REVIEW ARTICLE



Biotechnological potential of psychrophilic microorganisms as the source of cold-active enzymes in food processing applications

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Received: 1 July 2021 / Accepted: 29 September 2021 / Published online: 30 October 2021 © King Abdulaziz City for Science and Technology 2021

Abstract

Microorganisms striving in extreme environments and exhibiting optimal growth and reproduction at low temperatures, otherwise known as psychrophilic microorganisms, are potential sources of cold-active enzymes. Owing to higher stability and cold activity, these enzymes are gaining enormous attention in numerous industrial bioprocesses. Applications of several cold-active enzymes have been established in the food industry, e.g., β -galactosidase, pectinase, proteases, amylases, xylanases, pullulanases, lipases, and β -mannanases. The enzyme engineering approaches and the accumulating knowledge of protein structure and function have made it possible to improve the catalytic properties of interest and express the candidate enzyme in a heterologous host for a higher level of enzyme production. This review compiles the relevant and recent information on the potential uses of different cold-active enzymes in the food industry.

Keywords Psychrophiles · Cold-active enzymes · Food industry · Enzyme engineering

Introduction

The term "psychrophiles" is a Greek word coming from *psukhros* meaning 'cold' and *philein*, 'loving'. Psychrophiles are defined as microorganisms that can grow and sustain in cold environments, such as deep sea, high elevations and Polar Regions of the planet Earth, and can tolerate the temperature range 0–20 °C; however, the optimal temperature for their growth is 5 °C (Cavicchioli 2016; Salwan and Sharma 2020). The first known and taxonomically described species of psychrophiles are *Vibrio marinus* and *V. psychroerythrus* (Morita and Moyer 2001). Nonetheless, several species of bacteria (*Bacillus, Bacteroides, Arthrobacter, Clostridium, Pseudomonas*, and *Methanogenium*) and fungi (*Pseudogymnoascus*, and *Geomyces*) have also

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² Centre of Innovative and Applied Bioprocessing, Mohali, India been described to be psychrophilic (Morita and Moyer 2001; Meteyer and Verant 2019). Psychrophiles are well adapted to extreme environments and possess complex metabolic adaptations, such as altered nutrient transport mechanisms, intracellular ice formation and cold denaturation of proteins (Feller and Gerday 2003). Besides, these microorganisms are potentially rich enzymes that can maintain high activity even at low temperatures by reducing the temperature dependence of the reaction (Yayanos 2009; Cavicchioli et al. 2011; Irwin 2020; Rai and Rakshak 2015).

These cold-active or psychrophilic enzymes are advantageous for many industrial applications as they are (i) costeffective, (ii) able to catalyze reaction without any additional heat aid and (iii) can be inactivated selectively by gentle heat input. Cold-active enzymes are also used in several biotechnology applications to prevent many undesirable reactions and restrict the loss of volatile components (Gerday 2013; Kuddus 2018). Due to these inevitable applications, the role of cold-active enzymes in the coming years is expected to witness generous market growth (Santiago et al. 2016; Al-Ghanayem and Joseph 2020). Numerous possible prospects of these cold-active enzymes have also become evident in food processing applications as the low temperature reduces many adverse reactions and microbial contamination (Kuddus and Ramteke 2012; Javed and Qazi 2016). Production of fruit juices, alcoholic beverages, chocolates, sweeteners,



cheese, bakery products and milk products are typical applications of cold-active enzymes.

Despite the tremendous importance of cold-active enzymes, finding novel enzymes for their commercial applications in the food industry is a growing challenge (Kuddus 2018). On the other hand, less availability and low stability of these enzymes are significant signs. Furthermore, biotechnologists are more concerned about the specific catalytic activity of these cold-active enzymes, which need to be explored at the industrial level (Kuddus 2018; Hamid and Mohiddin 2018). Therefore, the development and invention of new technologies are needed to boost the quality and production of cold-active enzymes. Moreover, with the advent of approaches, such as protein engineering, rDNA technology and metagenomics, these enzymes can be devised to be used competently for diverse food applications (Thakur et al. 2021a). This review provides insights into applying cold-active enzymes in food processing, emphasizing the recombinant clones and their industrial significance (Fig. 1).

