

STUDIES ON PRODUCTION, CHARACTERIZATION AND APPLICATIONS OF MICROBIAL ALKALINE PROTEASES

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[Received-17/04/2012, Accepted-07/06/2012]

ABSTRACT

With the modern world focusing on eco-friendly products and product output, more and more chemical processes are being replaced by enzymatic methods. Alkaline proteases are one of the most important group of microbial enzymes that find varied uses in various industrial sectors such as leather, detergents, textile, food and feed etc. Industrially important alkaline proteases from bacterial sources have been studied extensively, of which *Bacillus* sp. is most reported. Most of the alkaline proteases that play a role in industries are thermostable as their optimal activity lies between 50⁰C to 70⁰C. The recently used statistical methods have given way to a more rapid optimization process for alkaline protease production. Other than traditional industrial uses, alkaline proteases have promising application in feather degradation and feather meal production for animal feed. This review highlights the alkaline proteases production, optimization of process parameters, characterization as well as their different applicability.

KEYWORDS: Alkaline proteases, *Bacillus*, Detergent, Keratin, Thermostable

INTRODUCTION

Proteases are enzymes occurring everywhere in nature be it inside or on the surface of living organisms such as plants, animals and microbes. These enzymes carry out proteolysis i.e. break down proteins by hydrolysis of the peptide bond that exists between two amino acids of a polypeptide chain. The proteases available today in the market are derived from microbial sources.

This is due to their high productivity, limited cultivation space requirement, easy genetic manipulation, broad biochemical diversity and desirable characteristics that make them suitable for biotechnological applications [1]. The world enzyme market is currently at \$5.1 billion and is expected to rise by 6.3% annually by 2013. Since proteases share a major part of the global enzyme market, an upward trend in their demand is

expected. Proteases can be classified according to their active pH range into neutral, acidic and alkaline proteases. Alkaline proteases (EC.3.4.21–24, 99) are those enzymes that are active at alkaline pH with optimum pH in between 9 to 11. Alkaline proteases are most widely used in industries. The largest application of proteases is in detergent industry where removal of proteinaceous substances occurs at alkaline pH. Another important application of alkaline proteases is in the leather industry for dehairing of hides and skins. Commercially successful uses of alkaline proteases are also mentioned in photographic, textile, chemical and food industry. They can be classified on the basis of chemical nature of the active site into: serine protease and metalloprotease.

Serine proteases are characterized by a serine residue forming a catalytic triad with aspartic acid and histidine in the active site. They are inactivated by organic phosphate esters. Organic phosphate esters acylate the active serine residue thereby rendering the enzyme inactive. They are strongly inhibited by phenylmethylsulfonyl fluoride (PMSF) and diisopropyl fluorophosphate (DFP) and chymostatin. Alkaline proteases from different microbial sources mainly belong to this group. Subtilisins are a major group of alkaline serine proteases mostly active at pH 10 that initiate the nucleophilic attack on the peptide (amide) bond through a serine residue at the active site. Subtilisin Carlsberg and subtilisin novo (sequence homology to subtilisin BPN') are two important types of alkaline proteases obtained from *Bacillus* sp. A subtilisin-like protease BPP-A produced by *Bacillus pumilus* MS-1 can be utilized as an industrial enzyme for the production of zein hydrolysates [2]. Serine alkaline proteases from halophilic sources *Natrialba magadii*, haloalkaliphilic bacterium sp. AH-6 and *Halobacterium* sp. have also been reported [3-5]. Metalloproteases are usually characterized by a catalytic divalent metal ion such as zinc ion in the active site. The ion coordinates to the protein via

three ligands (histidine, glutamate, aspartate, lysine or arginine) and a labile water molecule. They are inhibited by ethylenediaminetetraacetic acid (EDTA). The neutral metalloproteases show specificity for hydrophobic amino acids, while the alkaline metalloproteases possess a very broad specificity [1]. A metalloprotease secreted by *Pseudomonas aeruginosa* MTCC 7926 isolated from solvent-contaminated habitat is suggested to be useful for dehairing of animal skin, anti-staphylococcal activity and processing of X-ray film [6]. Another metalloprotease from *Streptomyces olivochromogenes* was reported to be useful in organic solvent-based enzymatic synthesis and detergent formulation [7]. Alkaline protease purified from entomopathogenic bacterium *Photorhabdus* sp. strain EK1 (PhPrTPI) has been classified as metalloprotease and it is Ca^{2+} dependent. The broad substrate specificity of the enzyme towards different biologically important proteins and peptides suggests the possibility of PhPrTPI being involved in degradation of insect tissues for providing nutrients to the associated nematode [8]. A metalloprotease secreted by *Salinivibrio* sp. strain AF-2004 exhibiting broad pH ranges (5.0–10.0), moderate thermoactivity and halotolerance is suggested for its high commercial value, it being a thermophilic and halophilic alkaline protease [9].

