

Review on bacterial production of alkaline pectinase with special emphasis on *Bacillus* species

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ABSTRACT

Pectinases consist of an exclusive group of enzymes which catalyze the degradation of pectic polymers present in the plant cell walls. Today, pectinases are the upcoming industrially important enzyme having major industrial importance and they hold a leading position among the commercially produced industrial enzymes. Microorganisms including yeast, bacteria, actinomycetes and a large number of filamentous fungi are commonly recognized as the best natural sources for the production of pectinase enzyme. The chief source of acidic pectinases is fungi but alkaline pectinases are produced from alkalophilic bacteria, primarily *Bacillus* spp. The alkaline pectinase has developed as important commercial enzymes with far-flung applications mainly in textile processing, bio-scouring of cotton fibers, degumming and retting of fiber crops, pretreatment of pectic wastewater etc. This review discusses the microbial production of pectinases with special emphasis on bacterial pectinase from *Bacillus* spp.

KEY WORDS: PECTINASE, *BACILLUS*, POLYGALACTURONASE, PECTATE LYASE, PECTIN LYASE

INTRODUCTION

Pectinases comprises of a unique group of enzymes which catalyze the degradation of pectic polymers present in the plant cell wall. They belong to the family of polysaccharidases that contribute to the breakdown of pectins from a variety of plants and are also known as pectolytic or pectic enzymes (Prathyusha & Suneetha, 2011). In the current biotechnological era, pectinase are one of the forthcoming enzymes showing progressive

increase in their market. They maintained the average annual growth rate of 2.86% from 27.6 million \$ in 2013 to 30.0 million \$ in 2016 and it is estimated that by 2021, the market size of the pectinase will reach 35.5 million \$ (Global Pectinase Market Research Report, 2017).

The pectinase enzymes are classified based on their mode of action against the galacturonan backbone of pectins. Primarily, there are two groups of pectic enzyme; the de-esterifying enzymes which catalyses the deesterification of pectins and the depolymeriz-

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ing enzymes which break the glycosidic α -(1-4) bonds between GalA residues either by hydrolysis (hydrolases) or by transesterification (lyases). Moreover, the latter two types of enzymes are classified on the basis of whether they exhibit a preferential hydrolytic /transesterification power against pectin, pectic acid or oligogalacturonate as the substrate and whether the pattern of action is random (endo-) or terminal (exo-). Another group of the pectic enzyme is protopectinase (PPase) which convert insoluble native protopectin into soluble pectins (Alkorta *et al.*, 1998; Kashyap *et al.*, 2001; Sharma *et al.*, 2013; Garg *et al.*, 2016; Hassan & Ali, 2016). The classifications of pectic enzymes acting on pectic substances are given in Figure 1.

Based on the optimum pH for enzyme activity, pectic enzymes are also classified into two, acidic and alkaline pectinase. Research and application of acidic pectinase are more established compared with alkaline pectinase. The chief source of acidic pectinase is from fungus *Aspergillus niger*. Acidic pectinases are used mainly in fruit juice industry for extraction and clarification of fruit juices, improvement of chromaticity and stability of red wines etc. They are also having application in maceration of plant tissue, liquefaction and saccharification of biomass, isolation of protoplasts. Whereas, alkaline pectinase are mostly used in the degumming and retting of fiber crops, textile processing and bio-scouring of cotton

fibers, pretreatment of pectic wastewater from fruit juice industries, paper making, coffee and tea fermentation, enzyme based oil extraction etc. The important applications of pectinase are shown in Figure 2. Bacteria, mainly *Bacillus* spp. are used for the production of alkaline pectinase (Kashyap *et al.*, 2001; Jayani *et al.*, 2005). Information regarding pectinase types, structure, applications, substrate etc. has been reviewed previously by many authors (Alkorta *et al.*, 1998; Kashyap *et al.*, 2001; Hoondal *et al.*, 2002; Jayani *et al.*, 2005; Prathyusha & Suneetha, 2011; Sharma *et al.*, 2013; Kohli *et al.*, 2015; Hassan & Ali, 2016). This review emphasizes reports on pectinase production by *Bacillus* spp.

Microbial production of pectinase: It is a well-known fact that, microbes are the prominent source of enzymes because they allow an economical technology with truncated resource consumption and low emission involving no social and political issues, as in the case of animal and plant sources (Chaplin & Bucke, 1990). It is reported that, fifty percent of accessible enzymes are initiated from fungi and yeast; 35 % from bacteria, while the remaining 15 % are either of plant or animal origin. Filamentous microorganisms are most widely used for pectinase production (Soares *et al.*, 1999). Microbes are chosen as a source of enzyme production compared to plants and animals because; (a) they produce a wide

