

A Comprehensive review on the properties and applications of extremozymes from extremophilic actinobacteria

Abstract

Actinobacteria are ubiquitous microorganisms found in soil, plants tissues and aquatic environments. They have an exceptional potential for producing secondary metabolites of pharmaceutical interest. In addition, they are good producers of enzymes effective in breaking down various organic substances. In recent years, the number of studies on enzymes produced by extremophilic actinobacteria has increased considerably thanks to their unusual mechanisms of action, their stability and resistance to denaturation as well as structural novelties allowing them to be promising and efficient for wide applications in many products and processes particularly in detergent, food and beverage, animal feed, leather, textile, pulp and paper, biofuel, cosmetic and pharmaceutical industries. This review summarizes the latest experimental data on extremozymes, produced by extremophilic actinobacteria. It focuses on the description of the producing actinobacteria, their biochemical and physicochemical properties and their potential industrial applications for the development of a bioeconomy, with particular emphasis on thermophilic, psychrophilic, acidophilic, alkaliphilic and halophilic hydrolases such as amylases, cellulases, xylanases, pectinases, chitinases, proteases, lipases and phospholipases.

Keywords: extremophilic actinobacteria, green biocatalysts, biobased economy, extremozymes, biochemical properties, industrial applications

Volume 8 Issue 1 - 2024

Afoua Gorrab,¹ Rania Ouertani,¹ Amal Souii,¹ Fatma Kallel,² Ahmed Slaheddine Masmoudi,¹ Ameer Cherif,¹ Mohamed Neifar^{2,3}

¹LBVBGR-LR11ES31, Higher Institute of Biotechnology of Sidi Thabet (ISBST), University of Manouba, Tunisia

²LAPVA-LR16ES20, National Engineering School of Sfax (ENIS), University of Sfax, Tunisia

³Common Services Unit "Bioreactor Coupled with an Ultrafilter", ENIS, University of Sfax, Tunisia

Correspondence: Mohamed Neifar, Laboratory of Plant Improvement and Agro-Resources Valorization (LAPVA-LR16ES20), National Engineering School of Sfax, University of Sfax, Road of Soukra km 4, Sfax, 3038 Tunisia, Tel 21628762783, Email mohamed.naifar@gmail.com

Received: August 19, 2024 | **Published:** September 6, 2024

Introduction

The global industrial enzymes market is estimated to be worth US\$5.6 billion in 2019, and this figure is expected to enhance at an average annual growth rate of approximately 6.4% from 2020 to 2027.¹ Enzymes are used as active ingredients, additives or manufacturing aids in many industrial sectors. Enzyme based technologies have been developed to meet the global needs of the detergent, leather, textile, energy, food and feed markets. One of the recent green and sustainable approaches to achieve catalysis in harsh industrial conditions is to use microbial extremozymes.²⁻⁴ Indeed, extremophile microorganisms (thermophiles, psychrophiles, acidophiles, alkaliphiles, halophiles, barophiles, metallophiles, xerophiles, radiophiles, metallophiles, etc.) have abilities to colonize the most hostile environments such as volcanic zones, hydrothermal sources, hypersaline and alkaline lakes, desert sediments, cold oceans, and industrial effluents. They develop several strategies to overcome several biotic stresses caused by diverse pathogens.³ They are also able to survive unusual conditions of temperatures, salinity, pressure, heavy metals, radiation, acidic or basic pH, drought, cold or a combination of different abiotic stresses (polyextremophiles).⁵⁻⁷

Actinobacteria play a vital role in many biological processes including biogeochemical cycles, bioweathering, bioremediation and plant growth promotion.⁸⁻¹⁰ They are widely recognized for their metabolic versatility and resistance to hazardous environmental conditions. Actinobacteria produce a broad spectrum of bioactive metabolites with high commercial value, e.g. pharmaceuticals, nutraceuticals, antitumor agents and plant growth regulators, making them a prime target for the discovery of novel bioactive compounds.¹¹⁻¹³ They are also a great source of several industrial enzymes, such as proteases, lipases, cellulases, amylases, pectinases, chitinases, among others.¹⁴⁻¹⁹ In recent years, with continued advances in sequencing

technology and bioinformatics tools, numerous studies have focused on exploring actinobacteria that thrive in extreme conditions to unearth the specific characteristics of their extremozymes.³ Their unique structural features and adaptations allow them to resist against denaturation at extremely conditions/environments, and efficiently catalyze reactions in organic solvents and unconventional media.²⁰⁻²⁶ This review summarizes, based on representative examples, the main extremozymes produced by extremophilic actinobacteria, focusing on their bio-physico-chemical properties and their industrial uses.

Biological and chemical diversity of actinobacteria

Actinobacteria represent one of the oldest phyla in the Bacteria domain.¹³ They have quite variable morphologies ranging from cocci to branched filaments. Members of this phylum are characterized by their high GC content ranging from 42% (*Gardnerella vaginalis*) to 74.4% (*Kineococcus radiotolerans*).^{27,28} The Actinobacteria phylum currently includes six classes, namely Actinobacteria, Acidimicrobiia, Coriobacteriia, Nitrospirales, Rubrobacteria and Thermoleophilia²⁹ (Figure 1). The sixteen orders of the class Actinobacteria are: Actinomycetales, Bifidobacteriales, Catenulisporales, Corynebacteriales, Cryptosporangiales, Frankiales, Glycomycetales, Jiangellales, Kineosporiales, Micrococcales, Micromonosporales, Propionibacteriales, Pseudonocardiales, Sporichthyales, Streptomycetales and Streptosporangiales.²⁷ The phylogeny of Actinobacteria remains controversial due, among other things, to the choice of datasets and phylogenetic methods.³⁰⁻³⁴ For example, according to the study by Sen et al.,³⁰ the order Micrococcales is subdivided into Micrococcales (*Kocuria*, *Rothia*, *Micrococcus*, *Arthrobacter*, *Tropheryma*, *Microbacterium*, *Leifsonia* and *Clavibacter*), Cellulomonales (*Beutenbergia*, *Cellulomonas*, *Xylanimonas*, *Jonesia* and *Sanguibacter*) and Brachyacteriales (*Brachyacterium*). Based on cell wall morphology and chemotype,

species of the genus *Streptomyces* of the family Streptomycetaceae are considered to be the most studied actinobacteria due to their interesting roles in ecology, industry and biotechnology.¹³

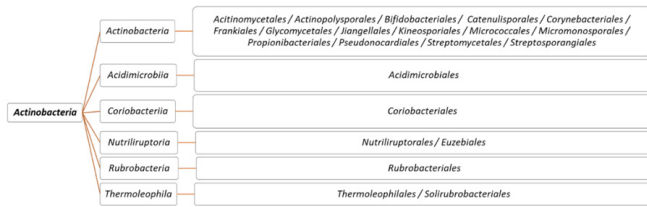


Figure 1 Proposed taxonomy for Actinobacteria in the forthcoming Bergey's Manual of Systematic Bacteriology.²⁹

Biotechnologically and industrially, actinobacteria are considered valuable bacteria since they are exploited for the production of biomolecules of interest (Figure 2). More than 10,000 bioactive molecules have been produced by actinobacteria, representing approximately 45% of all bioactive microbial metabolites discovered to date. They produce clinically useful antibiotics as well as other pharmaceutical products including anticancer agents, immunostimulants, immunosuppressants, antioxidants and antiparasitics.^{11–13,35–38}

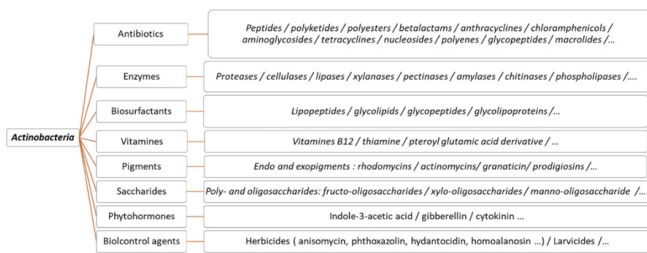


Figure 2 Important bioactive molecules produced by actinobacteria.^{11–13,35–38}

Actinobacteria are increasingly known for their capacity to produce many industrially important enzymes, such as ligninases, glycosyl hydrolases (cellulases, xylanase, amylases, etc.), proteases and lipases/esterases. These enzymes are used in the bioconversion of numerous agricultural and agro-industrial substrates and wastes into high added value bioproducts.^{14–19,39–44}