SOURCES OF ALKALINE PROTEASES

Alkaline proteases are obtained from various microbial sources such as bacteria, fungi and certain yeasts. Of all the microbial sources, bacterial proteases are of particular interest due to their various applications in industries such as detergent, textile, leather, food and feed industry. A major source of bacterial alkaline proteases is *Bacillus* species, which has been studied extensively. Some fungal species are also known to produce alkaline proteases of industrial use, of which *Aspergillus* species has been extensively studied. A very few studies exist on yeast species. Alkaline proteases from *Aureobasidium pullulans*, *Yarrowia lipolytica*, *Issatchenkia orientalis* and

Cryptococcus aureus with optimum pH of 9-10 and optimum temp of 45-50° C have been reported for their excellent bioactive peptide production properties [10]. Recently, alkaline proteases from mushrooms have also been studied and purified [11,12]. A list of various microorganisms producing alkaline proteases is shown in Table 1. Some halophilic sources have also been screened for the secretion of alkaline proteases (Table 2).

Halophilic enzymes find rapidly increasing use in biotechnological applications owing to their halotolerance, thermostability for long incubation periods and capability to retain activity in presence of high levels of organic solvents [13,14]. Extracellular alkaline proteases from halophilic bacteria with high pH and thermostability, organic solvent stability and compatibility with detergents have been reported [15]. Capiralla *et al.* [16] have reported alkaline protease from *Halobacterium halobium* S9 with potential application in debittering of protein hydrolysates.

PRODUCTION OF ALKALINE PROTEASES

For industrial use, only large scale production of alkaline proteases can suffice the requirement. Industrial production of alkaline proteases can be carried out by solid-state and submerged fermentations. Media composition [17], particularly carbon and nitrogen source [18] and process parameters such as temperature, pH, agitation speed [19], greatly influence the enzyme production. Each micro-organism producing alkaline protease requires a different medium and process conditions. The effect of carbon and nitrogen sources on the enzyme production has been studied extensively. Akhavan Sepahy and Jabalameli [20] studied the effect of various culture conditions on the production of an extracellular protease by *Bacillus* sp. and reported that sucrose and corn steep liquor are the best substrate for enzyme production. Some alkaline proteases also require metal ions in the form of salts in the production media. It has been shown that $\text{FeSO}_4 \cdot 7\text{H}_2\text{O}$ and $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$ enhanced the protease production by *Bacillus subtilis* RSKK96

[21]. Since cost of fermentation media is considerable, production of alkaline proteases have also been carried out using different agro industrial wastes (green gram husk, chick pea, wheat bran, rice husk, lentil husk, cotton stalk, crushed maize, millet cereal), tannery wastes, shrimp wastes, date wastes etc. [21-27]. Most of the microorganisms produce alkaline proteases at pH 8-9 and temperature of 32-45°C. Table 3 presents different physicochemical parameters for maximum production of alkaline proteases. Researchers are in a continuous process of optimizing production to achieve maximum yield and economical use of available resources. Traditionally, scientists have adopted “one variable at a time” strategy where each variable is optimized independently [28,29]. This is very time consuming, expensive and does not reflect true optimum when a large number of variables are involved because of interference by interaction between them. Recently, a number of statistical methods (Table 4) have been developed such as Taguchi methodology, Plackett–Burman design and response surface methodology (RSM) for optimization to achieve rapid and better understanding of interaction between various variables using a minimum number of experiments [30].

CHARACTERIZATION OF ALKALINE PROTEASES

Alkaline proteases from different sources have been characterized by various workers in order to use them for specific purposes. For example alkaline proteases with broad pH range activity, high thermostability and bleach stable find application in detergent and leather industry. The alkaline protease isolated from *Pseudomonas aeruginosa* is active at a broad pH range of 6–11 and a temperature range of 25–65°C. The studies showed that the purified enzyme retains its activity in surfactants and bleaching agents. These properties indicate its possible use in the detergent industry [6]. The optimum pH for caseinolysis by

most of the alkaline proteases lies in the range 8.0-11.0 at a temperature range of 37-60°C. Most of the bacterial alkaline proteases studied so far have been reported with optimum temperatures in the range of 50-70°C in buffers of pH 9-10, whereas fungal proteases show a pH optimum of 9-10 with optimum temperature in the range of 37-50°C. Alkaline proteases from bacterial sources exhibit higher thermostability than those obtained from fungal sources. Some alkaline proteases show exceptionally high thermostability with no fall in activity at 60-70°C upto 3 hours [31-33].

It has been also observed that alkaline proteases also require a divalent cation like Ca²⁺, Mg²⁺ and Mn²⁺ or a combination of these cations for maximum activity. Supposedly, cations protect the enzyme against thermal denaturation and play a vital role in maintaining the active conformation of the enzyme at high temperatures. Alkaline proteases are completely inhibited by phenylmethylsulfonyl fluoride (PMSF), diisopropyl fluorophosphates (DFP) and ethylenediaminetetraacetic acid (EDTA). Sulfonation of the essential serine residue in the active site by PMSF results in the complete loss of activity (Kumar and Takagi 1999). DFP inhibits the serine alkaline proteases by reacting with the serine residue in the active site followed by formation of a covalent adduct. EDTA inhibits metallotype alkaline proteases by chelating the divalent metal ion in its active site.