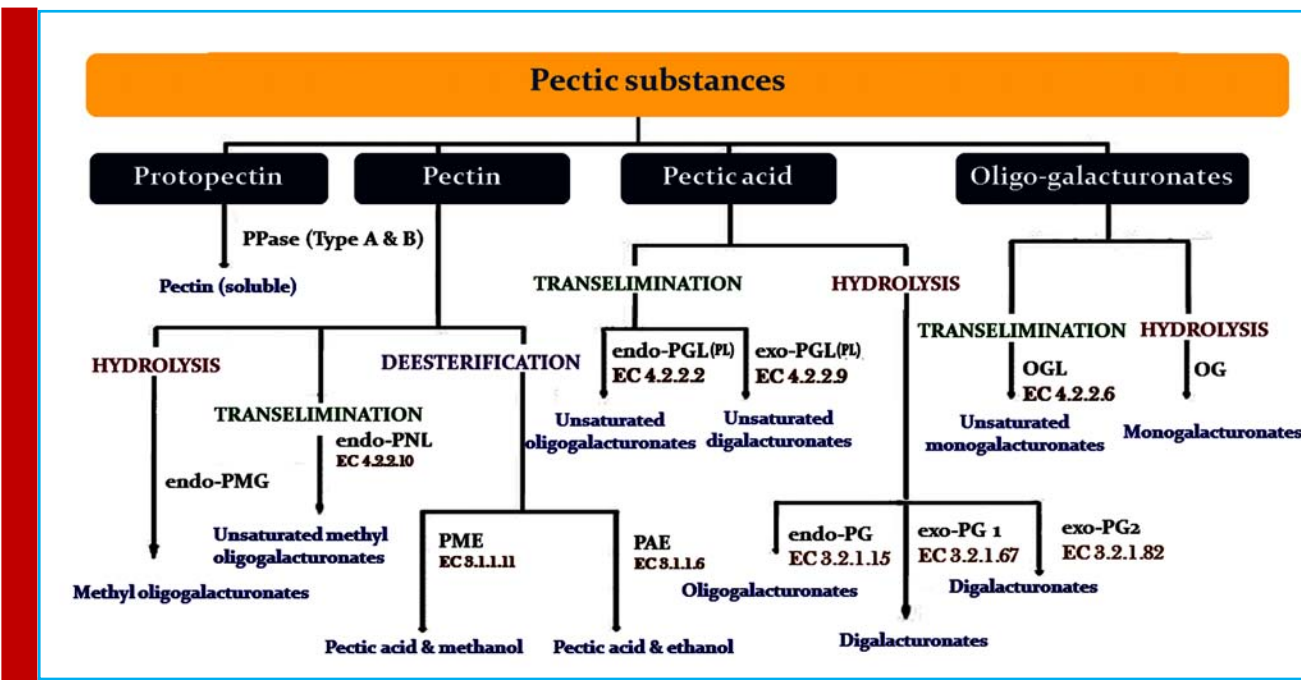


FIGURE 1. Classification of pectic enzymes acting on pectic substances – data modified from (Alkorta *et al.*, 1998; Jayani *et al.*, 2005) PMG: Polymethylgalacturonase, PNL: Pectinlyase, PGL: Polygalacturonatelyase, PN: Pectate lyase PME: Pectin methylesterase, PAE: Pectin acetyl esterase, OGL: Oligogalacturonidelyase, OG: Oligogalacturonase

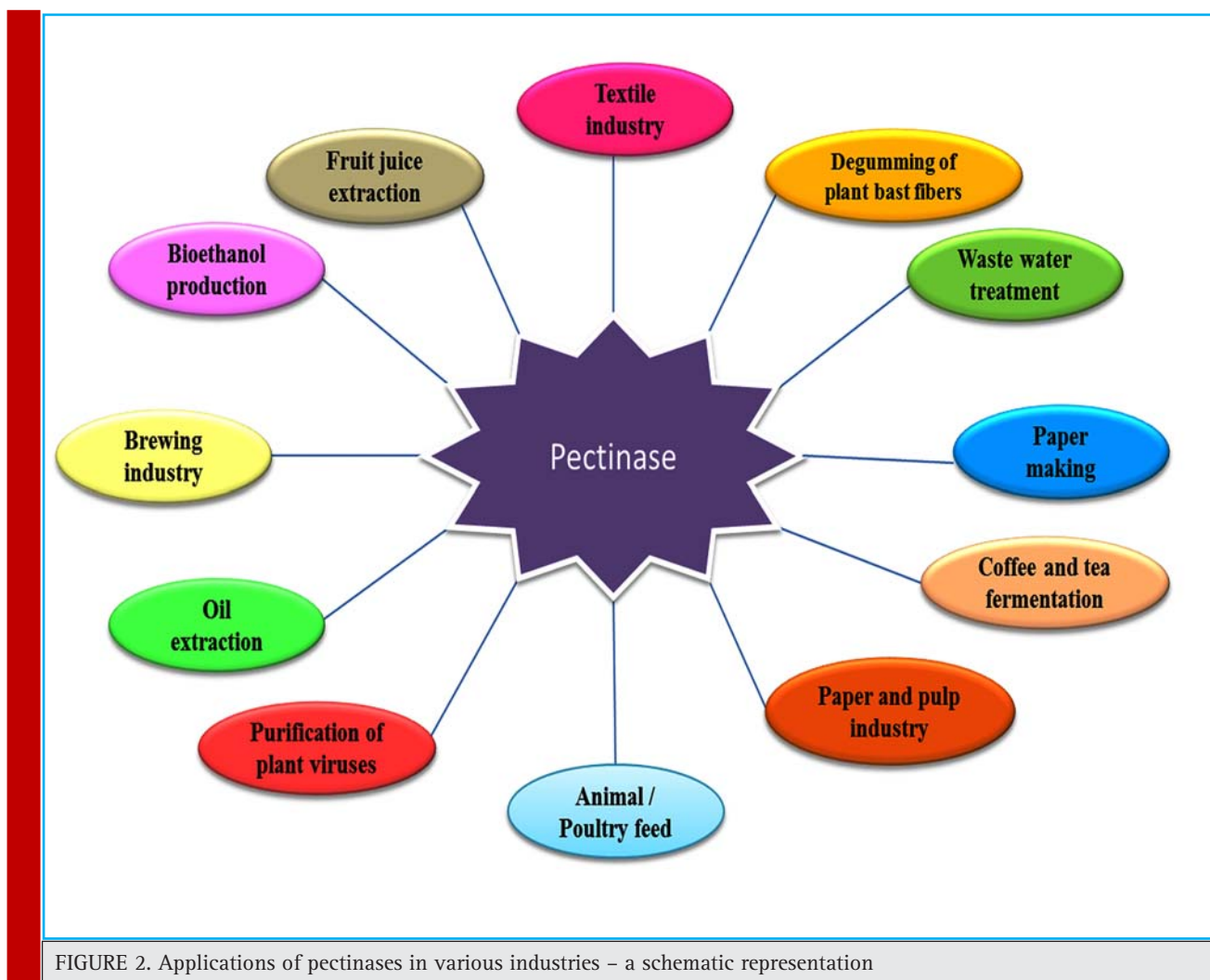


FIGURE 2. Applications of pectinases in various industries – a schematic representation