Strategies for discovering extremozymes producing extremophilic actinobacteria

Extremophilic microorganisms have the ability to live in various terrestrial and aquatic ecological niches under extreme conditions by adapting to ranges of temperature (55°C to 121°C and -2°C to 20°C), alkalinity or acidity (pH > 8, pH < 4), salinity (NaCl or KCl 2-5 M), pressure (>500 atmospheres), radiation (UV resistance > 600 J/m), heavy metals (arsenic, cadmium, copper, chromium, iron, nickel, lead, zinc and mercury), etc.³ These extremophiles are classified into thermophiles, psychrophiles, alkalophiles, acidophiles, halophiles, piezophiles, metalophiles, etc. depending on the extreme biota in which they grow and the abiotic environmental stressors they can resist and tolerate. The most representative actinobacterial genera for each extremophilic group were listed in figure 3. Extremophilic actinobacteria have developed several strategies of adaptation to extreme conditions such as antibiosis, variation of metabolic

modes (autotrophy, heterotrophy and saprobes) and the secretion of extremozymes stable at harsh conditions such as high temperature, alkalinity and salinity. Some actinobacteria are adapted to survive more than one type of extreme environment and are therefore called polyextremophiles.^{5–7}

Thermophilic actinobacteria	Psychrophilic actinobacteria	Alkaliphilic actinobacteria	Acidophilic Actinobacteria	Halophilic actinobacteria
<ul style="list-style-type: none"> • Thermoactinomyces • Thermomonospora • Microbispora • Saccharopolyspora • Saccharomonospora • Streptomyces • Geodermatophilus • ... 	<ul style="list-style-type: none"> • Streptomyces • Rhodococcus • Microbacterium • ... 	<ul style="list-style-type: none"> • Streptomyces • Micromonospora • Nocardoides • Microcella • Cellulomonas • Nesterenkonia • Streptosporangium • Corynebacterium • Georgania • Nocardopsis • Isapericcola • Nesterenkonia • ... 	<ul style="list-style-type: none"> • Streptomyces • Kitasatospora • ... 	<ul style="list-style-type: none"> • Micromonospora • Rhodococcus • Streptomyces • Dietzia • Salinispora • Marinophilus • Solwarospora • Salinibacterium • Aeromicrobium • Gardonia • Microbacterium • Mycobacterium • Nocardopsis • ...

Figure 3 Some representative genera of biotechnologically important extremophilic actinobacteria.

The classic approach to discover new extremozymes consists to cultivate microorganisms followed by a screening of the desired enzymes.^{27,45–47} Although many extremozymes with improved properties for industrial applications have been isolated from extremophiles using the classical culturable approach, approximately 99.9% of extremophilic actinobacteria (as for other microorganisms) cannot be grown using traditional laboratory techniques. Fortunately, metagenomic approaches have recently been developed to discover new extremozymes-encoding genes directly from uncultured microorganisms.³ Putative enzymes are identified on the basis of their conserved sequences or on the basis of their function and specific enzymatic activity. The latter is now the most frequently used technique to screen new extremozymes originating from extremophilic Actinobacteria. Furthermore, due to rapid advances in microbial genome sequencing technologies, increasing progress has been made in studying the diversity and biotechnological potential of actinobacteria. Indeed, the genomes of numerous terrestrial and marine extremophilic actinobacteria have been recently annotated, analyzed and evaluated, which provides valuable information related to these microbial extremophiles and their catalytic potential. Genomic annotation makes it possible to properly understand the mechanisms of adaptation of these extremophiles to extreme conditions and to identify the structural and catalytic specificities of their extremozymes. As an illustrated example, Figure 4 and 5 showed the genomic traits of the polyextremophilic actinobacterium *Microbacterium metallidurans* TL13 and the 3D structure of its thermostable heavy-metal reductases.⁴⁸

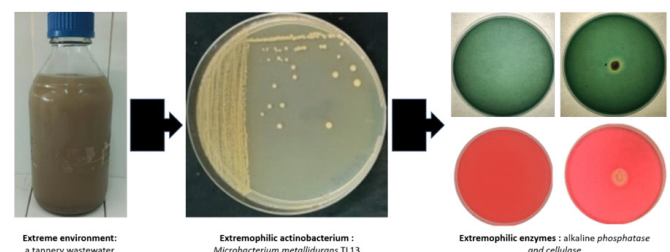


Figure 4 *Microbacterium metallidurans* TL13: an extremozyme-producing actinobacterium isolated from a tannery wastewater.

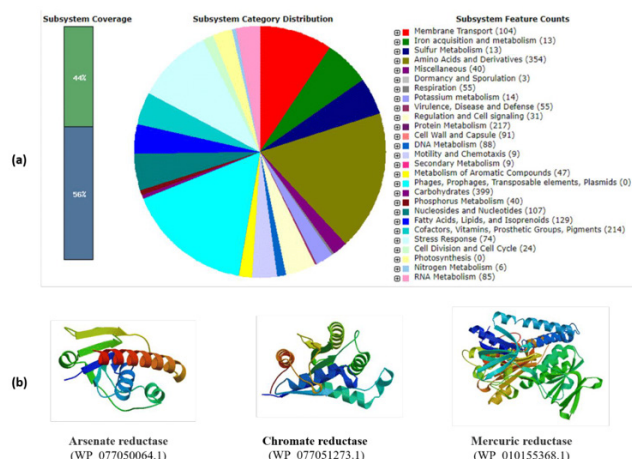


Figure 5 (a) RAST annotation summary of the genome of halo-thermo-alkali-tolerant, metallophilic plant growth promoting actinobacterium *Microbacterium metallidurans* TL13 (Taxonomy ID: 69370; genome size: 3,587,460 and GC Content: 70.7). The RAST annotation tool assigns names and functions to protein-coding genes via their subsystem technology. The green color represents features that are found in RAST subsystem. The blue color represents features not assigned to a subsystem. The draft genome sequence suggests potent metabolites candidates that are essential for survival under multiple environmental extreme conditions, such as high salinity, alkalinity and the presence of critical heavy metal concentrations.

(b) Swiss model 3D-structures of selected thermostable arsenate reductase (MM: 14.30 KDa; pl: 4.64; AI: 90.90), chromate reductase (MM: 20.20 KDa; pl: 4.69; AI: 92.00) and mercuric reductase (MM: 49.18 KDa; pl: 5.48; AI: 94.35) of the metallophilic strain TL13. The corresponding *in silico* properties were determined by ExPasy-Prot Param tool (<http://web.expasy.org/protparam/>). All proteins showed aliphatic index (AI) values of more than 90 indicated that they are highly thermo-stable.

Adaptation mechanisms and properties of actinobacterial extremozymes

The majority of enzymes currently on the market come from mesophilic micro-organisms, often inhibited by the extreme conditions of many industrial processes. Although these enzymes appear to perform well in certain industrial applications, the enzyme market remains insufficient to meet industrial demands, largely due to their poor stability under industrial conditions.⁴⁹ Industrial processes often require biocatalysts capable of withstanding a wide range of harsh conditions, including temperature, pH and salinity, while exhibiting high biocatalysis/bioconversion rate and reproducibility. Consequently, the enhanced stability and activity of extremozymes make them promising alternatives to ordinary processes conferring considerable economic potential in various industries. It's worth mentioning that microbial extremozymes use a variety of mechanisms to tolerate conditions. (Figure 6)

values below 3. They have acidic isoelectric points due to the abundance of acidic amino acids (aspartate, glutamate).⁵⁹ At very low pH, most surface amino acids are protonated, leading to a reduction in negative charges and protein stabilization.