The molecular mass reported for most of the alkaline proteases lie in the range 30-45 kDa. Bacterial alkaline proteases with molecular weight below or above this range have been rarely reported. Extracellular alkaline proteases with very low molecular mass of 23 kDa and 15 kDa have been isolated from *Streptomyces aurantiogriseus* EGS-5 and *Bacillus subtilis* PE-11 respectively [34,32]. Alkaline proteases with very high molecular masses of 51 and 66 kDa have also been reported [35,25]. The effect of various parameters on the enzyme characterization along with their molecular weight is summarized in Table 5.

IMMOBILIZATION OF ALKALINE PROTEASES

Despite of various advantages, the use of enzymes in industrial applications has been limited due to their high cost of production, instability and expensive recovery which restrict the repeated use of the enzyme. This led to technological developments in the field of immobilized enzymes/whole cells systems which offer the possibility of a wider and more economical use of enzymes. Several attempts have been made to increase stability to temperature, pH, and organic solvents; recovery and reuse of enzymes by using various techniques of enzyme immobilization. Whole cell immobilization technique has been employed for higher productivity of alkaline proteases by protecting the cells from shear forces, and advantage of reusability and easy recovery of products [36]. Cell free immobilization techniques have also been widely used for the production of proteases. The alkaline proteases are immobilized on solid support matrix by adsorption, covalent binding, ionic binding, cross linking and entrapment. A few examples of the immobilization of whole cells as well as cell free supernatants using various matrices are listed in Table 6.

MOLECULAR CHARACTERIZATION OF ALKALINE PROTEASES

Attempts have been made to deduce the nucleotide and amino acid sequences of alkaline proteases from various sources. These sequences not only help in deriving the primary structure, but also in identifying various functional regions of the proteases. Studies of DNA and protein sequence homology are important for a variety of purposes and have therefore become routine in computational molecular biology. They serve as a prelude to phylogenetic analysis of proteins and assist in predicting the secondary structure of DNA and proteins, and elucidating the structure-function relationship of proteases [1].

Alkaline protease from marine bacterium strain YS-80-122 with 463 amino acid residues and ANGTSSAFTQ as the N-terminal amino acid

sequence has been reported. Sequence alignment revealed that it displayed the highest percentage identity (91%) to an alkaline metalloprotease from psychrophilic *Pseudomonas* sp., with identity to serralyisin-type metalloproteases of other strains, *Pseudomonas fluorescens* Pf-5 and *Dickeya zeae* Ech1591 at 66% and 53% respectively. These results indicate that the alkaline protease belongs to the serralyisin-type metalloproteases [37]. Another alkaline protease BPP-A from *Bacillus pumilus* strain MS-1 consisting of 275 amino acids (mature protein), with a very high homology with that of subtilisin from *Bacillus pumilus* strain TYO-67 and strain UN-31-C-42 (99% amino acid identity) has been studied. Sequence of mature BPP-A contained a conserved essential catalytic triad made up of Asp140, His175 and Ser329 as well as three serine protease (subtilase) family signatures. High sequence homology and conserved sequences of the catalytic triad indicate that BPP-A is a typical subtilisin, which belong to the serine protease (subtilase) family [2].

Similarly, amino acid sequence of halotolerant alkaline proteases from *Bacillus subtilis* FP-133 was >98% identical to that of alkaline serine proteases identified as fibrinolytic enzymes. The mature protein consisted of 275 amino acid residues with a catalytic triad centre containing Asp, His and Ser residues, as identified by comparing the sequence with the homologous serine alkaline protease subtilisins of *Bacillus* and *Brevibacillus* sp.. Three dimensional modeling suggested that the acidic and polar amino acid residues located on the surface stabilize protein structure in the presence of relatively high NaCl concentrations [38].

Multiple sequence alignment of alkaline protease protein sequence of different *Aspergillus* species revealed a stretch of conserved region for amino acid residues from 69 to 110 and 130–204. A motif with a signature amino acid sequence of 50 amino acids was uniformly observed in proteases protein sequences indicating its involvement with the structure and enzymatic function. A

phylogenetic tree was constructed which indicated that several *Aspergillus* species-specific clusters for alkaline proteases exist from *Aspergillus fumigatus*, *Aspergillus niger*, *Aspergillus oryzae*, *Aspergillus clavatus* [39]. A similar result from sequence homology of alkaline protease from *Aspergillus clavatus* with alkaline serine proteases from *Aspergillus fumigatus* and *Verticillium chlamydosporium* revealed 90% and 100% amino acid identity respectively. It also showed high identity with the subtilisin-like protease of *Ophiostoma ulmi* and *Metarhizium anisopliae* (88% and 81% identity respectively), indicating that this enzyme belongs to a serine proteases group and it is a subtilisin-like proteases. These findings were supported by observations made by specific inhibitors and substrates specificity of the enzyme [40].

USES OF ALKALINE PROTEASES

Alkaline proteases are one of the most important classes of proteases from an industrial point of view, occupying a major share of the total enzyme market. Use of alkaline proteases as active ingredients in detergents is the largest application of this enzyme. They are also widely used in leather industry, medical diagnostics, recovery of silver from X-ray films, silk degumming, food and feed industry etc. Due to their vast applications in the industrial processes, many companies started manufacturing them at commercial level. The table 7 gives the commercial manufacturers of alkaline proteases with their product trade name and different applications.