Psychrophilic microbes as producers of cold-active enzymes

Psychrophiles represent an extensive range of microbial taxa such as bacteria, archaea, algae and yeasts that can thrive in permanently cold habitats, including glaciers, high mountains, ocean depths, shallow subterranean and polar regions. They experience severe physicochemical constraints, including decreased biochemical reaction rates, reduced membrane fluidity, altered transport systems and cold denaturation of proteins (D'Amico et al. 2006; Siddiqui and Cavicchioli 2006). Adaptive features to overcome these constraints have frequently been detected in the genomes of several psychrophiles sequenced so far (Saunders et al. 2003; Riley et al. 2008; Ayala-Del-Rio et al. 2010). Moreover, the production of cold-active enzymes that drive metabolism and the cell cycle in these microorganisms has been considered a characteristic adaptation. The enzymes offer high specific activity even at low temperatures, compensating for the

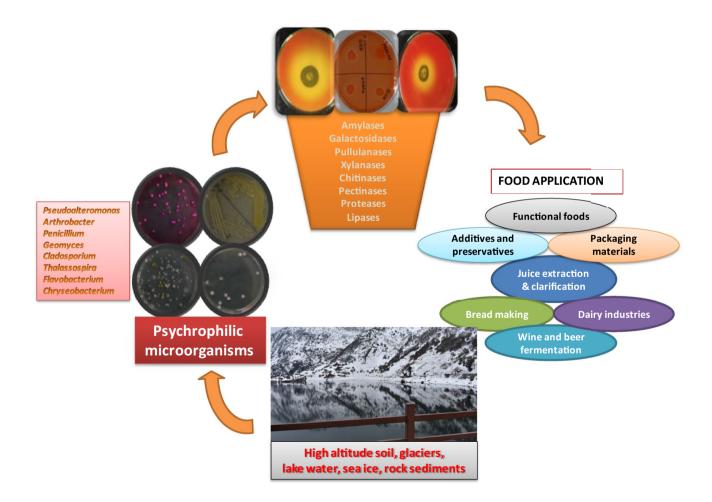


Fig.1 Schematic representation showing application of cold active enzymes produced by psychrophilic microorganisms in food processing industry



rapid decrease in chemical reaction rates. This adaptive feature is thought to be genetically encoded in the amino acid sequence and reportedly resulted from a long term selection process (Feller and Gerday 2003; Feller 2013).

Psychrophilic microorganisms contain many critical proteins and other metabolites as determinative factors in their adaptation to cold environments. They evolve mechanisms to maintain maximal translation and protein folding under cold conditions (Siddiqui and Cavicchioli 2006; Feller 2013). The protein components liable for the maintenance of RNA for carrying out the translation process, specifically the ribosomal proteins and RNA helicases, appeared to be overexpressed in many cold-adapted microorganisms (Lim et al. 2000; Jung et al. 2010). Furthermore, diminution in the number and strength of non-covalent interactions outside the catalytic cavity offers motional and dynamic strength to the active site amino acids upholding catalytic efficiency of the enzymes in cold temperatures (Feller 2013). More importantly, the chaperones including DnaK, GroEL, ribosomebound trigger factor (TF) and RNA chaperones, such as cold-shock protein A, which have reportedly played a crucial role in the folding of newly synthesized polypeptides, contribute significantly to the adaptive feature of psychrophilic microorganisms and conformational flexibility of the coldactive enzymes (Strocchi et al. 2006; Sung et al. 2011; Piette et al. 2011). Numerous microorganisms, including bacteria (species of Arthrobacter, Pseudolteromonas, Paracoccus, Pseudomonas, etc.) and fungi (species of Geomyces, Candida, Penicillium, Cladosporium, etc.) residing in the Antarctic and Arctic habitats, and high elevation regions have been investigated for the production of cold-active enzymes, which demonstrate the remarkable potential to be used in various industries (Santiago et al. 2016; Hamid and Mohiddin 2018; Bruno et al. 2019).

Cold-active enzymes in the food industry

Cold-active enzymes inherit flexible structures that presumably recompense for the low kinetic energy present in the cold habitats. Because of this reason, these enzymes often demonstrate a reduction in the activation enthalpy (Δ H) while showing more negative activation entropy (Δ S) as compared to their thermophilic and mesophilic counterparts. That is why the reaction rate of cold-active enzymes tends to decrease very slowly concerning a decrease in temperature around the surrounding. This balancing act of activation parameters is transformed to the high catalytic activity of these enzymes at low temperatures (Siddiqui and Cavicchioli 2006; Cavicchioli et al. 2011). Therefore, the biotechnological potential of cold-active enzymes reflects several factors, including their high activity in low to moderate temperatures, increasing thermolability at high temperatures, and the ability of these enzymes to carry out the reaction in organic solvents (Marx et al. 2007; Margesin and Feller 2010). Cold-active enzymes could also be able to catalyze reactions at temperatures that play down many undesirable and competitive chemical reactions. These low-temperature active enzymes are of particular interest for transforming heat-sensitive substrates (Jeon et al. 2009). Because of these unique features, cold-active enzymes are considered as most germane to the food and feed industry. Importance is being given to evade spoilage, change in flavour and nutritional parameters of the native thermolabile substrates and products. Cold-active enzymes are widely used in food applications, such as meat tenderization, baking, brewing, flavouring, cheese production, and animal feed processing (Tables 1 and 2).