Halophilic enzymes are characterized by maximum activity and stability at high NaCl concentrations exceeding 1.5 M. Halophilic enzymes generally have an acidic isoelectric point with an abundance of acidic amino acids (aspartic and glutamic acids) on the surface of the enzyme.⁶⁰ These amino acids have high water retention allowing the protein to remain soluble in conditions of dehydration caused by high salinity. Halophilic enzymes also tolerate organic solvents since

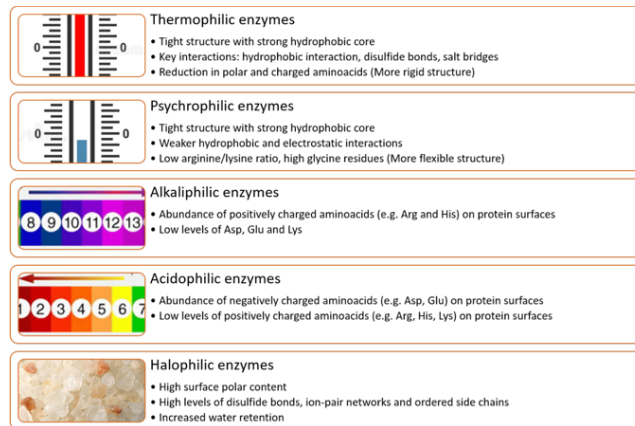


Figure 6 Protein adaptations and unique structural features of some extremozymes.^{49–62}

Thermophilic enzymes are characterized by maximum activity at temperatures between 50 and 125°C.⁵⁰ Disulfide bonds play a major role in thermal stability, due in part to the presence of disulfide bonds that reduce the entropy of the unfolded protein form;^{51–53} Other factors involved in maintaining the conformation of thermophilic extremozymes include the presence of loops and short helices, a good surface charge with internal hydrophobic amino acids.^{50,54} The three-dimensional structures of thermophilic enzymes are generally similar to those of their mesophilic counterparts, with some differences such as a greater presence of charged residues on their surface. Among the different types of extremozymes, thermophiles are the most studied scientifically and the ones that have attracted the most attention from manufacturers in view of their benefits for industrial processes. High temperatures increase the solubility of polymeric substrates, reduce the risk of contamination, promote faster reactions and improve solvent miscibility. Psychrophilic enzymes are active and stable at low temperatures between -20 and 10°C.⁵⁵ They preserve their enzymatic activities at low temperatures unlike mesophiles thanks to a low arginine/lysine ratio, little proline and plenty of glycine in the structure, increased surface hydrophobicity, few secondary structures, weaker interactions of hydrogen bonds, disulfide bridges and electrostatic interactions and.^{56,57} Psychrophilic enzymes improve the cost-effectiveness and efficiency of low-temperature industrial processes and reduce environmental impact.

Alkaliphilic enzymes have maximum activity and stability at pH values above 9. They have more alkaline isoelectric points than their neutrophilic counterparts, due to the abundance of basic amino acids such as arginine and histidine, and a reduction in acidic amino acids such as glutamate.⁴⁹ Arginine residues form ionic bonds with aspartate residues, enhancing enzyme stability under alkaline conditions.⁵⁸ Acidophilic enzymes have maximum activity and stability at pH

this lower water activity in the same way as high salinity. Another adaptive mechanism is their weak hydrophobic interactions which result in good protein flexibility at high NaCl concentrations.⁶¹ Structural analyzes revealed major differences between nonhalophilic and halophilic enzymes. The former contains higher percentages of protein surface serine and threonine and low hydrophobicity amino acids, aspartic and glutamic acids and lower percentages of basic amino acids such as lysine than enzymes non-halophilic. This favors the creation of a greater number of salt bridges and electrostatic interactions. In addition, negative surface charges are involved in protein solvation and prevention of protein denaturation and aggregation.⁴⁹

Polyextremophilic Enzymes are currently attractive for biotechnology because they resist to multiple harsh conditions, such as: pH, temperature and salinity.⁴⁹ The search for new unique polyextremophilic enzymes is a valuable activity for which extremophilic actinobacteria producers have been neglected. The structurally favored resistance of polyextremophilic enzymes is still poorly understood.⁶²

Industrially important hydrolytic actinobacterial extremozymes: *Streptomyces* species as model producers

A huge number of actinobacterial enzymes are being produced commercially and used in various industries (Figure 7).⁶³⁻⁶⁸ Some important extremophilic glycosyl hydrolases, lipase/esterases and proteases produced by actinobacteria particularly *Streptomyces* species are briefly described below (Table 1).

Amylases are enzymes that hydrolyze starch into glucose, maltose, and maltotriose units. Native or recombinant amylases are used in various industries such as the detergent industry, textile, leather, paper, food and feed industries and biofuel production. Enzymatic hydrolysis of starch produces simpler oligosaccharides and sugars

which find various industrial applications, some of which require a liquefaction or gelatinization step at high temperature (105-110°C) followed by saccharification at a lower temperature (55 to 60°C).⁶⁹

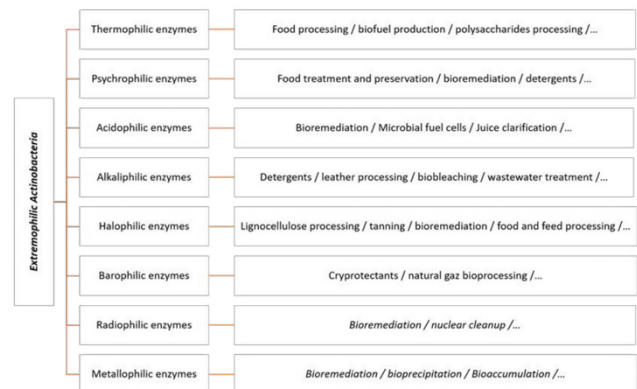


Figure 7 Actinobacterial Extremozymes and their potential applications.^{14-19,39-49}

Table 1 Properties and potential applications of various extremozymes produced by *Streptomyces* strains

Extremozymes	Producing strain	Enzymes properties	Potential applications	References
Extremophilic amylases	<i>S. fragilis</i> DA7-7	Thermo-halo-alkali-stable, retained activity in presence of detergents	Use in brewery, bakery, detergent, saccharification and starch liquefaction	69
	<i>Streptomyces</i> sp. TDI-10, <i>Streptomyces</i> sp. TDI-12 <i>Streptomyces</i> sp. TDI-13	Thermostable	Antifungal and bioleaching	70
	<i>Streptomyces</i> sp. No. 4	Thermo-alkali-stable	Hydrolyze of blue starch and production of glucose and maltose	71
	<i>Streptomyces</i> strain A3	Thermo-halo-alkali-stable, retained activity in presence of oxidants and detergents	Detergents, food and pharmaceutical industries	72
Extremophilic cellulases	<i>S. drozdowiczii</i> M7a	Thermostable, retained activity in the presence of detergents	Detergent and textile industries	73
	<i>Streptomyces</i> DSK59	Acido-thermostable	Hydrolyse of pretreated sorghum stover and liberation of reducing sugars	74
	<i>Streptomyces</i> sp. strain SirexAA-E (SACTE)	Thermostable	Deconstruction of lignocellulosic plant biomass into small polysaccharides and free sugars	75,76
	<i>S. griseorubens</i> LH-3	Thermostable	Biobleaching of eucalyptus kraft pulp Production of short-chain xylooligosaccharids	77
Extremophilic hemicellulases (xylanases and mannanases)	<i>S. ipomoea</i> CECT 3341	Thermo-alkalizable	Biobleaching of pine kraft pulps	78
	<i>Streptomyces galbus</i> NR	Thermoalkalizable	Liberation of reducing sugars and improvement of pulp bleachability	79
	<i>S. olivaceus</i> (MSU3)	Thermo-halo-alkalizable	Conversion of the pretreated agro-wastes into bioethanol	80
	<i>Streptomyces</i> sp. CS628	Alkali-thermostable	Biobleaching of kraft pulp and xylooligosaccharides production	81
	<i>S. chartreusis</i> LI 105	Alkali-thermostable	xylooligosaccharide production	82

Table I Continued..