Detergent industry

Alkaline proteases have contributed greatly to the development and improvement of modern household and industrial detergents. They are effective at the moderate temperature and pH values that characterize modern laundering conditions in industrial & institutional cleaning. Various enzymes used in laundry industry are proteases, lipase, cellulases, amylases etc. [41]. Of these, alkaline protease find a major application as

detergent additives because of their ability to hydrolyze and remove proteinaceous stains like blood, egg, gravy, milk etc in high pH conditions [42]. Proteases and other enzymes used in detergent formulations should have high activity and stability over a broad range of pH and temperature. The enzymes used should be effective at low levels (0.4–0.8%). One of the most difficult design challenges that biotechnologists face is that the protease should be compatible with various commercially available detergents and its function is not hampered in the vicinity of typical detergent ingredients, such as surfactants, builders, bleaching agents, bleach activators, fillers, fabric softeners and various other formulation aids. Recently, alkaline proteases from *Bacillus cereus*, *Bacillus pumilus* strain CBS, *Streptomyces* sp. strain AB1, *Bacillus licheniformis*, *Aspergillus flavus*, *Aspergillus niger*, *Bacillus brevis*, *Bacillus subtilis* AG-1 have exhibited excellent detergent compatibility in the presence of certain stabilizers such as CaCl₂ and glycine [43–49]. To survive the extremes of high alkalinity and chelator concentration in detergents, subtilisins have been improved with respect to their thermostability, resistance to chelators [50]. To prevent the loss of activity, several oxidatively stable serine proteases (OSPs) suitable for use in detergents have been isolated from alkaliphilic *Bacillus* strains [51].

Leather industry

Soaking, dehairing of hides and skins and bating have traditionally been carried out by using different chemicals which poses a high tannery waste pollution threat. Hence, proteases with a pH optimum around 9–10 are widely used in soaking to facilitate the water uptake of the hide or skin. Alkaline proteases with elastolytic and keratinolytic activity are used for dehairing and bating process to obtain a desired grain, softness and tightness of leather in a short time. Alkaline proteases with keratinolytic activity have been reported for remarkable dehairing properties [52–24]. A novel protease showing keratinolytic

activity from *B. subtilis* has been studied as a potential for replacing sodium sulfide in the dehairing process of leather industry [55]. Verma *et al.* [31] showed the use of protease from *Thermoactinomyces* sp. RM4 for dehairing goat hides.

Chemical Industry

A high stability in the presence of organic solvents is a feature which is highly desired in applications involving biocatalysis in non-aqueous medium for peptide synthesis. Alkaline proteases from *Aspergillus flavus*, *Bacillus pseudofirmus* SVB1, *Pseudomonas aeruginosa* PseA have shown promising results for potential of peptide synthesis due to their organic solvent stability [46,56,57]. In addition to demonstrating high organic tolerance, alkaline proteases from *B. pumilus* strain CBS and *Streptomyces* sp. strain AB1 are potential strong candidates for use in peptide synthesis in low water systems [44]. Apart from peptide synthesis, they have also been reported for organic synthesis. Alcalase, an industrial alkaline protease has been used for the synthesis of Bz-Arg-Gly-NH₂ (N-benzoylargininylglycinamide), precursor dipeptide of RGDS (Arg-Gly-Asp-Ser) catalysed in water/organic co-solvent systems [58]. Synthesis of 2H-1-benzopyran-2-one derivatives using alkaline protease from *Bacillus licheniformis* has been reported by Wang *et al.* [59]. Regioselective syntheses of polymerizable vinyl guaifenesin ester has been studied by an alkaline protease from *Bacillus subtilis* [60].

Medical Uses

The use of immobilized alkaline protease from *Bacillus subtilis* possessing therapeutic properties has been studied for development of soft gel-based medicinal formulas, ointment compositions, gauze, non-woven tissues and new bandage materials [61]. Oral administration of proteases from *Aspergillus oryzae* has been used as a diagnostic aid to correct certain lytic enzyme deficiency syndromes [1]. Alkaline-fibrinolytic protease have been reported to preferentially

degrade fibrin suggesting its future application in thrombolytic therapy and anticancer drugs [62,35]

Waste management

Wastes from poultry processing industry and leather industry are recalcitrant to commonly known proteolytic enzymes due to presence of keratin-rich wastes whose polypeptide is densely packed and strongly stabilized by several hydrogen bonds and hydrophobic interactions in addition to several disulfide bonds [63]. Chemical and mechanical hydrolysis of keratin wastes is successful but they have several disadvantages of being energy intensive, polluting and leading to loss of essential amino acids. Hence, enzymatic degradation using alkaline proteases with keratinolytic activity (keratinases) is an attractive method [64]. *Bacillus* species is the most widely reported bacterial source of keratinases for feather degradation [65-67]. Other reported bacterial sources of keratinases are *Pseudomonas* sp. MS21, *Microbacterium* sp., *Chryseobacterium* sp. and *Streptomyces* sp. [68-71]. Fungal keratinases from *Aspergillus Oryzae*, *Chrysosporium indicum*, *Trichophyton mentagrophytes*, *Microsporum* sp., *Trichophyton* sp., *Aspergillus terreus*, *Scopulariopsis* sp., *Fusarium oxysporum* have also been studied towards the degradation of keratin [72,73]. Feather hydrolysates obtained after feather degradation can be used as additives for feedstuffs, fertilizers, glues and films or used for the production of the rare amino acids -serine, cysteine, and proline [74].