According to the International Union of Biochemistry and Molecular Biology (IUBMB), enzymes are classified into six major categories based on the type of reactions they catalyze: oxidoreductases, transferases, hydrolases, lyases, isomerases, and ligases. Though these enzymatic reactions are vital in food production, food biotechnologists are mainly concerned with oxidoreductases and hydrolases (Chourasia et al. 2020; Raveendran et al. 2018; Bruno et al. 2019). Hydrolases are given preference, considering their significant roles in cheese production, malting and brewing, conversion of starch to glucose and fructose, and human metabolism (Fernandes 2010; Adrio and Demain 2014; Sarmiento et al. 2015). The anticipated food applications of cold-active hydrolytic enzymes are discussed in the subsequence sections.

Hydrolases

Hydrolases constitute a very composite collection of enzymes that catalyze bond cleavages upon reaction with water. The innate role of most hydrolases is digestion that is breaking down nutrients into smaller units suitable for digestion, and because of this, they usually have a broad range of substrate specificity. Examples include proteases (hydrolyze proteins to oligopeptides and amino acids), lipases (hydrolyze triglycerides to glycerol and fatty acids), phosphatases, pectinases, etc.

β-Galactosidases

 β -Galactosidases offer an obvious example of glycosidases, with the advantage of being active at low temperatures and cost-effective (Bruno et al. 2019). β -Galactosidases are used to remove lactose from milk and dairy products, where the enzyme hydrolyzes lactose into glucose and galactose. With more than half of the world's population suffering from lactose intolerance and also because of the difficulties in crystallizing lactose, β -galactosidases



Cintinase	and marine sediment (Antarctica)	r seudomonus sp.	spoilage of cold-stored foods and against vegetable spoilage pathogenic fungi	Liue
Protease	Deep sea sediment (China)	Pseudoaltermonas sp. SM9913	Taste enhancement of cold- stored meat	He e
	Sausage	Penicillium nalgiovense PNA9	Meat ripening	Papa (20
	NS	Arsukibacteriumikkense	Production of bioactive dairy products and other functional foods	De C
	East Rathong Glacier, Sik- kim	Chryseobacterium polytri- chastri	Production of antioxidant peptides	Muk
	Sea ice (Antarctica)	<i>Pseudoalteromonas</i> sp. NJ276	As additives in baking indus- try, food processing and preservation industries	Wan
	Refrigerator of a meat fac- tory (China)	Serratia sp. WJ39	Food processing industry	Ji et :
Phytase	Deep sea sediment (Antarc- tica)	Rhodotorula mucilaginosa JMUY14	Feed industry especially aquaculture feed process- ing	Yu e
	Antarctic sample	Cryptococcus laurentii AL27	Feed industry	Pavlo
Pectinase	Marine sponge (Antarctica)	Geomyces sp. F09-T3-2	Production of white wine	Pove
	Fruit orchard soil and spoiled refrigerated fruits and vegetables (Himalaya)	Saccharomyces sp.	Fruit juice clarification	Naga
	Grapes (Argentina)	Bacillus sp. CH15	Red wine making	Mart An
	Grape wine and wineries (Argentina)	Aureobasidium pullulans strains	Cold-wine making	Merí (<mark>20</mark>
NS not specified				
(Dalmaso et al.	prowing attention of food bi 2015). Cold-active β-galac) can improve the technica	tosidases (act- and ma	l-active β-galactosidases from the Antarctic any are from the Antarctic (2013) and Laye et al. (2017)	and A

Table 1 Cold-active enzymes produced by psychrophilic microorganisms and their potential applications in food industry

Microorganisms

Pseudoalteromonas halo-

Potential/recommended

Xylo-oligosaccharides

application

large intestine. Several studies have reported the isolation

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sychrophilic bacteria, Arctic regions. Karan scribed the cold-active 1 /) de β-galactosidases from Halorubrum lacusprofundi, a bacterial psychrophile isolated from the hypersaline Deep Lake of Antarctica. Several β-galactosidases, isolated from species of Paracoccus, Arthrobacter and Pseudoalteromonas (P. haloplanktis TAE 79 and LMGP-19143) of the Antarctic region, are functional at low temperatures and have potential application in the milk and dairy industry (Trimbur et al. 1994; Hoyoux et al. 2001; Turkiewicz et al. 2003; Cieslinski et al. 2005; Makowski et al. 2007; Hildebrandt et al. 2009). Cold-active β -galactosidases showing considerable lactose