Extremozymes	Producing strain	Enzymes properties	Potential applications	References
Extremophilic pectinases	<i>S. hydrogenans</i> YAM1	Thermostable	Production of active compound from fruit wastes and removal of fibrous pollutants from different environments	84
	<i>Streptomyces</i> sp. S27	Alkalistable	Use in alkaline industries such as textile	85
	<i>S. lydicus</i>	Thermostable and active at neutral pH	Banana fiber processing	86
	<i>S. halstedii</i> ATCC 10897	Thermostable	Juice clarification	87
	<i>S. chilikensis</i> RC1830	Thermostable	Production of chito-oligo-saccharides	88
Extremophilic chitinases	<i>St. venezuelae</i> P10	Thermal stability, pH tolerant, and antifungal properties	Potential use as biocontrol agents	89
	<i>Streptomyces mutabilis</i>	Thermostable	Synthesis of N-acetylD-glucosamine, in production of single cell protein (SCP) and for bioconversion of chitin wastes	90
	<i>S. violascens</i> NRRL B2700	Thermostable	Applications in industrial, medical and commercial fields	91
	<i>S. violaceusniger</i>	Thermostable	Antagonistic effect of this bacterium against wood rotting fungi	92
	<i>S. fungicidicus</i> MML1614	Alkali-thermostable	Leather, pharmaceutical detergents, food and brewing	93
Extremophilic proteases	<i>S. koyangensis</i> TN650	Alkaline, detergent-stable, solvent-tolerant	Application in detergent formulations and non-aqueous peptide biocatalysis	94
	<i>Streptomyces</i> sp. MAB18	Alkali-thermostable, antioxydant properties	Use as supplementary protein and antioxidant in the animal feed formulations	95
	<i>Streptomyces</i> sp. AB1	Alkali-thermostable, stability in organic solvents	Applications in detergent formulations, dehairing during leather processing, and non-aqueous peptide biocatalysis	96
	<i>Streptomyces</i> sp.	Alkali-thermostable	Useful for biotechnological process involving keratin hydrolysis or in the leather industry	97
	<i>S. gulbargensis</i>	Alkali-thermostable	Valorization of keratin-containing wastes or in the leather industry	98
	<i>Streptomyces</i> sp. DP2.	Alkali-thermostable	Production of protein hydrolysates, for the film industry, and in waste processing	101
	<i>Streptomyces</i> sp. OCI19-7	Organic solvent-tolerant	Biodiesel production	102
	<i>Streptomyces</i> sp. CS273	Alkaline, organic solvent-tolerant	Transesterification of waste cooking oil	103
	<i>S. violascens</i> OCI25-8	Alkali-thermostable	Oily wastewater treatment	104
	<i>S. pratensis</i> MVI	Organic solvent-tolerant	Enzymatic improvement of n6/n3 ratio in polyunsaturated fatty acids from fenugreek seed oil	105
Lipases/esterases	<i>Streptomyces</i> sp. WV007	Thermostable, organic solvent tolerant	Potential applications in industries for oil modification and production of partial glyceride	106
	<i>S. thermocarboxydus</i> ME168	Thermostable	Sugar ester synthesis	107
	<i>S. lividans</i> TK64	Alkali-thermostable	Synthesis of polyesters, optical prodrugs, and food additives	110
	<i>S. coelicolor</i> A3(2)	Cold-active, alcohol and organic solvent tolerant	Organic synthesis of short-chain esters such as flavors	112
	<i>S. cinnamoneum</i> SK43.003	Acido-thermostable	Conversion of phosphatidylcholine to phosphatidylserine	114
Extremophilic phospholipases	<i>S. olivochromogenes</i> CS528	Alkali-thermostable	Lipid industry	115
	<i>S. violaceoruber</i> AS-10	Thermostable	Food manufacturing processes, i.e. egg processing, baking processes, degumming of fats and oils and milk processing for cheese production	116

Extremophilic amylases, particularly thermostable ones, derived from thermophilic actinobacteria of the genus *Streptomyces*, are in high demand to reduce energy input and make the whole process more profitable and economical. Certain extremophilic actinobacteria have the capacity to produce active and stable amylases at alkaline pH, which are very useful for detergent formulations. Modern laundries prefer detergent formulations based on psychrophilic amylases to wash clothes at low temperatures to save energy.^{69–72}

Cellulases are enzymes that hydrolyze cellulose into glucose, cellobiose and cello-oligosaccharides. Cellulases are commonly used in various industries, including food and feed, textile and laundry, pulp and paper industries as well as the production of biomass-based biofuels.^{73–76} There is an increased demand for extremophilic cellulases to be exploited in food industries and biorefineries, where thermal processes are used for the bioconversion of cellulose-derived materials. Most cellulases from extremophilic actinobacteria have highly sought-after properties such as good thermostability, strong resistance to acids, high stability to alkalis and excellent stability to detergents. Cellulase from *S. drozdowiczii* showed enormous potential in the textile and detergent industries thanks to its high activity and stability even in the presence of commercial detergents.⁷³

The acidothermophilic endoglucanase from the extremophilic bacteria *Streptomyces* DSK59 hydrolyzes pretreated sorghum stalks more efficiently than fungal enzymes, thus releasing a higher level of total reducing sugars.⁷⁴ *Streptomyces* sp. strain SirexAA-E (SACTE), isolated from the pine-boring woodwasp *Sirex noctilio*⁷⁵ produced and secreted extremophilic cellulases that efficiently degrade and convert plant biomass into small polysaccharides and free sugars.⁷⁶

Hemicellulases are enzymes that hydrolyze hemicelluloses, such as xylans, xyloglucans, arabinoxylans, and glucomannans from plant biomass. The most influential hemicellulose degrading enzymes are the xylanases and mannanases. They are applied in various fields such as in animal feed, paper and pulp, waste treatment, baking, brewing, and biofuel industries. Xylanases with high stability against temperature, pH and salinity are widely produced by extremophilic actinobacteria.^{77–82} Extremophilic xylanases and mannanases from *S. griseorubens* LH-3, *S. ipomoea* CECT 3341 and *S. galbus* NR were successfully applied to biobleaching of kraft pulps,^{77–79} that of *S. olivaceus* (MSU3) was used for the hydrolysis of agricultural waste for subsequent production of bioethanol.⁸⁰ The extremophilic xylanases from *Streptomyces* sp. CS802 and *S. chartreusis* L1105 showed potential for the synthesis of xylooligosaccharide-type prebiotics.^{81,82}

Pectinases are a group of complex enzymes that degrade the complex pectin. They are used in various industrial processes such as the clarification of fruit juices, the production of functional foods and animal feed, the liquefaction and saccharification of plant biomass, the bio-scouring of cotton fiber, the fermentation of coffee and tea, the production of paper and bioethanol, the retting and degumming of textile fibers, and the treatment of pectic wastewaters. The demand of extremophilic pectinases are continuously increasing. Several researchers succeed in the purification, characterization and applications of many extremophilic pectinases from *Streptomyces* species.^{83,87}

Chitinases, enzymes that hydrolyze chitin, have attracted interest in numerous industrial applications given their effectiveness in the degradation of chitin in fungal walls and insect exoskeletons. Because of its catalytic potential, they have been applied for the production of

chitin oligomers with bioactivities as well as biological control agents against nematodes and phytopathogenic fungi. *Streptomyces* species are hyperproducers of highly extremophilic extracellular chitinases useful for biocontrol applications in sustainable agriculture.^{88–92}

Proteases or proteolytic enzymes catalyze proteolysis by hydrolysis of peptide bonds. The search for new proteases and their formulations is continuous as they have many applications in industries such as detergents, animal feed, and breweries. *Streptomyces* have proven to be good producers of proteases with great potential for industrial applications.^{93–102} Marine *S. fungicidicus* MML1614 produced thermostable alkaline protease and applied to remove blood stains more effectively than conventional detergents.⁹³ *S. koyangensis* TN650 produced a detergent-stable and solvent-tolerant alkaline protease, a potential candidate for future application in detergent formulations and non-aqueous peptide biocatalysis.⁹⁵ *Streptomyces* sp. MAB18 produced a protease derived from poultry waste that may be useful as a supplemental protein and antioxidant in animal feed formulations.⁹⁵