Feed and Food industry

Alkaline proteases are widely used for production of protein hydrolysates for more than 40 years. Hydrolysates can be used as additives to food and mixed feed to improve their nutritional value. In medicine, they are administered to patients with digestive disorders and food allergies [75]. Protein hydrolysates can be obtained from a variety of substrates such as whey, meat, soy and casein. It is well known that meat hydrolysates usually taste bitter when the degree of hydrolysis (DH) is above

the 10% required for satisfactory solubilization. With Novozymes's Flavourzyme®, it is possible to degrade the bitter peptide groups and obtain a degree of hydrolysis of 20% without bitterness. Alkaline proteases also find their use in meat processing. SEB Tender 70, commercially available proteases is extensively used in meat tenderization to break down collagens in meat to make it more palatable for consumption.

Silver Recovery

Silver is one of the precious and noble metals used in large quantities for many purposes, particularly in the photographic industry. The waste X-ray/photographic films containing black metallic silver spread in gelatin are very good source for silver recovery compared to other types of film. The amount of silver in the X-ray film varies between 1.5 and 2.0% (w/w). Various traditional methods for silver recovery are burning the films directly, oxidation of the metallic silver following electrolysis, stripping the gelatin-silver layer using different chemical solutions. But this method poses serious environmental considerations. For this reason, enzymatic hydrolysis of gelatin is an alternative option to minimize the impact on environment [127]. Successful recovery of silver from X-ray films have been reported by alkaline proteases derived from *Bacillus subtilis*, *Conidiobolus coronatus*, *Streptomyces avermectinus* [76-78]. Recovery of silver and polyethylene terephthalate from used lith film for printing using the thermostabilized mutant enzyme of the alkaline protease from alkaliphilic *Bacillus* sp. B21-2 has also been reported [79].

Silk Degumming

Threads of raw silk must be degummed to remove sericin, a proteinaceous substance that covers the silk fiber. Traditionally, degumming is performed in an alkaline solution containing soap. This is a harsh treatment because the fiber itself is attacked. It also have other disadvantages of high energy consumption, time consumption and also a loss in luster of silk due to the amounts of water used in

this process [80]. However, the use of selected alkaline proteolytic enzymes is a better method because they remove the sericin without attacking the fibre. Tests with high concentrations of enzymes show that there is no fiber damage and the silk threads are stronger than with traditional treatment. In a recent study by Romsomsa *et al.* [81], optimization of silk degumming protease production from *Bacillus subtilis* C4 has been carried out using Plackett-Burman design and response surface methodology.

CONCLUSION

Keeping in view the industrial importance of alkaline proteases, researchers are more and more focusing on discovering alkaline proteases with novel properties to meet the industrial requirements as well as increasing demand of global enzyme market. Other than traditional uses in detergent industry for removing protein stains, leather industry for dehairing, textile industry for silk degumming, photographic industry for silver recovery, food and feed industry, chemical industry for peptide synthesis, they are increasingly finding potential future applications in pharmaceutical industry. Recently, alkaline proteases with keratinolytic activity are finding an interesting application in the feather degradation for reducing poultry waste and feather meal production for fertilizers, glues, biodegradable films, and foils. Through this review, an attempt has been made to highlight the recent advances in the field of alkaline proteases and their applicability.

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Tables:**Table 1. Alkaline proteases from various sources.**

| Microorganisms | Applications | References |
|--|--|--------------------------------|
| <i>Pseudomonas aeruginosa</i> PD 100 | Collagen replacement therapy, waste treatment, removal of blood stains and dehairing | Najafi <i>et al.</i> [82] |
| <i>Thermoactinomyces</i> sp. RM 4 | Leather industry for dehairing process | Verma <i>et al.</i> [31] |
| <i>Bacillus</i> sp. Tk1 and Tk2 | Detergent industries, food industries and pharmaceutical industries | Kuberan <i>et al.</i> [83] |
| <i>Bacillus pumilus</i> Cbs | Detergents, dehairing | Jaouadi <i>et al.</i> [44] |
| <i>Streptomyces</i> sp. Ab1 | Dehairing, feather hydrosylation | Jaouadi <i>et al.</i> [44] |
| <i>Bacillus subtilis</i> PE-11 | Detergents | Adinarayana <i>et al.</i> [32] |
| <i>Bacillus</i> sp. SSR1 | Detergent industry | Singh <i>et al.</i> [84] |
| <i>Bacillus licheniformis</i> RP1 | Chitin extraction, chicken feather degradation and dehairing | Haddar <i>et al.</i> [26] |
| <i>Bacillus coagulans</i> and <i>licheniformis</i> | Industrial sector | Asokan and Jayanthi [85] |
| <i>Bacillus clausii</i> | Industrial sector | Vadlamani <i>et al.</i> [86] |
| <i>Bacillus cereus</i> | Detergents and removal of blood stains | Abou-Elela <i>et al.</i> [43] |
| <i>Bacillus</i> sp. K-30 | Deproteinisation of rice bran and detergents | Naidu <i>et al.</i> [87] |
| <i>Bacillus cereus</i> 1173900 | Dehairing | Ravindran <i>et al.</i> [25] |
| <i>Bacillus circulans</i> | - | Prakasham <i>et al.</i> [23] |
| <i>Bacillus subtilis</i> | Depilating agent in leather processing | Mukhtar and Haq [88] |
| <i>Pseudomonas fluorescens</i> | Detergent and textile industry | Kalaiarasi and Sunitha [89] |
| <i>Microbacterium</i> strain AR-68 | Detergent and leather industry | Gessesse and Gashe [90] |
| <i>Streptomyces pulvereus</i> | - | Jayasree <i>et al.</i> [29] |
| <i>Stenotrophomonas maltophilia</i> (MTCC 7528) | Detergent additive for cold washing and environmental bioremediation in cold regions | Kuddus and Ramteke [91] |
| <i>Shewanella oneidensis</i> MR-1 | Industrial sector | Anbu <i>et al.</i> [92] |
| <i>Streptomyces aurantiogriseus</i> EGS-5 | High temperature industrial process | Ahmad [34] |
| <i>Aspergillus niger</i> | Detergents | Devi <i>et al.</i> [47] |