Xylanase

Cold-active enzymes Sample source

Soil sample (Antarctica)

References

Collins et al. (2002)

planktis production β-Galactosidase Sea sediment (Antarctica) Guehomyces pullulans 17-1 Dairy industry for pro-Song et al. (2010) duction of lactose free products Soil (Antarctica) Rahnellainusitata -Do-Núñez-Montero et al. (2021) Chitinase Soil, seal and penguin feces, Pseudomonas sp. Biocontrol of microbial Liu et al. (2019) et al. (2004) agianni and Sergelidis 2014) Gobba et al. (2014) khia et al. (2021) ng et al. (2008) t al. (2014) et al. (2015) lova et al. (2008) veda et al. (2018) ga Padma et al. (2011) rtín and Morata de mbrosini (2013) erín and de Ambrosini 2015)

of whey by producing syrups rich in glucose and galactose

that can be utilized in a variety of food products (Gerday

et al. 2000). These enzymes also possess transglycosyla-

tion activities to catalyze the hydrolysis of lactose with the

instantaneous transfer of the monosaccharides to higher

oligosaccharides forming tri and tetrasaccharides (Kar-

asova-Lipovova et al. 2003; Benesova et al. 2005). Such

galacto-oligosaccharides have potential uses as prebiotic

additives that enhance the growth of Bifidobacteria in the

Cold-active enzymes	Sample source	Source microorganisms	Cloning/expression host	Expression vector	Potential/recommended applications	References
Xylanase	Sea water (China)	Zunongwangia profunda	Escherichia coli BL21, E. coli DH5a	pGEX-6p-1	Food industry for process- ing of sea and saline foods	Liu et al. (2014)
	NS	Saccharophagus- degradans 2–40	<i>E. col</i> i DH5α, <i>E. coli</i> BL21	pET21a	Baking industry	Ko et al. (2016)
	SN	Pseudoalteromons halo- planktis TAH3A, Flavo- bacterium sp. MSY-2	E. coli BL21	pET22b-XFH-His6, pET22b-Xyn8-His6	Baking industry	Dornez et al. (2011)
	NS	Glaciecola mesophile	E. coli BL21	pET22b	Baking industry	Zheng et al. (2011)
	NS	Flavobacterium johnso- niae	<i>E. coli</i> JM109, <i>E. coli</i> DH5α, <i>E. coli</i> BL21	pSCH602, pSCH643	Food and feed industry	Chen et al. (2013)
β-Galactosidase	Calcium carbonate (Ikka columns, South-West Greenland)	Alkalilactibacillusikkense	E. coli TOP10	pUC18dLacZ	-D0-	Schmidt and Stougaard (2010)
	Soil (Antarctica)	Arthrobacter sp. 20B	E. coli TOP10F'	NS	-Do-	Białkowska et al. (2009)
	Soil (Antarctica)	Paracoccus sp. 32d	E. coli LMG 194	pBAD/Myc-His A	-Do-	Wierzbicka-Woś et al. (2011)
	Deep sea water (Mariana Trench)	Alteromonas sp. ML52	E. coli BL21	pET-gal	-Do-	Sun et al. (2018)
	Alimentary tract (krill Thyssanoessamacrura) (Southern Shetlands)	Pseudoalteromonas sp. 22b	E. coli ER2566	pETbeta22b	-D0-	Cieśliński et al. (2005)
	Frozen soil (China)	Rahnella sp. R3	E. coli BL21	pCold I	-Do-	Fan et al. (2015)
a-amylase	Surface sea water (China)	Zunongwangia profunda (MCCC 1A01486)	E. coli DH5a, E. coli BL21	pGEX-6P-1	Baking industry	Qin et al. (2014)
	Deep sea sediment (China)	Bacillus sp. dsh19-1	E. coli JM109, E. coli BL21	pColdI	Starch hydrolysis	Dou et al. (2018)
	Sea ice (Antarctica)	Pseudoalteromonas sp. M175	E. coli BL21	pET-amy175	Food industry	Wang et al. (2018a)
	Soil (Tehran)	Exiguobacterium sp. SH3	E. coli DH5a, E. coli BL21	pEamy	Food industry	Mojallali et al. (2014), Emampour et al. (2015)
	Soil (King George Island, Antarctica)	Geomycespannorum	E. coli DH5a, Aspergillus pBC-hygro oryzae	pBC-hygro	Food industry	Krishnan et al. (2011), Mao et al. (2015)
	NS	Bifidobacterium longum	E. coli MC1061	NS	In food processing indus- try for production of slowly digestible starch	Lee et al. (2016)
β-Glucosidase	Konjac field (South Korea)	Paenibacillusxylanilyticus KJ-03	E. coli JM109, E. coli BL21	pColdI	Food processing industry	Park et al. (2013)