Streptomyces strains produced keratinolytic alkaline protease and are good candidates for in detergent formulations, hair removal during leather processing and non-aqueous peptides biocatalysis.^{96–99} Stable alkaline protease from soil isolates of *S. clavuligerus* strain Mit-1 and stable and thermostable alkaline protease from *Streptomyces* sp. DP2 isolates from dairy protein factories and slaughterhouses were characterized. Their alkaline stability and thermostability make them excellent candidates for multiple industrial applications.^{100,101}

Lipases or lipolytic enzymes are ubiquitous triacylglycerol hydrolases that catalyze hydrolysis reactions of triglycerides to glycerol and fatty acids but also esterification, interesterification, transesterification, acidolysis and aminolysis reactions in aqueous and organic media.^{102–110} Thanks to these multiple and versatile reactions, lipases find various industrial and biotechnological applications, covering different areas such as the production of detergents, foods, polymers, biofuels, fine chemicals, diagnostics and bioremediation. Recently, several studies have focused on the isolation, purification, characterization and applications of extremophilic lipases and esterases from *Streptomyces* species.^{102–113} Phospholipases are lipolytic enzymes that hydrolyze the ester bonds of phospholipids. Phospholipases find numerous applications in the food industry, particularly in the bakery, egg and dairy industries since they help reinforce emulsifying properties by acting on the phospholipids already present in food ingredients. Phospholipases D were easy to produce *Streptomyces* species, catalyze transesterification reactions and, more importantly, were stable under various storage conditions.^{114,115} The food enzyme phospholipase A2, produced with the genetically modified strain *S. violaceoruber* AS-10, was distributed on the market under the name PLA2 NAGASE. It was intended for use as an alternative to emulsifiers in many food manufacturing processes, namely the processing of eggs, pastry and bakery processes such as cakes, breads, the degumming of fats and oils and the processing of milk for the production of cheese.¹¹⁶

Finally, it should be noted that certain species of *Streptomyces* have been recently exploited as promising cell factories for the simultaneous production of numerous secondary metabolites and extracellular extremozymes. To further illustrate this point, we can cite the case of the strain *Streptomyces coelicolor* A3(2) which has an efficient enzymatic machinery to break down cellulose, hemicelluloses and lignin in plant biomasses. This polyextremophilic strain was also capable of producing more than 17 distinct families of active bioproducts such as actinorhodin, nogalamycin,

geosmin and coelimecin P1.^{117,118} A second example concerns the rhizospheric strain *Streptomyces* sp. CRPSP2-6A1 which was able to (i) completely metabolize lignocellulosic biomass thanks to the presence of cellulases (endoglucanase and exoglucanase, as well as glucosidases and galactosidases), xylanases (arabinofuranosidases, mannosidases, endohemicellulases, and carboxylesterases) and lignin-degrading peroxidases, (ii) catabolize chitin and chitosan contributing to the metabolism of fungal and insect biomass wastes and (iii) efficiently metabolize various harmful xenobiotic compounds using many other enzymes, such as haloalkane dehalogenase, alkanal monooxygenase, 4-nitrophenol 4-monooxygenase, biphenyl 2,3-dioxygenase, pentachlorophenol 4-monooxygenase, anthranilate 3-monooxygenase, limonene 1-2 monooxygenase, and steroid C26-monooxygenase. The availability of genome sequences of extremophilic *Streptomyces* strains not only helps in understanding the mechanism of cellulose degradation and stress resistance, but also provides information on secondary metabolic potentials important for the production of various classes of bioactive compounds.

Conclusion

The phylum Actinobacteria represents the largest taxonomic group among the microbial community. The bacteria of this phylum are promising sources for the production of bioactive pharmaceutical compounds but are also considered as biofactories of enzymes with interesting new properties such as great stability in extreme environmental conditions, good tolerance to solvents, remarkable specificity and selectivity of substrates, which open the way to various applications in textiles, tanneries, refineries, human and animal food, detergency, paper industry, pharmaceutical industry, etc. These extremozymes can be used in faster biocatalytic processes, more precise, more specific and more environmentally friendly. However, only a few of the characterized enzymes from actinobacteria in particular belonging to the genus *Streptomyces* have been studied for real industrial applications. Therefore, it will be interesting to focus in the future on species from other genera of the phylum Actinobacteria like *Microbacterium* species that produce hyperstable enzymes with concrete examples of applications of these extremozymes in various industrial processes. Moreover, the use of metagenomic approaches will help to better identify and characterize species not yet culturable and their genes encoding extremozymes. Furthermore, the discovery of genomic sequences of polyextremophilic actinobacteria should encourage biotechnologists and manufacturers to embark on the exploration and industrial exploitation of extremozymes hyperproducing-Actinobacteria. The cost of production of these green catalysts could decrease considerably by using cheap substrates or agro-industrial and agricultural wastes, applying the response surface methodology for cultivation process optimization and scaling-up bioreactor production. Developments in protein engineering and directed evolution technologies will enable adaptation and improvement of biocatalytic characteristics, which will increase the application of extremophile-derived enzymes in industry.

Acknowledgements

This study was supported by the Tunisian Ministry of Higher Education and Scientific Research.

Funding

None.

Conflicts of interest

The authors declare, that there is no conflict of interest.

References

1. Adiguzel G, Faiz O, Sisecioglu M, et al. A novel endo- β -1,4-xylanase from *Pediococcus acidilactici* GC25; Purification, characterization and application in clarification of fruit juices. *Int J Biol Macromol.* 2019;129:571–578.
2. Jorquera MA, Graether SP, Maruyama F. Bioprospecting and biotechnology of extremophiles. *Front Bioeng Biotechnol.* 2019;7:204.
3. Sysoev M, Grötzinger SW, Renn D, et al. Bioprospecting of novel extremozymes from prokaryotes-the advent of culture-independent methods. *Front Microbiol.* 2021;12:630013.
4. Escudero-Agudelo J, Martínez-Villalobos J, Arocha-Garza H, et al. Systematic bioprospection for cellulolytic actinomycetes in the Chihuahuan Desert: isolation and enzymatic profiling. *PeerJ.* 2023;11:e16119.
5. Karan R, Mathew S, Muhammad R, et al. Understanding high-salt and cold adaptation of a polyextremophilic enzyme. *Microorganisms.* 2020;8(10):1594.
6. Laye VJ, Solieva S, Voelz VA, et al. Effects of salinity and temperature on the flexibility and function of a polyextremophilic enzyme. *Int J Mol Sci.* 2022;23(24):15620.
7. Pasqualetti M, Gorrasi S, Giovannini V, et al. Polyextremophilic chitinolytic activity by a marine strain (IG119) of *Clonostachys rosea*. *Molecules.* 2022;27(3):688.
8. Doumbou CL, Salovey MKH, Crawford DL, et al. Actinomycetes, promising tools to control plant diseases and to promote plant growth. *Phytoprotection.* 2001;82(3):85–102.
9. Lacombe-Harvey MÈ, Brzezinski R, Beaulieu C. Chitinolytic functions in actinobacteria: ecology, enzymes, and evolution. *Appl Microbiol Biotechnol.* 2018;102(17):7219–7230.
10. Lewin GR, Carlos C, Chevrette MG, et al. Evolution and ecology of actinobacteria and their bioenergy applications. *Annu Rev Microbiol.* 2016;70:235–254.
11. Hong K, Gao A, Xie Q, et al. Actinomycetes for marine drug discovery isolation from mangrove soils and plants in China. *Mar Drugs.* 2009;7(1):24–44.
12. Sharma P, Dutta J, Thakur D. Chapter 21 - *Future prospects of actinobacteria in health and industry. New and future developments in microbial biotechnology and bioengineering.* Editor(s): Bhim Pratap Singh, Vijai Kumar Gupta, Ajit Kumar Passari. Elsevier. 2018:305–324.
13. Selim MSM, Abdelhamid SA, Mohamed SS. Secondary metabolites and biodiversity of actinomycetes. *J Genet Eng Biotechnol.* 2021;19(1):72.
14. Salwan R, Sharma V. Chapter 11 - *The role of actinobacteria in the production of industrial enzymes. New and future developments in microbial biotechnology and bioengineering.* Editor(s): Bhim Pratap Singh, Vijai Kumar Gupta, Ajit Kumar Passari, Elsevier. 2018 :165–177.
15. Walia A, Guleria S, Mehta P, et al. Microbial xylanases and their industrial application in pulp and paper biobleaching: a review. *3 Biotech.* 2017;7(1):11.
16. Leo VV, Asem D, Zothanpuia, et al. Chapter 13 - *Actinobacteria: a highly potent source for holocellulose degrading enzymes. New and future developments in microbial biotechnology and bioengineering.* Editor(s): Bhim Pratap Singh, Vijai Kumar Gupta, Ajit Kumar Passari, Elsevier. 2018:191–205.