| | | |
|--|--|-------------------------------|
| Genetically engineered <i>Aspergillus oryzae</i> U1521 | Animal food and feed processing | Samarntarn <i>et al.</i> [93] |
| <i>Aspergillus flavus</i> and <i>Aspergillus terreus</i> | Leather and food industry | Chellapandi [94] |
| <i>Aspergillus nidulans</i> HA-10 | Food processing, pharmaceutical and leather industries | Charles <i>et al.</i> [95] |
| <i>Engyodontium album</i> BTMFS10 | Detergents and silver recovery | Jasmin <i>et al.</i> [96] |
| <i>Aspergillus flavus</i> | Detergent industry, peptide synthesis | Yadav <i>et al.</i> [46] |

Table 2. Halophilic sources of alkaline proteases.

| Microorganisms | Optimum pH | Optimum Temperature (°C) | References |
|---|------------|--------------------------|-----------------------------------|
| <i>Chromohalobacter</i> TVSP101 | 8 | 75 | Vidyasagar <i>et al.</i> [7] |
| <i>Virgibacillus marismortui</i> NB2-1 | 10 | 50 | Chamroensaksri <i>et al.</i> [98] |
| <i>Natronococcus occultus</i> | 7-9 | 60 | Studdert <i>et al.</i> [99] |
| <i>Natrialba magadii</i> | 8-10 | 60 | Giménez <i>et al.</i> [3] |
| <i>Bacillus</i> sp. HS-4 | 8 | 37 | Sehar and Hameed [100] |
| <i>Halobacterium</i> sp Js1 | 7.5-8 | 40 | Vijayanand <i>et al.</i> [101] |
| Haloalkaliphilic Bacteria | 9 | 50 | Dodia <i>et al.</i> [102] |
| <i>Bacillus halodurans</i> SK5 | 10 | 70 | Ibrahim <i>et al.</i> [103] |
| <i>Halobacterium mediterranei</i> | 8-8.5 | 55 | Stepanov <i>et al.</i> [104] |
| <i>Streptomyces clavuligerus</i> strain Mit-1 | 9 | - | Thumar and Singh [105] |
| <i>Bacillus licheniformis</i> | 10 | 60 | Öztürk <i>et al.</i> [106] |

Table 3. Physiochemical parameters for alkaline proteases production.

| Microorganisms | Carbon source | Nitrogen source | pH | Temperature (°C) | Agitation speed (rpm) | References |
|---|------------------|------------------------|------|------------------|-----------------------|------------------------------------|
| <i>Bacillus licheniformis</i> N-2 | Glucose | Defatted Soyabean meal | 10 | 37 | 140 | Nadeem <i>et al.</i> [22] |
| <i>Bacillus</i> sp. strain CR-179 | Starch, Maltose | Corn steep liquor | 8 | 45 | 150 | Akhavan Sepahy and Jabalameli [20] |
| <i>Aspergillus niger</i> | Glucose | Ammonium sulphate | 8.5 | 45 | 150 | Devi <i>et al.</i> [47] |
| <i>Halobacterium</i> sp JS1 | Rice bran | Defatted Soyabean meal | 7 | 40 | 200 | Vijayanand <i>et al.</i> [101] |
| <i>Bacillus subtilis</i> SVR-07 | Glucose | Soyabean meal | 9 | 55 | 150 | Reddy <i>et al.</i> [107] |
| <i>Bacillus brevis</i> | Lactose | Soyabean meal | 10.5 | 37 | 200 | Banerjee <i>et al.</i> [108] |
| <i>Bacillus licheniformis</i> NCIM-2042 | Starch | Soyabean meal | 7 | 37 | 180 | Bhunia <i>et al.</i> [28] |
| <i>Bacillus subtilis</i> RSKK96 | Arabinose | Beef extract | 9 | 37 | 150 | Akcan and Uyar [21] |
| <i>Pseudomonas fluorescens</i> | Wheat bran | Peptone | 9 | 37 | -- | Kalaiarasi and Sunitha [89] |
| <i>Microbacterium</i> strain AR-68 | Glucose, Sucrose | Peptone | 10.3 | 32 | -- | Gessesse and Gashe [90] |
| <i>Streptomyces pulvereceus</i> | Starch | Casein | 9 | 33 | -- | Jayasree <i>et al.</i> [29] |