Cold-active enzymes	Sample source	Source microorganisms	Cloning/expression host	Expression vector	Potential/recommended applications	References
Pullulanase	Soil (Tehran)	Exiguobacterium sp. SH3	E. coli BL21	pET-26b (+)	Cold hydrolysis of starch and in sugar syrup production	Mojallali et al. (2014), Rajaei et al. (2015)
	Soil (China)	Paenibacilluspolymyx- aNws-pp2	E. coli DH5α, E. coli BL21	pET-28a, pET-32a and pET-42a	Food processing industry	Wei et al. (2015)
	Hot-spring (Sikkim, India) Hot-spring metagenome	Hot-spring metagenome	E. coli BL21	pET28a(+)	Production of resistant starch	Thakur et al. (2021b)
	Waste water (Europe)	Bacillus methanolicus PB1	<i>E. col</i> i DH5α, <i>E. coli</i> BL21	pET-28a	Cold hydrolysis of starch	Heggeset et al. (2012), Zhang et al. (2020)
Chitinase	Sea sediment (China)	Pseudoalteromonas sp. DL-6	E. coli BL21	pET28a	Biocontrol of microbial spoilage of cold-stored foods	Wang et al. (2014a; b)
	Sea ice (Antarctica)	Glaciozyma antarctica PI12	E. coli JM109, Pichia pastoris	pPICZ α A	Biocontrol of microbial spoilage of cold-stored foods	Ramli et al. (2013)
Lipase/esterase	Deep sea water (Indian ocean)	Aeromicrobium sp. SCSIO 25071	<i>E. coli</i> DH5α, <i>E. coli</i> Rosetta (DE3)	pET28a	As food additives	Su et al. (2016)
	Marine sediment (Arctic)	Halocynthiibacter arcti- cus	E. coli BL21	pET-21a	Production of flavored compounds	Le et al. (2020)
	NS	Streptomyces coelicolor A3(2)	<i>E. coli</i> BL21 (DE3)	pET16b	Production of flavored compounds	Brault et al. (2012)
	NS	Malassezia globose	E. coli Top10, P. pastoris X-33	pGAPZaA	Oil modification and food processing industry	Xu et al. (2015)
	Car service contaminated soil (Malaysia)	Staphylococcus epider- midis AT2	E. coli Tuner (DE3)pLacI	pTrcHis2-TOPO	Food processing industry	Kamarudin et al. (2014)
	Cold seafloor dwelling Atlantic hagfish stomach (North Norway)	Rhodococcus sp. AW25M09	E. coli DH5-α, E. coli BL21 (DE3)	pET-22b	Food industry for preserv- ing food flavor and integrity	De Santi et al. (2014)
	NS	Neisseria meningitides	E. coli	pQE-30	Production of organic esters for use in food industry	Yoo et al. (2019)
	NS	Lactobacillus acidophilus NCFM	<i>E. coli</i> DH5-α, <i>E. coli</i> BL21 (DE3)	pET-21a	Dairy and food processing industry	Wang et al. (2018b)
	NS	Bacillus licheniformis ATCC 14580	E. coli BL21	pGEMT-Easy Vector	Food processing industry	Borgi et al. (2014)
Pectinase	Soil (Sub-Antarctic	Tetracladium sp.	E. coli, P. pastoris	pPinka-HC	Production of fruit juice	Carrasco et al. (2019)

NS not specified

References	Huang et al. (2012)
Potential/recommended applications	Food and feed industry
Expression vector	pET-32a
Cloning/expression host	E. coli JM109, E. coli Rosetta-gami (DE3)
Source microorganisms	Bacillus subtilis Bs5
Cold-active enzymes Sample source	Soil (China)
Cold-active enzyme	Mannanase

hydrolyzing efficiency were isolated from *Enterobacter ludwigii* and *Alkalilactibacillus ikkense* inhabiting the Arctic region (Schmidt and Stougaard 2010; Alikkunju et al. 2016). Besides, filamentous psychrophilic fungi colonizing the Antarctic environments hold the similar potential of producing β -galactosidases as those of bacterial psychrophiles. For instance, β -galactosidases isolated from *Tausonia pullulans* and *Cryptococcus albidus* colonizing Antarctic sea sediments reportedly hydrolyze lactose (Koleva et al. 2006; Song et al. 2010; Zhang et al. 2012).