17. Pizarro LF, Souza HF, Ferreira JS, et al. *Production of Lipase by Actinobacteria*. Dharumadurai D (Ed.), *Methods in Actinobacteriology*. Springer protocols handbooks. Humana, New York, NY. 2022:505–512.
18. Aguiar MS, Maldonado RR, Carvalho AL, et al. *Production of actinobacteria amylase by fermentation in solid state using residues of licuri palm (Syagrus coronata)*. D. Dharumadurai (Ed.). *Methods in Actinobacteriology*. Springer Protocols Handbooks. Humana, New York, NY. 2022:495–503.
19. Fernandes de Souza H, Aguiar Borges L, Dédalo Di Próspero Gonçalves V, et al. Recent advances in the application of xylanases in the food industry and production by actinobacteria: A review. *Food Res Int*. 2022;162(Pt B):112103.
20. Shivilata L, Satyanarayana T. Thermophilic and alkaliphilic actinobacteria: biology and potential applications. *Front Microbiol*. 2015;6:1014.
21. Kleeberg I, Welzel K, Vandenheuvel J, et al. Characterization of a new extracellular hydrolase from *thermobifida fusca* degrading aliphatic-aromatic copolyesters. *Biomacromol*. 2005;6(1):262–270.
22. Chen S, Su L, Billig S, et al. Biochemical characterization of the cutinases from *thermobifida fusca*. *J Mol Catal B Enzym*. 2010;63(3–4):121–127.
23. Ugur A, Boran R. Production and characterization of a cold-active and n-hexane activated lipase from a newly isolated *serratia grimesii* RB06-22. *Biocatal Biotransform*. 2014;32(4):222–230.
24. Huang YC, Chen GH, Chen YF, et al. Heterologous expression of thermostable acetylxyylan esterase gene from *thermobifida fusca* and its synergistic action with xylanase for the production of xylooligosaccharides. *Biochem Biophys Res Commun*. 2010;400(4):718–723.
25. Zhang F, Hu S-N, Chena J-J, et al. Purification and partial characterisation of a thermostable xylanase from salt-tolerant *thermobifida halotolerans* YIM 90462^T. *Process Biochem*. 2012;47(2):225–228.
26. Hui ML, Tan LT, Letchumanan V, et al. The extremophilic actinobacteria: from microbes to medicine. *Antibiotics (Basel)*. 2021;10(6):682.
27. Nouioui I, Carro L, Garcia-Lopez M, et al. Genome-based taxonomic classification of the phylum actinobacteria. *Front Microbiol*. 2018;9:2007.
28. Verma M, Lal D, Kaur J, et al. Phylogenetic analyses of phylum actinobacteria based on whole genome sequences. *Res Microbiol*. 2013;164(7):718–728.
29. Ludwig W, Euzeby J, Schumann P, et al. *Whitman road map of the actinobacteria*. Bergey's Manual of Systematic Bacteriology. vol. 5, Springer, New York, NY. 2012:1–28
30. Sen A, Daubin V, Abrouk D, et al. Phylogeny of the class actinobacteria revisited in the light of complete genomes. the orders 'frankiales' and micrococcales should be split into coherent entities: proposal of frankiales ord. nov., geodermatophilales ord. nov., acidothermales ord. nov. and nakamurellales ord. nov. *Int J Syst Evol Microbiol*. 2014;64(Pt 11):3821–3832.
31. Zhi XY, Li WJ, Stackebrandt E. An update of the structure and 16S rRNA gene sequence-based definition of higher ranks of the class actinobacteria, with the proposal of two new suborders and four new families and emended descriptions of the existing higher taxa. *Int J Syst Evol Microbiol*. 2009;59(Pt 3):589–608.
32. Oren A, Garrity GM. Notification of changes in taxonomic opinion previously published outside the IJSEM. *Int J Syst Evol Microbiol*. 2019;69(1):13–32.
33. Gao B, Gupta RS. Phylogenetic framework and molecular signatures for the main clades of the phylum actinobacteria. *Microbiol Mol Biol Rev*. 2012;76(1):66–112.
34. Barka EA, Vatsa P, Sanchez L, et al. Taxonomy, physiology, and natural products of actinobacteria. *Microbiol Mol Biol Rev*. 2016;80(1):1–43.
35. Anandan R, Dharumadurai D, Manogaran GP. *An introduction to actinobacteria*. Actinobacteria - basics and biotechnological applications. Edited by Dharumadurai Dhanasekaran and Yi Jiang. Intech. 2016.
36. Jose PA, Jha B. Intertidal marine sediment harbours Actinobacteria with promising bioactive and biosynthetic potential. *Sci Rep*. 2017;7(1):10041.
37. Naligama KN, Weerasinghe KE, Halmillawewa AP. Characterization of bioactive actinomycetes isolated from kadolkele mangrove sediments, Sri Lanka. *Pol J Microbiol*. 2022;71(2):191–204.
38. Grover M, Bodhankar S, Maheswari M, et al. *Actinomycetes as mitigators of climate change and abiotic stress*. Singapore: Springer Singapore. *Plant growth promoting actinobacteria*. 2016:203–212.
39. Mukhtar S, Zaheer A, Aiysha D, et al. Actinomycetes: a source of industrially important enzymes. *J Proteomics Bioinform*. 2017;10(12):316–319.
40. Casas-Godoy L, Gasteazoro F, Duquesne S, et al. Lipases: An overview. *Methods Mol Biol*. 2018;1835:3–38.
41. Jaouani A, Neifar M, Hamza A, et al. Purification and characterization of a highly thermostable esterase from the actinobacterium *geodermatophilus obscurus* strain G20. *J Basic Microbiol*. 2012;52(6):653–660.
42. Nandimath AP, Karad DD, Gupta SG, et al. Consortium inoculum of five thermo-tolerant phosphate solubilizing actinomycetes for multipurpose biofertilizer preparation. *Iran J Microbiol*. 2017;9(5):295–304.
43. Wei R, Oeser T, Zimmermann W. Synthetic polyester-hydrolyzing enzymes from thermophilic actinomycetes. *Adv Appl Microbiol*. 2014;89:267–305.
44. Silveira MAV, Batista Dos Santos SM, Okamoto DN, et al. Atlantic Forest's and Caatinga's semiarid soils and their potential as a source for halothermotolerant actinomycetes and proteolytic enzymes. *Environ Technol*. 2023;44(11):1566–1578.
45. Jose PA, Jha B. Intertidal marine sediment harbours Actinobacteria with promising bioactive and biosynthetic potential. *Sci Rep*. 2017;7(1):10041.
46. Puttaswamygowda GH, Olakkaran S, Antony A, et al. *Chapter 22 - Present status and future perspectives of marine actinobacterial metabolites*. In: Buddolla, V. (Ed.), *Recent Developments in Applied Microbiology and Biochemistry*. Academic Press. 2019:307–319.
47. Jose PA, Maharshi A, Jha B. Actinobacteria in natural products research: Progress and prospects. *Microbiol Res*. 2021;246:126708.
48. Ouertani R, Ouertani A, Mahjoubi M, et al. New plant growth-promoting, chromium-detoxifying microbacterium species isolated from a tannery wastewater: Performance and genomic insights. *Front Bioeng Biotechnol*. 2020;8:521.
49. Mesbah NM. Industrial biotechnology based on enzymes from extreme environments. *Front Bioeng Biotechnol*. 2022;10:870083.
50. Han H, Ling Z, Khan A, et al. Improvements of thermophilic enzymes: from genetic modifications to applications. *Bioresour Technol*. 2019;279:350–361.
51. Liu T, Wang Y, Luo X, et al. Enhancing protein stability with extended disulfide bonds. *Proc Natl Acad Sci USA*. 2016;113(21):5910–5915.
52. Tang F, Chen D, Yu B, et al. Improving the thermostability of *Trichoderma reesei* xylanase 2 by introducing disulfide bonds. *Electron J Biotechnol*. 2017;26:52–59.
53. Landeta C, Boyd D, Beckwith J. Disulfide bond formation in prokaryotes. *Nat Microbiol*. 2018;3(3):270–280.