variety of enzymes and their enzyme contents are more predictable and controllable, (b) generally economical in bulk production and dependable provisions for raw material of constant composition, (c) high productivity rate and enzymes obtained via microbial source are greater in volume, (d) microbes are easy to manipulate to derive enzymes of desired nature and they can be cultured in large quantities in a relatively short period of time by the established method of fermentation using sophisticated tools, (e) they can be made to produce enzymes over wide range of environmental condition and (f) plant and animal tissues contain more potentially harmful materials than microbes, including phenolic compounds (from plants), endogenous enzyme inhibitors and proteases (Chaplin & Bucke, 1990).

The production of pectin degrading enzymes has been extensively reported and meticulously studied in bacteria and filamentous fungi because they play an essential role in the phytopathogenesis (Blanco *et al.*, 1999). Several reports are available regarding pectinase enzymes

by the microorganism such as bacteria, fungi, yeast and actinomycetes. They are also distributed in higher plants and in some protozoa, nematodes and insects but they are not found in higher animals (Jayani *et al.*, 2005; Pedrolli *et al.*, 2009; Sharma *et al.*, 2013).

Pectinase production by fungi: Filamentous fungi have been used for more than 50 years in the production of industrial enzymes (Dalboge, 1997). Many extracellular enzymes are produced by fungi which are capable of decomposing organic matter and one such enzyme is pectinolytic enzymes. Filamentous fungi are considered as one of the most potent producers of pectinases and they can be employed extensively in SSF for the economic production process. Various types of fungal species have been reported to be employed for the production of pectinases. *Aspergillus niger* is the most commonly used fungal species for industrial production of pectinolytic enzymes (Gummadi & Panda, 2003). Findings on the isolation, characterization, selection, properties and fermen-

tation of *A. niger* strains for the production of pectinolytic enzymes using different substrates was made by different workers (Finkler *et al.*, 2017). Other species of *Aspergillus* were also reported to produce pectinase including *A. oryzae*, *A. fumigatus*, *A. terreus*, *A. sojae*, *A. awamori* etc (Pedrolli *et al.*, 2009; Garg *et al.*, 2016). Production of endo-PGL was firstly reported in *A. giganteus* (Pedrolli & Carmona, 2014). Further, species of *Penicillium*, *Fusarium*, *Mucor*, *Neurospora crassa*, *Sclerotinia sclerotiorum* etc also have role in pectinase production (Pedrolli *et al.*, 2009; Pedrolli & Carmona, 2014; Garg *et al.*, 2016). The fungus produces these enzymes to break down the middle lamella in plants so that it can extract nutrients from the plant tissues and insert fungal hyphae.

Pectic enzyme production in yeasts: The pectinolytic enzyme production in yeasts has received less consideration and only a few yeast species show this ability (Blanco *et al.*, 1999). The first reports of pectinolytic (endo-PG) production by yeasts were described by Luh and Phaff in 1951 in *Saccharomyces fragilis*. Among different yeast species representing all yeast genera, they found that only six (*S. fragilis*, *S. fragilis* var. no. 351, *S. thermantitonus*, *Torulopsis kefyri*, *Candida pseudotropicalis* var. *lactosa*, and *Candida pseudotropicalis*) were capable of causing a noticeable change in pectin (Luh & Phaff, 1951). Recent reports are also added to describe the ability of Yeast species for pectinase production these includes; *Saccharomyces sp.*, *Cryptococcus sp.*, *Aureobasidium pullulans*, *Rhodotorula dairenensis*, *Kluyveromyces marxianus*, *Geotrichum klebahnii*, *Wickerhamomyces anomalus* etc. (Alimardani Theuil *et al.*, 2011; Merin *et al.*, 2015; Hassan & Ali, 2016; Naumov *et al.*, 2016).

Bacterial production of pectinase: A review of the currently available literature reveals little quantitative information about the diversity of bacterial genera having pectinolytic properties. Bacterial pectinase is produced mainly by bacteria belonging to genera *Bacillus* and *Erwinia*. Elyrod in 1942 reported that the bacterium *Erwinia* sp. can degrade pectin with the aid of pectin degrading enzymes (Elrod, 1942) and in the past decade, bacterial pectinase biosynthesis has been extensively studied in phytopathogens, especially in the soft-rotting *Erwinia* species *E. carotovora* and *E. chrysanthemi* which are reported to produce a set of pectinolytic enzymes such as PL, PG, PME, and PAE (Matsumoto *et al.*, 2003).

Chessen *et al.*, 1980 reviewed that bacteria like *Bacillus*, *Pseudomonas* and *Micrococcus* isolated from retting flax, jute, sisal and coir, and *Erwinia* from coffee fruits have shown to possess the ability to degrade pectin by producing pectinolytic enzymes (Chesson, 1980). Other bacterial genera reported to have pectinolytic properties include species of *Pseudomonas* (Sohail & Latif, 2016),

Streptomyces (Ramirez-Tapias *et al.*, 2015) *Lactobacillus* (Karam & Belarbi, 1995) etc.