Table 4. Statistical methods for optimizing production of alkaline proteases.

| Optimization methods | Micro organism | Parameters Optimized | Yield improvement | References |
|--------------------------------|---|--|-------------------|--------------------------------------|
| Plackett–Burman design and RSM | <i>Bacillus licheniformis</i> NCIM 2042 | Inoculum size, temperature, pH and agitation | 1.71 fold | Bhunia and Dey [109] |
| Plackett–Burman design | <i>Bacillus subtilis</i> DKMNR | Carbon and nitrogen source | 4.4 fold | Kezia <i>et al.</i> [110] |
| Plackett–Burman design and RSM | <i>Bacillus subtilis</i> C4 | Medium components and agitation | 2.2 fold | Romsomsa <i>et al.</i> [81] |
| Response surface methodology | <i>Bacillus subtilis</i> HB04 | Medium components, temperature and agitation | 8 fold | Venil and Lakshmanaperumalsamy [111] |
| Plackett–Burman design and RSM | <i>Aspergillus clavatus</i> ES1 | Media constituents, pH and temperature | 14 fold | Hajji <i>et al.</i> [30] |
| Plackett–Burman and RSM | <i>Bacillus</i> sp. RKY3 | Inoculum size and medium components | 2 fold | Reddy <i>et al.</i> [112] |
| Plackett–Burman method | <i>Halobacterium</i> sp. SP1 | Medium components | 3.9 fold | Akolkar <i>et al.</i> [113] |
| Taguchi design of experiments | <i>Bacillus clausii</i> | Medium components and agitation | 4 fold | Oskouie <i>et al.</i> [114] |
| Box-Behnken design | <i>Aspergillus oryzae</i> | Carbon and nitrogen source, temperature, pH | 3 fold | Babu <i>et al.</i> [115] |

Table 5. Characterization of Alkaline proteases.

| Sources | Mol. Wt. (kDa) | Optimum pH | Optimum Temp. (°C) | Effect of metal ions | | Inhibitors | References |
|---|----------------|------------|--------------------|--|---|-----------------------------|--------------------------------|
| | | | | Stimulatory | Inhibitory | | |
| <i>Bacillus subtilis</i> PE-11 | 11 | 10 | 60 | Ca ²⁺ , Mg ²⁺ , Mn ²⁺ | - | PMSF, DFP | Adinarayana <i>et al.</i> [32] |
| <i>Aspergillus flavus</i> | -- | 9 | 37 | Fe ³⁺ | Hg ²⁺ , Cu ²⁺ | PMSF, EDTA | Yadav <i>et al.</i> [46] |
| <i>Aspergillus niger</i> | 38 | 10 | 50 | Ca ²⁺ | Zn ²⁺ , Co ²⁺ | EDTA | Devi <i>et al.</i> [47] |
| <i>Bacillus cereus</i> | 31 | 10 | 50 | Cu ²⁺ | Ca ²⁺ , Mg ²⁺ , and Mn ²⁺ | EDTA | Abou-Elela <i>et al.</i> [43] |
| <i>Streptomyces aurantiogriseus</i> EGS-5 | 23 | 10-11 | 40 | - | - | DFP | Ahmad [34] |
| <i>Aspergillus nidulans</i> HA-10 | 42 | 8 | 35 | - | - | mercaptoethanol, PMSF, EDTA | Charles <i>et al.</i> [95] |
| <i>Bacillus cereus</i> | 66 | 8 | 40 | Ca ²⁺ , Mg ²⁺ , Mn ²⁺ | Zn ²⁺ , Cu ²⁺ and Fe ²⁺ | EDTA and EGTA | Ravindran <i>et al.</i> [25] |
| <i>Natrialba magadii</i> | 45 | 8 | 45 | - | - | DFP, PMSF, chymostatin | Giménez <i>et al.</i> [3] |
| <i>Streptomyces olivochromogen</i> | -- | 7.5 | 50 | Ca ²⁺ | Co ²⁺ , Cu ²⁺ , Zn ²⁺ , Mn ²⁺ | EDTA | Simkhada <i>et al.</i> [7] |

| | | | | | | | |
|---------------------|----|-----|----|--------------------------------------|--------------------------------------|----------------------------|-----------------------------|
| <i>es</i> | | | | | and Fe ²⁺ | | |
| <i>Bacillus</i> sp. | -- | 9.5 | 60 | Ca ²⁺ and M ²⁺ | Ag ⁺ and Co ²⁺ | DFP, PMSF, Iodoacetic acid | Saurabh <i>et al.</i> [116] |

Table 6. Immobilization of alkaline proteases in different matrices.