54. Karshikoff A, Nilsson L, Ladenstein R. Rigidity versus flexibility: the dilemma of understanding protein thermal stability. *FEBS J.* 2015;282(20):3899–3917.
55. Santiago M, Ramirez-Sarmiento CA, Zamora RA, et al. Discovery, molecular mechanisms, and industrial applications of cold-active enzymes. *Front Microbiol.* 2016;7:1408.
56. Sarmiento F, Peralta R, Blamey JM. Cold and hot extremozymes: industrial relevance and current trends. *Front Bioeng Biotechnol.* 2015;3:148.
57. Mangiagalli M, Brocca S, Orlando M, et al. The «cold revolution». Present and future applications of cold-active enzymes and ice-binding proteins. *N Biotechnol.* 2020;55:5–11.
58. Fujinami S, Fujisawa M. Industrial applications of alkaliphiles and their enzymes-past, present and future. *Environ Technol.* 2010;31(8–9):845–856.
59. Parashar D, Satyanarayana T. An insight into ameliorating production, catalytic efficiency, thermostability and starch saccharification of acid-stable α -amylases from acidophiles. *Front Bioeng Biotechnol.* 2018;6:125.
60. Oren A. Life at high salt concentrations, intracellular KCl concentrations, and acidic proteomes. *Front Microbiol.* 2013;4:315.
61. Brininger C, Spradlin S, Cobani L, et al. The more adaptive to change, the more likely you are to survive: protein adaptation in extremophiles. *Semin Cell Develop Biol.* 2018;84:158–169.
62. Akal AL, Karan R, Hohl A, et al. A polyextremophilic alcohol dehydrogenase from the Atlantis II deep red sea brine pool. *FEBS Open Bio.* 2018;9(2):194–205.
63. El-Ahmady El-Naggar N. Chapter 11 - *Streptomyces*-based cell factories for production of biomolecules and bioactive metabolites. *Microbial Cell Factories Engineering for Production of Biomolecules.* 2021;183–234.
64. Otani H, Uduary DW, Mouncey NJ. Comparative and pangenomic analysis of the genus *Streptomyces*. *Sci Rep.* 2022;12(1):18909.
65. Cuebas-Irizarry MF, Grunden AM. *Streptomyces* spp. as biocatalyst sources in pulp and paper and textile industries: Biodegradation, bioconversion and valorization of waste. *Microb Biotechnol.* 2024;17(1):e14258.
66. Vojnovic S, Aleksic I, Ilic-Tomic T, et al. *Bacillus* and *Streptomyces* spp. as hosts for production of industrially relevant enzymes. *Appl Microbiol Biotechnol.* 2024;108(1):185.
67. Besaury L, Fromentin J, Detain J, et al. Transcriptomic analysis of lignocellulose degradation by *Streptomyces coelicolor* A3(2) and elicitation of secondary metabolites production. *FEMS Microbiol Lett.* 2022;369(1):fnac101.
68. Challis GL. Exploitation of the *Streptomyces coelicolor* A3(2) genome sequence for discovery of new natural products and biosynthetic pathways. *J Ind Microbiol Biotechnol.* 2014;41(2):219–232.
69. Nithya K, Muthukumar C, Kadaikunnan S, et al. Purification, characterization, and statistical optimization of a thermostable α -amylase from desert actinobacterium *Streptomyces fragilis* DA7-7. *3 Biotech.* 2017;7(5):350.
70. Ruchika S. Isolation and characterization of thermophilic actinomycetes with extracellular enzyme and bio-surfactant production potential from thar desert, India. *Int J Microbiol Res.* 2016;8(4):743–746.
71. Primarini D, Ohta Y. Some enzyme properties of raw starch digesting amylases from *Streptomyces* sp. No. 4. *Starch.* 2000;52(1):28–32.
72. Chakraborty S, Raut G, Khopade A, et al. Study on calcium ion independent α -amylase from haloalkaliphilic marine *Streptomyces* strain A3. *Indian J Biotechnol.* 2012;11:427–437.
73. Lima A, Nascimento R, Bon E, et al. *Streptomyces drozdowiczii* cellulase production using agro-industrial by-products and its potential use in the detergent and textile industries. *Enz Microb Technol.* 2005;37(2):272–277.
74. Budihal SR, Agsar D, Patil SR. Enhanced production and application of acidothermophilic *Streptomyces* cellulase. *Bioresour Technol.* 2016;200:706–712.
75. Adams AS, Jordan MS, Adams SM, et al. Cellulose-degrading bacteria associated with the invasive woodwasp *Sirex noctilio*. *ISME J.* 2011;5:1323–1331.
76. Takasuka TE, Book AJ, Lewin GR, et al. Aerobic deconstruction of cellulosic biomass by an insect-associated *Streptomyces*. *Sci Rep.* 2013;3:1030.
77. Wu H, Cheng X, Zhu Y, et al. Purification and characterization of a cellulase-free, thermostable endo-xylanase from *Streptomyces griseorubens* LH-3 and its use in biobleaching on eucalyptus kraft pulp. *J Biosci Bioeng.* 2018;125(1):46–51.
78. Montiel MD, Hernández M, Rodríguez J, et al. Evaluation of an endo-beta-mannanase produced by *Streptomyces ipomoea* CECT 3341 for the biobleaching of pine kraft pulps. *Appl Microbiol Biotechnol.* 2002;58(1):67–72.
79. Kansoh AL, Nagieb ZA. Xylanase and mannanase enzymes from *Streptomyces galbus* NR and their use in biobleaching of softwood kraft pulp. *Antonie Van Leeuwenhoek.* 2004;85(2):103–114.
80. Sanjivkumar M, Silambarasan T, Palavesam A, et al. Biosynthesis, purification and characterization of β -1,4-xylanase from a novel mangrove associated actinobacterium *Streptomyces olivaceus* (MSU3) and its applications. *Protein Expr Purif.* 2017;130:1–12.
81. Rahman MA, Choi YH, Pradeep GC. A novel low molecular weight endo-xylanase from *Streptomyces* sp. CS628 cultivated in wheat bran. *Appl Biochem Biotechnol.* 2014;173(6):1469–1480.
82. Zhu Y, Li X, Sun B, et al. Properties of an alkaline-tolerant, thermostable xylanase from *Streptomyces chartreusis* L1105, suitable for xylooligosaccharide production. *J Food Sci.* 2012;77(5):C506–C511.
83. Shrestha S, Chio C, Khatiwada JR, et al. Optimization of cultural conditions for pectinase production by *Streptomyces* sp. and characterization of partially purified enzymes. *Microb Physiol.* 2023;33(1):12–26.
84. Hosseini Abari A, Amini Rourani H, Ghasemi SM, et al. Investigation of antioxidant and anticancer activities of unsaturated oligo-galacturonic acids produced by pectinase of *Streptomyces hydrogenans* YAM1. *Sci Rep.* 2021;11(1):8491.
85. Yuan P, Meng K, Shi P, et al. An alkaline-active and alkali-stable pectate lyase from *Streptomyces* sp. S27 with potential in textile industry. *J Ind Microbiol Biotechnol.* 2012;39(6):909–915.
86. Jacob N, Asha Poorna C, Prema P. Purification and partial characterization of polygalacturonase from *Streptomyces lydicus*. *Bioresour Technol.* 2008;99(14):6697–6701.
87. Ramirez Tapias YA, Rivero CW, Gallego FL, et al. Stabilization by multipoint covalent attachment of a biocatalyst with polygalacturonase activity used for juice clarification. *Food Chem.* 2016;208:252–257.
88. Tanaya Behera H, Mojumdar A, Kumari K, et al. Exploration of genomic and functional features of chitinolytic bacterium *Streptomyces chilikensis* RC1830, isolated from Chilika Lake, India. *3 Biotech.* 2022;12(5):120.
89. Mukherjee G, Sen SK. Purification, characterization, and antifungal activity of chitinase from *Streptomyces venezuelae* P10. *Curr Microbiol.* 2006;53(4):265–269.
90. Rajendran K, Krishnamoorthy M, Karuppiiah K, et al. Chitinase from *Streptomyces mutabilis* as an effective eco-friendly biocontrol agent. *Appl Biochem Biotechnol.* 2024;196(1):18–31.