Xylanases

Xylanases that catalyze the endohydrolysis of $1,4-\beta$ -Dxylosidic linkages in xylan grasp the enormous potential for extensive use in biomass processing and the food industry (Joshi et al. 2020; Phukon et al. 2020a). They have an essential role in bread-making, since they transform insoluble hemicellulose in the dough to soluble sugars, giving in elastic and soft bread. Few cold-active enzymes, purified from psychrophilic bacteria inhabiting Antarctic habitats, such as Pseudoalteromonas haloplanktis and Flavobacterium sp., have been observed to improve the dough properties along with the bread volume (Collins et al. 2002; Dornez et al. 2011). Few shreds of evidence of filamentous fungi (e.g., Geomyces pannorum and Cladosporium sp.), colonizing extreme habitats such as Polar Regions, producing xylanases have also been described (Del-Cid et al. 2014). The xylanolytic activity of the fungi was reportedly intact at low temperatures with very low stability; however, detailed characterization of xylanases and their physicochemical properties from fungi are truly limited. Therefore, further studies on the psychrophilic xylanases from microorganisms of diverse cold habitats should be driven forward, and efforts must be assumed to facilitate them as model candidates for different food processing applications.

Amylases

Amylases are another kind of glycoside hydrolases that catalyze starch hydrolysis by acting upon α -1,4-glycosidic bonds, a covalent bond joining two α -D-glucose together. They hydrolyze starch to form malotriose, maltose, glucose monomers, and limit dextrins. Amylases are further classified based on the specificity of the reactions catalyzed by them, and are exoamylases (e.g., β -amylases, glucoamylases and α -glucosidases), endoamylases (e.g., α -amylases), debranching amylases (e.g., pullulanases, isoamylases, and dextrinases), and transferases (e.g., 4- α -glucanotransferases, and cyclodextrin glycosyltransferases) (van der Maarel et al. 2002). Amylases have tremendous potential for use in food applications, including wine and beer fermentation and bread and fruit juices.



Several cold-active amylases have been purified from psychrophilic microorganisms (Srimathi et al. 2007; Roohi et al. 2013; Ramli et al. 2013; Qin et al. 2014; Rajaei et al. 2015). The most studied cold-active amylases are a-amylases, but only a few have been proposed for possible application in the food industry, including a novel a-amylase isolated from a psychrophilic fungus, *Geomyces panorum*, that is reported to have latent application for the baking industry (He et al. 2017).

Chitinases

Chitinases can break down chitin by hydrolyzing N-acetyl- β -D-glucosaminide (1 \rightarrow 4)- β -linkages randomly into oligo- and monomeric components. Chitinases have been influential in various biotechnological sectors because of their role in the bioconversion of chitinous biomass into several value-added products. Although poorly studied, cold-active chitinases and their high activity at fairly low temperatures can be used in the production of chitooligosaccharides (COS), the oligomers of N-acetyl-D-glucosamine (GlcNAc) and D-glucosamine (GlcN) linked by β-1,4-O-glycoside bond released as a result of chitin degradation (Le and Yang 2019; Rajabi 2019). COS are low weight water-soluble substances having interesting bioactivities and can easily be absorbed by the small intestine. COS can find application in the food industry as a food additive and packaging materials as its antimicrobial and antioxidant properties enable protection from deterioration (Liu et al. 2019; Rajabi 2019). Cold-active chitinases can be used to produce single-cell proteins for use as cheaper and alternative protein dietary supplements to soymeal and fish meal (Patil and Jadhav 2014; Wadhwa and Bakshi 2016). In addition, cold-active chitinases offer antifungal activity, which could be helpful in bio-control of postharvest fungal pathogens and, therefore, demonstrating the considerable potential for the food industry in safe storage of vegetables and fruits (Castillo et al. 2016; Le and Yang 2019). Reports on cold-active chitinase from psychrophilic microorganisms are not many except a cellbound cold-active chitobiase, exochitinase isolated from psychrophilic Arthrobacter sp. TAD20 (Lohienne et al. 2001). Despite the tremendous significance, the use of cold-active chitinases has not witnessed any considerable progress in food industries, which could be because of their less availability and specificity. We must prioritize their isolation and exploration from varied sources, including the psychrophilic microorganisms. Furthermore, they can be tailored and expressed in selected hosts to improve their stability and yield towards enhanced applicability in food biotechnology.



