of collagen fibre. Results

Fig. 3 Microscopic examination of buffalo hides. L.S. of (a) chemically dehaired pelt, (b) enzymatically dehaired pelt. Epidermis (→), Hair follicle (—●), Empty follicle (.....▶), Corium (C)

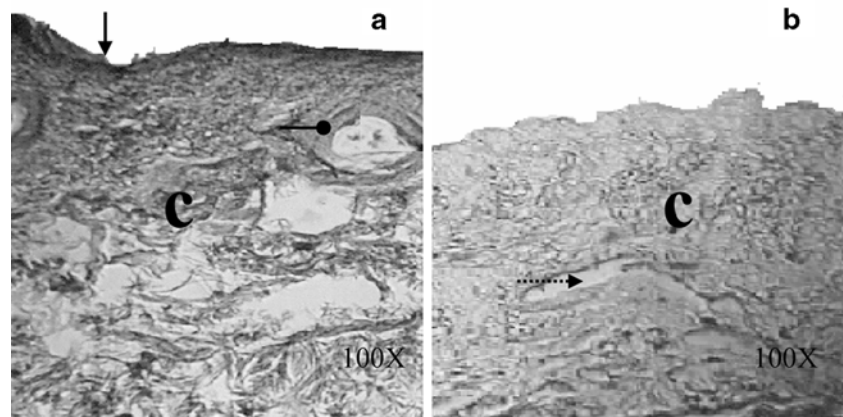
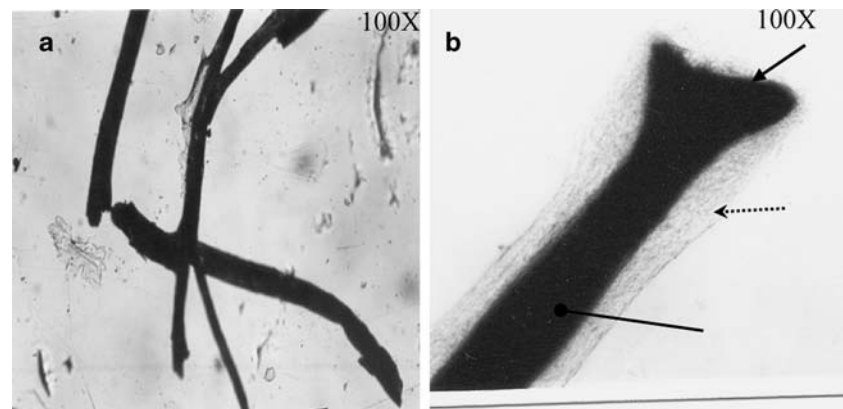


Fig. 4 Microscopy of hair, released in dehairing. (a) Chemical treated: broken hair; (b) Enzyme treated: intact hair. Hair root (←), Root sheath (←.....), Hair shaft (●—)



showed that for *Rhizopus oryzae*, the protease required liming treatment for desired swelling (Pal et al. 1996).

Enzymatic dehairing did not damage the collagen layer. Likewise, a novel keratinase obtained from *B. subtilis* S14 exhibited remarkable dehairing capabilities without damage to bovine skin collagen, which was confirmed on SDS-PAGE using control collagen, and collagen treated with standard keratinase, collagenase and test enzymes (Alexandre et al. 2005).

When hair is removed from buffalo hide by chemical method, it was observed microscopically that the hair was distorted or broken into small pieces (Fig. 4a) while enzyme-treated hair was intact, with hair root, shaft, and sheath (Fig. 4b). In previous reports, the epidermis, hair shaft and bulbs were completely removed with better splitting of fibre bundles by the action of multiple protease concentrate (Chandrasekaran and Dhar 1985). Our study indicates that the hair loosening was due to enzyme action and not due to the swelling up or modification of the sheath.

Conclusion

Production of an efficient dehairing enzyme from *B. cereus* MCM B-326 with inexpensive and easily available

substrates is reported. Enzymatically removed intact hair can be used as a value-added byproduct. Along with dehairing, the enzyme exhibited opening up of collagen bundles. Use of this enzyme in the dehairing process will reduce the time of the process, and yield better quality leather. Simultaneously, it will reduce the pollution load of leather manufacturing process.

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