91. Gangwar M, Singh V, Pandey AK, et al. Purification and characterization of chitinase from *Streptomyces violascens* NRRL B2700. *Indian J Exp Biol*. 2016;54(1):64–71.
92. Nagpure A, Gupta RK. Purification and characterization of an extracellular chitinase from antagonistic *Streptomyces violaceusniger*. *J Basic Microbiol*. 2013;53(5):429–439.
93. Ramesh S, Rajesh M, Mathivanan N. Characterization of a thermostable alkaline protease produced by marine *Streptomyces fungicidicus* MML1614. *Bioprocess Biosyst Eng*. 2009;32(6):791–800.
94. Ben Elhoul MB, Jaouadi NZ, Rekika H, et al. A novel detergent-stable solvent-tolerant serine thiol alkaline protease from *Streptomyces koyangensis* TN650. *Int J Biol Macromol*. 2015;79:871–882.
95. Manivasagan P, Venkatesan J, Sivakumar K, et al. Production, characterization and antioxidant potential of protease from *Streptomyces* sp. MAB18 using poultry wastes. *Biomed Res Int*. 2013;2013:496586.
96. Jaouadi B, Abdelmalek B, Fodil D, et al. Purification and characterization of a thermostable keratinolytic serine alkaline proteinase from *Streptomyces* sp. strain AB1 with high stability in organic solvents. *Bioresour Technol*. 2010;101(21):8361–8369.
97. Tatineni R, Doddapaneni KK, Potumarthi RC, et al. Purification and characterization of an alkaline keratinase from *Streptomyces* sp. *Bioresour Technol*. 2008;99(6):1596–1602.
98. Syed DG, Lee JC, Li WJ, et al. Production, characterization and application of keratinase from *Streptomyces gulbargensis*. *Bioresour Technol*. 2009;100(5):1868–1871.
99. Chao YP, Xie FH, Yang J, et al. Screening for a new *Streptomyces* strain capable of efficient keratin degradation. *J Environ Sci (China)*. 2007;19(9):1125–1128.
100. Thumar JT, Singh SP. Secretion of an alkaline protease from a salt-tolerant and alkaliphilic, *Streptomyces clavuligerus* strain MIT-1. *Braz J Microbiol*. 2007;38(4):766–772.
101. Bajaj BK, Sharma P. An alkali-thermotolerant extracellular protease from a newly isolated *Streptomyces* sp. DP2. *N Biotechnol*. 2011;28(6):725–732.
102. Ayaz B, Ugur A, Boran R. Purification and characterization of organic solvent-tolerant lipase from *Streptomyces* sp. OC119-7 for biodiesel production. *Biocatal Agric Biotechnol*. 2015;4(1):103–108.
103. Mander P, Yoo H-Y, Kim SW, et al. Transesterification of waste cooking oil by an organic solvent-tolerant alkaline lipase from *Streptomyces* sp. CS273. *Appl Biochem Biotechnol*. 2014;172(3):1377–1389.
104. Boran R, Ugur A, Sarac N, et al. Characterisation of *Streptomyces violascens* OC125-8 lipase for oily wastewater treatment. *3 Biotech*. 2019;9(1):5.
105. Vahidi M, Imanparast S, Jahandar H, et al. An organic solvent-tolerant lipase of *Streptomyces pratensis* MV1 with the potential application for enzymatic improvement of n6/n3 ratio in polyunsaturated fatty acids from fenugreek seed oil. *J Food Sci Technol*. 2021;58(7):2761–2772.
106. Yuan D, Lan D, Xin R, et al. Screening and characterization of a thermostable lipase from marine *Streptomyces* sp. strain W007. *Biotechnol Appl Biochem*. 2016;63(1):41–50.
107. H-Kittikun A, Prasertsan P, Zimmermann W, et al. Sugar ester synthesis by thermostable lipase from *Streptomyces thermocarboxydus* ME168. *Appl Biochem Biotechnol*. 2012;166(8):1969–1982.
108. Mander P, Yoo HY, Kim SW, et al. Transesterification of waste cooking oil by an organic solvent-tolerant alkaline lipase from *Streptomyces* sp. CS273. *Appl Biochem Biotechnol*. 2014;172(3):1377–1389.
109. Côté A, Shareck F. Expression and characterization of a novel heterologous moderately thermostable lipase derived from metagenomics in *Streptomyces lividans*. *J Ind Microbiol Biotechnol*. 2010;37(9):883–891.
110. Wang B, Wang A, Cao Z, et al. Characterization of a novel highly thermostable esterase from the Gram-positive soil bacterium *Streptomyces lividans* TK64. *Biotechnol Appl Biochem*. 2016;63(3):334–343.
111. Zhou X, Huang J, Ou Z, et al. Conditions of enzyme production and properties of alkaline lipase by *Streptomyces* Z94-2. *Wei Sheng Wu Xue Bao*. 2000;40(1):75–79.
112. Brault G, Shareck F, Hurtubise Y, et al. Isolation and characterization of EstC, a new cold-active esterase from *Streptomyces coelicolor* A3(2). *PLoS One*. 2012;7(3):e32041.
113. Rodríguez-Alonso G, Toledo-Marcos J, Serrano-Aguirre L, et al. A novel lipase from *Streptomyces exfoliatus* DSMZ 41693 for biotechnological applications. *Int J Mol Sci*. 2023;24(23):17071.
114. Li M, Zhou Y, Duan X, et al. Characterization of a phospholipase D from *Streptomyces cinnamoneum* SK43.003 suitable for phosphatidylserine synthesis. *Biotechnol Appl Biochem*. 2022;69(5):1917–1928.
115. Simkhada JR, Lee HJ, Jang SY, et al. A novel alkalo- and thermostable phospholipase D from *Streptomyces olivochromogenes*. *Biotechnol Lett*. 2009;31(3):429–435.
116. EFSA Panel on Food Contact Materials, Enzymes and Processing Aids (CEP); Lambré C, Barat Baviera JM, et al. Safety evaluation of the food enzyme phospholipase A2 from the genetically modified *Streptomyces violaceoruber* strain AS-10. *EFSA J*. 2023;21(2):e07458.
117. Nivedita S, Behera SS, Behera HT, et al. Comparative genome-wide analysis of novel *Streptomyces* isolates RC1831 and RC1832: deciphering the role of functional carbohydrate (CAZy) active genes including chitinase for production of chitosan. *3 Biotech*. 2024;14(4):114.
118. Escudero-Agudelo J, Martínez-Villalobos J, Arocha-Garza H, et al. Systematic bioprospection for cellulolytic actinomycetes in the Chihuahuan Desert: isolation and enzymatic profiling. *Peer J*. 2023;11:e16119.