Importance of bacterial production of pectinase: *Aspergillus niger*, a GRAS microorganism is the major organism used for the industrial production of pectinase. However, this mold also secretes several other enzymes which may trigger collateral reactions such as the release of volatile phenols less desirable for the production of wine or fruit juices, for instance arabinofuranosidase, which can cause turbidity (Whitaker, 1990). Pectinases from fungal sources are produced best under acidic pH and low temperature conditions and can therefore not be used in various industrial bioprocesses which utilize neutral to alkaline pH with high temperatures exceeding 45°C. It has been shown that bacteria produce pectinase that withstands high pH and temperature (Hoondal *et al.*, 2002; Andrade *et al.*, 2011). Also, it is easy to harvest pectinase than fungus as it is an extracellular product in bacterial culture (Sohail & Latif, 2016). Bacterial strains producing commercial enzymes are always preferred over fungal strains because of ease of fermentation process (for production) and implementation of strain improvement techniques or any modern technique to increase the production yield (Prathyusha & Suneetha, 2011). Moreover, bacterial pectinases with novel properties have the added advantage that enzyme production is achieved in less time as compared to fungal sources (Joshi *et al.*, 2015).

Bacillus species for industrial enzyme production: Among the diverse types of microorganisms inhabiting in the soil, bacteria are the amplest and major organism. Considering the bacterial genera of soil, Bacilli are most abundant followed by Cocci and Spirilla. The genus *Bacillus* and *Cocci* comprises several varieties of industrially important species contributing approximately half of the existing commercial production of bulk enzymes (Priest, 1977; Aisha & Barate, 2016). *Bacillus* species have been the imperative industrial enzyme producers with roles in applied microbiology for over a millennium.

Because of several reasons *Bacillus* species continue to be the predominant bacterial workhorses in microbial fermentations (Schallmeyer *et al.*, 2004; Satyanarayana.T *et al.*, 2005). They produce more than two dozen of biologically active molecules generating a high potential for biotechnological and biopharmaceutical applications (Stein, 2005). Also, it is estimated that enzymes from *Bacillus* spp. makeup about 50 % of the total enzyme market (Schallmeyer *et al.*, 2004). Another major feature that makes these groups predominant is that most of them are environmental friendly, don't have fastidious nutritional requirements and are easy to grow and handle (Sohail & Latif, 2016). *Bacillus* species such as *B. subtilis* and *B. licheniformis* are on the Food and Drug

Administration's GRAS (generally regarded as safe) status (Food and Drug Administration, 1999; Schallmeyer *et al.*, 2004). Moreover, the biochemistry, physiology, and genetics of *B. subtilis* and other species are well studied and the complete genome sequence of *B. subtilis* 168 comprising of 4100 protein-coding genes has been published in 1997 (Kunst *et al.*, 1997).

Various properties of *Bacillus* strains are reported which make them superior in industrial biotechnology, including their high growth rates leading to short fermentation cycle times, ability to secrete proteins into the extracellular medium (Acton, 2012; Barros *et al.*, 2013) and their ability to adapt with changing environmental and nutritional conditions (Buescher *et al.*, 2012; Nicolas *et al.*, 2012). The capacity of selected *Bacillus* strains to produce and secrete large quantities (20–25 g/L) of extracellular enzymes has placed them among the most important industrial enzyme producers (Satyanarayana *et al.*, 2005). Moreover, many researchers used strains belonging to *Bacillus*, because this genus includes strains (such as *B. subtilis*) that can grow on cheap substrates such as agro-wastes (Sakai *et al.*, 1989).

Pectinase production by *Bacillus* spp.: Pectic enzymes are of functional relevance in the retting process and evidence regarding pectinolytic properties of *Bacillus* spp. was recorded years ago. Different species of the genus *Bacillus* have been reported to be retting agents and active against pectic materials (Potter & McCoy, 1955). Nortje and Vaughn, 1953 tested the pectinolytic activity of *B. subtilis* and *B. pumilus* in relation to the softening of olives and pickles (Nortje & Vaughn, 1953). The first *in vitro* fermentation studies of pectin and pectic acid was reported in 1955 using *B. polymyxa* strain 30 (Potter & McCoy, 1955). Over the past few years, pectinolytic properties have been described in several *Bacillus* species.

It is evident from many research works that, among different bacterial isolates screened for pectinolytic properties *Bacillus* strains were selected as the most potent enzyme producers (Soares *et al.*, 1999; Jayani *et al.*, 2010; Rehman *et al.*, 2012; Kavuthodi *et al.*, 2015; Sohail & Latif, 2016). As mentioned in the introduction, alkaline pectinases have a wide variety of industrial applications and, bacteria mainly *Bacillus* spp. are the chief producers. Apart from this fact, there are also some other reasons for researchers to focus on pectinase from *Bacillus* spp. These include; (i) they produce all class of pectic enzymes, (ii) have short fermentation period for enzyme production, (iii) can produce enzymes very economically by using different agro-wastes as cheap substrates, (iv) fermentation can be attained by either SSF, Smf and (v) genetic information regarding pectinase genes of many *Bacillus* spp. are available in various nucleotide sequence databases. Thus it supports

successful cloning and expression of pectinase gene in other organisms.