| Microorganisms | Matrix used | Stability and activity of enzymes | References |
|---|---|---|--------------------------------|
| <i>Bacillus subtilis</i> # | Agar, gelatin, sodium alginate and polyacrylamide | Increased activity | Kumar and Vats [117] |
| <i>Bacillus subtilis</i> PE-11# | Calcium alginate, k-Carrageenan, polyacrylamide, agar-agar, and gelatin | Higher volumetric activities, long life-term stability, reusability | Adinarayana <i>et al.</i> [36] |
| <i>Bacillus circulans</i> # | agar-agar and calcium alginate | Increased activity | Kocher and Mishra [118] |
| <i>Bacillus subtilis</i> -K 30# | calcium alginate, polyacrylamide, agar-agar and gelatin | Higher volumetric activities , long life-term stability, reusability | Naidu <i>et al.</i> [119] |
| <i>Bacillus licheniformis</i> # | Calcium alginate, k-Carrageenan, agar-agar | Increased activity | Ahmed and Abdel-Fattah [120] |
| <i>Bacillus pumilis</i> # | calcium alginate, polyacrylamide and agar-agar | Increased activity | Kumari <i>et al.</i> [121] |
| <i>Bacillus sphaericus</i> [^] | Sawdust, perlite and silica | Increased stability | Surendran <i>et al.</i> [122] |
| <i>Myceliophthora</i> sp [^] | Calcium alginate beads | Increase in optimum temperature , thermal stability and stability for up to 7 reuses | Zanphorlin <i>et al.</i> [123] |
| <i>Streptomyces avermectinus</i> [^] | Sponge | Increased thermostability, half life, activation energy , reuse after 5 cycles | Ahmed <i>et al.</i> [78] |
| <i>Aspergillus oryzae</i> [^] | Calcium alginate beads | Increase in Km value, optimum pH, enhanced stability in acidic as well as alkaline environments | Sharma <i>et al.</i> [124] |
| <i>Bacillus licheniformis</i> [^] | Loofa | Increased activity, half life, activation energy, stability in EDTA | Ahmed <i>et al.</i> [125] |
| <i>Bacillus polymyxa</i> [^] | Perlite | Increased stability | Emtiazi <i>et al.</i> [126] |

whole cell immobilization, ^ cell free immobilization

Table 7. Commercial manufacturers, trade name and applications of alkaline proteases.

| Manufacturers | Trade name of product | Industrial Uses | Applications |
|--------------------|-----------------------|------------------------|---|
| Novozymes, Denmark | RONOZYME ProAct | Feed industry | Maximize protein utilization and improving nutritional value of animal feed |
| | Novozym 37020 | Meat and feed industry | Enhance protein quality |
| | Savinase, Everlase | Detergent industry | Remove protein based stains |
| | Ovozyme | | Remove egg soils |
| | Polarzyme | | Cold water hand wash |
| | Alcalase | Chemical industry | Resolution of N-protected amino acids , peptide synthesis |
| | Textile | Silk degumming | |

| | | | |
|---|----------------------|---------------------------------|--|
| | | Detergent | Remove protein based stains |
| | Novobate 1547 | Leather industry | Bating |
| | NovoLime | | liming |
| | NovoCor SG | | Soaking |
| Genecor, USA | Purafast | Detergent industry | cold wash conditions |
| | Purafect Prime L | | High alkaline performance at lower dose rate |
| | Purafect OX L | | High alkaline protease with superior storage stability |
| | PrimaTan | Leather industry | Bating |
| | Purafect 4000L | | Soaking |
| | Protex 6L | Food industry | Hydrolysis of milk fractions, baking, protein processing, pet food production |
| | | Photographic industry | Recovery of silver from films |
| Advanced Enzymes, India | SEBsoak | Leather industry | Soaking |
| | SEBlime | | Dehairing, liming |
| | SEBate Alakli | | Bating |
| | Protosol | Detergent industry | Cleaning and washing |
| Amno Enzyme, Japan | Peptidase R, ProteAX | Feed industry | Bitterless protein hydrosylate |
| | THERMOASE PC10F | Leather industry | Bating and dehairing |
| | PROTIN SD-AY10 | Photographic industry | Recovery of waste silver in the film industry |
| Maps Enzyme Limited, India | Palkobate | Leather industry | Bating |
| | Palkosoak | | Soaking |
| | Palkodehair | | Dehairing |
| | Palkogent | Detergent industry | Removal of protein stains |
| AB Enzymes, Germany | COROLASE H-PH | Food and Feed industry | Protein hydrolysis |
| | COROLASE LAP | | Bitterless Protein hydrolysis of animal or vegetable raw materials, e.g. whey, casein, soy |
| Speciality enzymes and biotechnologies, USA | SEBalase BP | Detergent industry | Removal of protein stains |
| | SEB Tender 70 | Meat industry | Tenderization of meat |
| | SEBDigest | Food and Feed industry | Protein hydrolysis |
| | SEBpro A | | develop savory flavors and debitterment of protein hydrolysates |
| Rossari Biotech, India | Lyserine ESD Powder | Silk industry | Degumming of silk |
| Kao Corp., Tokyo, Japan | Proteinase K-16 | Detergent industry | Protein removal |
| Bachauna Technology, India | Verma Batzyme | Leather industry | Bating |
| Sunson, China | PRAL800 | Leather, silk and food industry | Leather depilation, silk degumming and protein hydrolysis |