Other hydrolases

Pullulanases are critical glycosidase enzymes in starch processing and are primarily used in making maltose and glucose syrups (Thakur et al. 2021a). Pullulanases catalyze the hydrolysis of α -1,6-and α -1,4-linkages in starch, pullulan, amylopectin and other related oligosaccharides (Hii et al. 2012). Many cold-active pullulanases have been identified and characterized and shown their efficiency for other biotechnology applications, such as starch degradation for bioethanol production and transformation of biomass-derived starch into resistant starch (Elleuche et al. 2014; Thakur et al. 2021a, b).

Other glycosidases that represent a prospective candidate for the food industry are cellulases. They catalyze total hydrolysis of cellulose into sugars and are used in coffee processing and making wine (Kumar et al. 2018; Jayasekara and Ratnayake 2019). Cellulases, pectinases, and hemicellulase are also used to extract fruit juices and minimize food spoilage (Kuhad et al. 2011). Three cellulolytic activities have been cited from cellulases of natural origin: endoglucanase, β -glucosidase, and β -1,4-cellobiohydrolase. Over the decades, several cold-active cellulases sourced from psychrophilic microorganisms have been purified and characterized (Benesova et al. 2005; Shipkowski and Brenchley 2005; Zeng et al. 2006; Fu et al. 2010); however, their potential applicability in the food industry is scarcely known. Conversely, bioprospecting novel enzymes of these kinds from diverse cold-adapted microbial resources could address the present limitations of cost and energy associated with the use of thermophilic and mesophilic enzymes in various industrial processes.

Proteases

Proteases or peptidases are an interesting class of hydrolyzing enzymes that catalyze the hydrolysis of large proteins into minor peptides. Basing upon their ability to hydrolyze N- or C-terminal peptide bonds and internal peptide bonds, proteases are classified, respectively, into exopeptidases and endopeptidases (Chourasia et al. 2021). Exopeptidases cleaving N-terminal peptide bonds are called aminopeptidases, whereas those hydrolyze C terminal peptide linkage are called carboxypeptidases. Cold-active proteases hold enormous potential due to their several unique features, including optimal activity at low temperature, enabling an easy transformation of thermolabile products (Kuddus and Ramteke 2012; Kuddus 2018). In addition, they have specific applications in processes, such as the fermentation of fish and soya sauce, contributing nil change in nutritional value and flavour. They can also be considered alternatives to rennet, speeding up the ripening process of slow ripening cheeses when used with lipase. Cold-active proteases can also be profitable in taste development and softening of frozen meat products (Joshi and Satyanarayana 2013). Several proteases have been recovered from psychrophilic microorganisms, including bacteria, fungi and algae (Peterson et al. 2013; Vaca et al. 2013; Joshi and Satyanarayana 2013). A cold-active protease isolated from psychrophilic *Pseudoalteromonas* sp. has been reported to secrete essential amino acids that selectively enhance the taste of frozen meat (He et al. 2004). The psychrophilic serine protease belonging to *Chryseobacterium* sp. has been proposed for its applicability in improving meat quality because of its tolerance to salt and optimal activity at low temperature (Mageswari et al. 2017).

Cold-active protease obtained from Pseudoalteromonas sp. produced free amino acids from milk protein at 4 °C, implicating its significance in low-temperature food processing (Wang et al. 2008). A metalloprotease isolated from psychrophilic Enterococcus faecalis displayed no side effects while administered orally and can be used in functional foods, increasing their stability and solubility (Yuan et al. 2009). Similarly, cold-adapted proteases from Arsukibacterium ikkense, releasing bioactive peptides from milk caseins, have been accounted for indicating the practical suitability of these enzymes in dairy industry (De Gobba et al. 2014). Besides psychrophilic bacteria, fungi colonizing the extreme habitats have been known to produce proteases active at low to moderate temperatures (Duarte et al. 2018; Lario et al. 2015). One such protease (aspartic protease) purified from psychrotrophic yeast Sporobolomyces roseus was proposed as a potent biocatalyst in the production of soy-sauce and cheese, meat tenderization, and also as bread additive (Bialkowska et al. 2018). In a recent study, cold-active protease from Chryseobacterium polytrichastri isolated from East Rathong Glacier has been reported to produce soybean derived bioactive peptides (Mukhia et al. 2021).