Pectic enzymes from *Bacillus* sp.: *Bacillus* spp. are renowned producers of all classes of pectinases including hydrolases, lyases, esterases and protopectinases. The important strains of *Bacillus* reported for different groups of pectinases are listed under;

Hydrolases: The pectic hydrolases are a type of depolymerases that split the α -(1,4)-glycosidic bonds between galacturonic monomers by hydrolytic cleavage releasing *mono*-, *di*-, *oligo* galacturonates. These enzymes have been divided into four groups; those acts on pectate were called polygalacturonases (PG), while those preferentially hydrolyze pectin were called polymethylgalacturonases (PMG). In connection with their mode of action on the galacturonic acid backbone, the prefixes *endo*- and *exo*- used (Alkorta *et al.*, 1998). It is a member of glycosyl-hydrolases family 28 (Pedrolli *et al.*, 2009). These are one of the most studied and widely used pectinases which has high potential in commercial applications especially in fruit juice industry.

Bacillus strains are inferred to be the potent sources of *exo*-PG (Nagel & Vaughn, 1961; Kobayashi *et al.*, 2001; Jahan *et al.*, 2017). The ability of *B. polymyxa* in PG production was reported in 1961 (Nagel & Vaughn, 1961). A number of *Bacillus* species including *B. subtilis*, *B. pumilus* and *B. licheniformis* were reported to have PG production ability. A novel, alkaline and thermostable PG produced from an environmental isolate *Bacillus* sp. MG-cp-2 was reported by Kapoor *et al.*, in 2000 and its application in degumming of ramie and sunn hemp bast fibres was also revealed in the next year (Kapoor *et al.*, 2000; Kapoor *et al.*, 2001). Further, the ability of this isolate to produce PG by both SmF and SSF was also confirmed (Kapoor & Kuhad, 2002). Reports regarding a high molecular weight (115000 Da) alkaline *exo*-PG in a culture of *Bacillus* sp. strain KSM-P576 was publicized in 2001 (Kobayashi *et al.*, 2001). The ability of *B. sphaericus* for PG production was initially reported by Jayani *et al.*, 2010 (Jayani *et al.*, 2010).

Among several bacterial strains isolated from soil and rotten vegetables, the strain which produced maximum PG was identified as *B. licheniformis* KIBGE IB-2 (Rehman *et al.*, 2012). Also, the PG produced from this isolate was immobilized in chitosan for continuous degradation of pectin polymers (Rehman *et al.*, 2014). Further, properties of the enzyme were characterized and reported that the PG is having thermal stability and is able to perform its catalytic activity in a diversified environment (Rehman *et al.*, 2015). Pectinolytic (PG) properties of a new soil isolate *B. subtilis* C4 was reported in 2014 by Kusuma and reddy (Kusuma & Reddy, 2014a). Further, after purification and characterization of the enzyme,

they suggested that there was a threefold increase in the specific activity of PG produced by this isolate (Kusuma & Reddy, 2014b). PMG activity from *Bacillus* sp. strain BR1390, a novel environmental isolate was presented in the same year (Rastegari & Karbalaei, 2014).

Bacillus spp. were also reported for endo PG production. Endo-PG, exo-PG and PNL activities of five *Bacillus* strains isolated from decaying vegetable material was studied by Sores *et al.*, 2001 and these enzymatic solutions resulted in maximal reduction of the solution of citrus pectin viscosity, between 80 and 97 % (Soares *et al.*, 2001). The molecular weight of endo-PG produced by *B. subtilis* was found around the range of 67 kDa (Munir & Haidri, 2015).

Lyases/trans-eliminases: Lyases or trans-eliminases cleave α (1, 4)-glycosidic linkages by transelimination resulting in galacturonide with a double bond between C-4 and C-5 at the non-reducing end. They split the glycosidic bonds of either pectate (endo- and exopolygalacturonate lyase -PGL) or pectin (polymethylgalacturonate lyase - PMGL). Up to the recent times, all described pectin lyases were endo-PGL (Alkorta *et al.*, 1998). The first report of an exo-pectin lyase was in 2014, in the fungus *A. giganteus* (Pedrolli & Carmona, 2014). But the enzyme has not yet been reported in *Bacillus* sp. These enzymes are classified into the polysaccharide lyase family 1 (CAZy database). PGL has an absolute requirement of Ca²⁺ ions whereas PMGL requires Ca²⁺ and other cations only for its stimulation (Jayani *et al.*, 2005).

Production, applications, biochemical and molecular characteristics of lyases acting on pectic polymers have been reviewed previously (Dubey *et al.*, 2016). Kelly and Fogarty, 1978 reported PL production from an isolated soil bacterium *Bacillus* sp. RK9 and noted that hydrolysis of substrate occurred in a random fashion and the enzyme was 50% more active towards acid soluble pectic acid than towards sodium polypectate (Kelly & Fogarty, 1978). *B. stearothermophilus* with pectinolytic activity has been isolated by Karbassi and Vaughn, 1980 and found that it is producing a considerable amount of endo-polygalacturonic acid *trans*-eliminase (endo-PATE) (Karbassi & Vaughn, 1980). They also noted that enzyme produced by this organism was much more heat stable than previously reported similar enzymes from the mesophilic *B. polymyxa* (Nagel & Vaughn, 1961) and the thermotolerant *B. pumilus* (Dave *et al.*, 1976).

Details regarding a bacterial strain (*Bacillus* sp. PN33) producing large amounts of extracellular PNL was revealed in 1998 where, the maximum activity was found at acidic pH of 6 and is an unusual example for bacterial PNL (Kim *et al.*, 1998). Another high yielding pectinase strain, *Bacillus* sp. DT7 producing alkalothermophilic PNL with a shorter incubation period of

24 hr was reported in 2000 (Kashyap *et al.*, 2000). A low-molecular-weight (20,300 Da), high-alkaline PL was found in an alkaline culture of *Bacillus* sp. strain KSM-P15 by Kobayashi *et al.*, 1999 and suggested that this may be a novel enzyme that belongs to a new family (Kobayashi *et al.*, 1999). Later, it was also reported that the strain KSM-P15 produces high alkaline PL with high molecular weight (70,000 Da) (Ogawa *et al.*, 2000).

A thermophilic bacterial strain of *Bacillus* sp. with endo-PL activity has been isolated by Tako *et al.*, 2000 and noted that it had PPase activity, besides PL activity on lemon protopectin and cotton fibers (Takao *et al.*, 2000). A novel alkalophilic strain of *B. pumilus* BK2 producing a new type of extracellular endo- PL with high pl and a high pH optimum was reported in 2006 (Klug-Santner *et al.*, 2006). The PL producing ability tested from a group of six *Bacillus* species (*B. subtilis*, *B. pumilus*, *B. sphaericus*, *B. cereus*, *B. thuringiensis*, and *B. fusiformis*) isolated from cocoa fermentation, it was revealed that *B. fusiformis*, *B. subtilis*, and *B. pumilus* species were the best PL producers compared to other species (Ouattara *et al.*, 2011).

Esterases: The major component of pectic polysaccharides is homogalacturonan (HG), constituting about 65% of the pectin and contains (1→4) linked α -D-galacturonic acids (1,4-d-GalpA) which can be acetylated or methyl esterified, called smooth regions of pectin. The group pectin esterases comprises of pectin methylesterase (PME) and pectin acetyl esterase (PAE). The enzyme PME catalyzes reactions according to the double-displacement mechanism and its types, mode of action, structure etc. were reviewed previously (Kohli *et al.*, 2015) whereas PAE hydrolyzes the acetyl ester from the HG region of pectin forming pectic acid and acetate (Remoroza *et al.*, 2014). These enzymes act before the action of PG and PL which need non-esterified substrates (Kashyap *et al.*, 2001). These are a well-studied group of enzymes, which belong to carbohydrate esterase (CE) family 8 of CAZy database (Remoroza *et al.*, 2015).

The PME gene of *E. crysanthemii* B374 was successfully cloned in *B. subtilis* in 1991 (Heikinheimo *et al.*, 1991) and its purification and characterization were also reported (Pitkanen *et al.*, 1992). A recombinant acetyl esterase from *B. licheniformis* DSM13 was purified and biochemically characterized by Remoroza *et al.*, 2014 and reported to deacetylates a wide range of acetyl-rich pectins (Remoroza *et al.*, 2014). Later they also reported PME from the same organism that de-methylesterifies lemon pectin and sugar beet pectin in a stepwise manner (Remoroza *et al.*, 2015).

Protopectinases: *Bacillus* sp. was also reported to produce PPase, an enzyme that liberates water-soluble pec-

tic substances by restricted hydrolysis from water-insoluble protopectin in plant cell walls (Sakai *et al.*, 1989; Sakai & Sakamoto, 1990; Sakamoto *et al.*, 1994; Takao *et al.*, 2000). There are two types of protopetinases based on their reaction mechanism; A-type and B-type. A-type PPases react with the polygalacturonic acid region of protopectin (inner site) and the B-type PPases react with the polysaccharide chains that may connect the polygalacturonic acid chain and cell wall constituents (outer site) (Sakai *et al.*, 1993). A-type PPases are again classified into PPase-F, -L and -S based on the organism they isolated and all these types are having an approximate molecular weight of 30 kDa. B-type PPases are also in different types such as PPase-B, -C and -T. PPase-B, -C and -T have molecular weights of 45, 30, and 55 kDa, respectively. PPase-B and -C have an isoelectric point (pI) of around 9.0 whereas PPase-T has a pI of 8.1 (Gundala & Chinthala, 2017).

Saki *et al.*, (1989) used some strains belonging to the genus *Bacillus* such as; *B. subtilis*, *B. amyloliquefaciens*, *B. cereus*, *B. circulans*, *B. coagulans*, *B. firmus*, *B. licheniformis*, *B. macerans*, and *B. pumilus*, to study the production of PPase and concluded that *B. subtilis* IFO 12113 produce a new type of 'PPase -B' in respect to its reaction mechanism of the solubilization of protopectin, since it does not catalyze the degradation of polygalacturonic acid (Sakai *et al.*, 1989). Further, they also reported 'PPase-C' (does not react with rabbit antiserum against PPase -B) from *B. subtilis* IFO 3134, (Sakai & Sakamoto, 1990). Later, from the culture filtrate of *B. subtilis* IFO 3134 Sakamoto *et al.*, 1994 discovered two pectinolytic enzymes PPase-N and PPase-R. But, according to their substrate specificities and modes of action, these could be respectively classified as endo-PL and endo - PNL (Sakamoto *et al.*, 1994).

The major strains of *Bacillus* spp. that has been reported recently for pectinase production with their characteristic temperature and pH are shown in Table 1.

Shorter fermentation period for enzyme production: The time required for the production of pectinase by *Bacillus* spp. are remarkably less when compared to other microbial pectinases reported in the literature (Kashyap *et al.*, 2000). Literature review indicates that maximum pectinase production through many strains of *Bacillus* can be attained within 48 h of fermentation period. The optimum condition for pectinase production by *B. firmus* was recorded at a fermentation time for 18 h and the enzyme production was declined after 30 h (Roosdiana *et al.*, 2013). Whereas, pectinase production by *B. subtilis* SS started after 18 h of incubation and the production reached highest at 24 h (Ahlawat *et al.*, 2009). Alkaline pectinase production by *B. subtilis* WSHB04-02 showed an optimum fermentation time of

25 h (Wang *et al.*, 2007). Maximum pectinase production by *B. pumilus* dcsr1 was attained in 30 h through SmF using fermenter (Sharma & Satyanarayana, 2006) while enzyme production by *B. gibsonii* S-2 reached highest in 48h (at 35 °C) through SSF (Li *et al.*, 2005).

Economical enzyme production utilizing agro-wastes as substrates: Pectinase production from *Bacillus* spp. can be also achieved economically by utilizing different agro-waste as substrate. It is proved that orange peel waste can be used as a sole carbon source for pectinase production by various strains of *Bacillus* (Kapoor *et al.*, 2000; Embaby *et al.*, 2014; Tepe & Dursun, 2014; Kaur & Gupta, 2017). Wheat bran is another substrate, proven as a cheap and easily available source throughout the year for higher pectinase production and many researchers used wheat bran as an economical carbon source for pectinase production by *Bacillus* spp. Among various agro-byproducts studied for PG production by *Bacillus* sp. MG-cp-2, it is found that PG production level was boosted significantly by using wheat bran and ramie fibre in the production media (Kapoor *et al.*, 2000). *Bacillus* strains cultivated on wheat bran produced endo-PG, exo-PG and PNL in the crude enzymatic solution (Soares *et al.*, 2001). A high yield of pectinase (PG) was attained from *B. licheniformis* KIBGE IB-21 (Rehman *et al.*, 2012) and *B. licheniformis* KIBE-IB3 using wheat bran as substrate (Jahan *et al.*, 2017). Various other agro-industrial wastes such as; rice bran, cassava bagasse, sugar beet pulp, carrot peels etc. are also exploited for pectinase production by *Bacillus* spp. (Ghazala *et al.*, 2015; Nawawi *et al.*, 2017).

Mode of fermentation by either SmF or SSF: Pectinase production using *Bacillus* sp. can be attained by either SSF, SmF or by both. High quantities of PG is produced by *Bacillus* strains cultivated by SmF and semi-SSF (Soares *et al.*, 1999). Multiple pectinase enzymes activities such as endo-PG, exo-PG and PNL was showed by SSF culture filtrate of *Bacillus* strains (Soares *et al.*, 2001). Improved PG from *Bacillus* sp.MG-cp-2 under SmF and SSF and effect of amino acids and their analogues, vitamins and surfactants in fermentation was reported in 2002 (Kapoor & Kuhad, 2002). Medium optimization was found to effective in both SSF and Smf for pectinase production using *Bacillus* strains (Ghazala *et al.*, 2015; Bibi *et al.*, 2016; Kaur & Gupta, 2017). Zou *et al.*, 2014 developed a new Fed-Batch fermentation for enhanced production of alkaline PGL using *B. subtilis* 7-3-3. The process combines the enzymatic pretreatment of the carbon source with controlled pH of the fermentative broth to enhance enzyme production in a cheap manner (Zou *et al.*, 2014).

Cloning and expression of pectinase gene: Genetic information regarding pectinase genes of many *Bacillus*

Table 1. Recently reported *Bacillus* spp. producing different types of pectinase

No.	<i>Bacillus</i> species	Type of pectinase	Characteristics		References
			pH	Temp (°C)	
1	<i>B. subtilis</i>	Pectinase	8.0	50	(Torimiro & Okonji, 2013)
		Endo -PG	5	60	(Munir & Haidri, 2015)
2	<i>B. subtilis</i> KSM-P358	Exo -PG	8	55	(Sawada <i>et al.</i> , 2001)
3	<i>B. subtilis</i> EFRL 01	PG	8	45	(Qureshi <i>et al.</i> , 2012)
4	<i>B. subtilis</i> (TCCC11286)	PL	9	50	(Liu <i>et al.</i> , 2012)
5	<i>B. subtilis</i> 168	PL	9.5	50	(Zhang <i>et al.</i> , 2013)
6	<i>B. subtilis</i> C4	PG	9	60	(Kusuma & Reddy, 2014b)
7	<i>B. subtilis</i> 7-3-3	PGL	6.5	34	(Zou <i>et al.</i> , 2014)
8	<i>B. subtilis</i> BKDS1	PG, PNL, PL	8	40	(Kavuthodi <i>et al.</i> , 2015)
9	<i>B. subtilis</i> AD11	Pectinase	8.42	30	(Nawawi <i>et al.</i> , 2017)
10	<i>B. subtilis</i> SAV-21	Pectinase, PNL			(Kaur & Gupta, 2017)
11	<i>B. subtilis</i> Btk27	Pectinase	7.5	50	(Oumer & Abate, 2017)
12	<i>Bacillus</i> sp. N16-5	PL	11.5	50	(Li <i>et al.</i> , 2010)
13	<i>Bacillus</i> sp. SMIA-2		10	60-70	(Andrade <i>et al.</i> , 2011)
14	<i>Bacillus</i> sp. strain BR1390	PMG	6	60	(Rastegari & Karbalaei, 2014)
15	<i>Bacillus</i> sp. ZGL14	pectinase	8.6	50	(Yu <i>et al.</i> , 2017)
16	<i>Bacillus</i> sp ZJ1407	pectinase	5	37	(Yu & Xu, 2018)
17	<i>B. pumilus</i> (NRRL B-212)	Exo-pectinase	8	30	(Tepe & Dursun, 2014)
18	<i>B. pumilus</i>	Exo-pectinase	8	30	(Tepe & Dursun, 2014)
19	<i>B. pumilus</i> (ATCC 7061)	PL	8	65	(Liang <i>et al.</i> , 2015)
20	<i>B. licheniformis</i> KIBGE IB-21	PG	7	37	(Rehman <i>et al.</i> , 2012)
			8-10	45	(Rehman <i>et al.</i> , 2015)
21	<i>B. licheniformis</i> SHG10	PG	8	37.8	(Embaby <i>et al.</i> , 2014)
22	<i>B. licheniformis</i> DSM-13	PAE	8	50	(Remoroza <i>et al.</i> , 2014)
		PME	8	50	(Remoroza <i>et al.</i> , 2015)
23	<i>B. licheniformis</i> KIBGE IB-3	PG	7	37	(Jahan <i>et al.</i> , 2017)
24	<i>B. stearothermophilus</i>	Pectinase	7.5	60	(Torimiro & Okonji, 2013)
25	<i>B. cereus</i>	Pectinase	8.5	37	(Namasivayam <i>et al.</i> , 2011)
26	<i>B. cereus</i>	Pectinase	8	50	(Torimiro & Okonji, 2013)
27	<i>B. sphaericus</i> (MTCC 7542)	PG	6.8	30	(Jayani <i>et al.</i> , 2010)
28	<i>B. megaterium</i> AK2	PL	8.5	50	(Mukhopadhyay <i>et al.</i> , 2012)
29	<i>B. clausii</i>	PNL	10	60	(Li <i>et al.</i> , 2012)
30	<i>B. firmus</i>	PG	7	50	(Roosdiana <i>et al.</i> , 2013)
31	<i>B. halodurans</i> M29	Pectinase	10	80	(Mei <i>et al.</i> , 2013)
32	<i>B. tequilensis</i> SV11	PL	9	60	(Chiliveri & Linga, 2014)
33	<i>B. mojavensis</i> I4	Pectinase	8	60	(Ghazala <i>et al.</i> , 2015)
34	<i>B. vallismortis</i> (JQ990307)	PG			(Sohail & Latif, 2016)

spp. are available in various nucleotide sequence databases. Thus it supports successful cloning and expression of pectinase gene in other organisms. The most cloned pectinase gene from *Bacillus* sp. is PL gene '*pel*'. The first cloning and characterization of a *pel* gene from the *Bacillus* genus was reported in *B. subtilis*. The cloned

gene indicated a 1,260 bp open reading frame (ORF) encoding a 420 amino acid polypeptide which includes a 21 amino acid (aa) signal sequence. Molecular weight is found to be 45,605 Da and the purified enzyme had similar properties to the PL isolated from extracellular media of the organism (Nasser *et al.*, 1993). Since

then many researchers reported the successful cloning, expression, sequencing and characterization of pectinase genes (especially PL) from several *Bacillus* strains (Li *et al.*, 2010; Dubey *et al.*, 2016).

The gene encoding the PNL of *B. subtilis* IFO 3134, has been cloned, sequenced, and characterized in 1996 and it consists of a coding sequence of 345 aa and expression of the PNL in *E. coli* was also reported (Sakamoto *et al.*, 1996). The gene encoding the exo-PG (pehK) of *Bacillus* isolate was cloned and characterized in 2001. The cloned gene revealed a 2940 ORF consist of 980 aa (signal sequence -27 aa and mature protein -953 aa) (Sawada *et al.*, 2001). A highly alkaline thermostable pectinase from *B. halodurans* M29 was cloned and expressed in *E. coli*. The expressed enzyme showed high thermostability and long half-life and only 54 % sequence similarity to known enzymes and thus considered novel (Mei *et al.*, 2013). The gene encoding *Bacillus* PAE was also cloned and expressed in *E. coli* (Bolvig *et al.*, 2003; Remoroza *et al.*, 2014).

PRESENT SCENARIO

The reports on pectinase production by *Bacillus* spp. are still continuing with latest findings. Kavuthodi *et al.*, 2015 isolated and identified the most potent pectinolytic bacterial strain as *B. subtilis* BKDS1 and found to produce pectinases (PG, PL & PNL) also, coproduction of biosurfactant along with pectinase was reported (Kavuthodi *et al.*, 2015). Pectinase production by *B. mojavensis* I4 using carrot peels and its application in sesame seeds oil extraction was reported in 2016 (Ghazala *et al.*, 2015). In the same year, it was reported that among different microbial species screened for pectinase production, most prominent pectinase producing isolates were *Bacillus* sp. and identified as *B. firmus*, *B. coagulans*, *B. endophyticus* and *B. vietnamsensis* (Aaisha & Barate, 2016). Extracellular pectinase production and its purification from a new strain of isolated *B. subtilis* were also published in 2016 (Mercimek Takcı & Turkmen, 2016).

The results of a recent study indicated that out of 20 isolates screened for xylanopectinolytic enzyme activity, the most prominent strain was identified as *B. subtilis* ADI1 (Nawawi *et al.*, 2017). Also, pectinolytic enzyme production by *B. subtilis* SAV-21 was also reported in 2017 (Kaur & Gupta, 2017). The study conducted by Jahan *et al.*, in 2017 revealed that *B. licheniformis* KIBGE IB-3 has potential to produce a high amount of pectinase by utilizing different agro-wastes (Jahan *et al.*, 2017). Thermo acidic pectinase production from *Bacillus* sp. ZJ1407 has a good acidic and thermal stability within a pH range of 3.0-5.0 and at 80-90 °C (Yu & Xu, 2018).

CONCLUSION

As revealed in this review, *Bacillus* spp. are highly efficient for pectinase production mainly by alkaline pectinase and upholds their position as dominant pectinase producer among bacterial genera. They have proved to produce almost all classes of pectinase enzymes based on the substrate provided. It is quite evident from the literature that *B. subtilis* is the leading bacterium reported to have pectinolytic property followed by *B. pumilus*. So, considering the importance of alkaline pectinase, new bacterial strains especially from the genus *Bacillus* need to be identified for the industrial production of pectinase.

Conflict of Interests: There are no conflicts of interest.

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