In the present biotechnological era, cold-active proteases can be applied in exciting food bioprocess industries if they can be economically produced at a large scale. There is a need of exploring novel cold-active proteases that can find applications in cheese ripening, recovering heat-sensitive nutrients, such as lipids rich in polyunsaturated fatty acids (Rai et al. 2012, 2017; Hathwar et al. 2011). Cold-active proteases can be applied in the recovery of protein hydrolysates from food processing byproducts that are rich in heat-sensitive lipids and antioxidants (Rai et al. 2013). Furthermore, they can be explored for softening of meat products and winemaking against haze-producing proteins. Due to their tremendous potential in the food industry, novel cold-active enzymes producing isolates from unexplored niches and genomic resources need to be studied. Extensive research is needed on factors affecting the enhanced the production of these cold-active proteases and molecular mechanisms for improvement of the activity to meet their demand at the industrial level.

Esterases and lipases

Esterases, also referred to as carboxyl ester hydrolases, are drawing vast attention for biotechnological applications. They catalyze synthesis and hydrolysis of ester bonds and are classified into lipases (act on lipids) and non-lipolytic esterases (act on water-soluble ester substrates) (Thierry et al. 2017). The curiosity on cold-active lipases is associated with their inherent physiological and structural adaptive mechanisms (Phukon et al. 2020b). Cold-active lipases are becoming a fundamental part of the modern food industry. where they can be used in protein polymerization and gelling in fish, clearing of drains clogged by lipids in food processing, upgrading food texture and modifying flavor, and also in the production of fatty acids and interesterification of fats (Joseph et al. 2008). A cold-active lipase from psychrophilic Pseudomonas fluorescence has been used for the synthesis of flavoring compound butyl caprylate (Tan et al. 1996). Similarly, cold-active lipases from fungal psychrophiles, such as Candida antarctica, C. cylindracea, Hansinuela lanuginose, and Geotrichum candidum, have been applied for the esterification of functionalized phenols giving inlipophilic antioxidants for use in sunflower oil (Buisman et al. 1998; Pandey et al. 1999). Esterases that catalyze simple esters with short-chain fatty acids (e.g., triglycerides) are attracting candidates for use in the cheese ripening process. Several enzymes with cold-adapted esterase activity have been identified from psychrophilic bacteria, such as Pseudoalteromonas spp., and Thalassospira sp., Oleispira sp. (D'Auria et al. 2009; Al-Khudary et al. 2010; Lemak et al. 2012). As a whole, cold-active lipases and esterases offer various advantages as an alternate to the usual biochemical processes in the food industry. However, several challenges have to be surmounted in prioritizing their enhanced utility in diverse biotechnology applications, including the food biotechnology sector, which is discussed further in this review.

Phytases

Phytases are gaining importance in the food and feed industries as they catalyse phosphate removal from phytate, which is considered antinutrient as they bind to divalent minerals (Pable et al. 2019). For decades, they have increasingly been used to enrich absorbable phosphate groups in animal foods, particularly for non-ruminant livestock and fish (Kumar et al. 2012). Numerous studies have reported the isolation and identification of cold-active phytases of psychrophilic microbial origin (Yu et al. 2015; Park and Cho 2011), which can be used in the food manufacturing and processing industry as they mediate phytate degradation increasing the bioavailability of minerals. Because of this property, phytases may also find an application in functional food production (Hamid et al. 2014). Nevertheless, little is known regarding



cold-active phytases real-time application within the food industry.

Pectinases

Pectinases or pectinolytic enzymes are an assorted group of enzymes that collectively break down pectin, a significant polysaccharide present in the plant cell wall (Patidar et al. 2018). Pectinolytic enzymes are classified based on the cleavage site (polygalacturonases, lyase/trans-eliminases including pectinlyase, pectate lyase, and pectin esterases). On the pH range, they may be categorized into alkaline and acid active enzymes. They are an integral part of the food industry and contribute more than 40% of the total share of food enzymes (Adapa et al. 2014). Most pectinases used in the food industry are originated from thermophilic or mesophilic microorganisms. Recently, efforts have been growing to modify the thermophilic enzymes for developing cold-active counterparts because of certain unique features, including their high catalytic activity at cold temperature (< 25 °C) and easy inactivation upon mild heat treatment (Truong et al. 2001; Pulicherla et al. 2011). Cold catalysis offers less energy input in the enzymatic reaction. Cold-active pectinases can be used in efficient fruit juice extraction, settling the suspended particles in the fermented mash, winemaking, secreting polymeric colour pigments, releasing terphenols for aroma and purification of coffee and tea (Blanco et al. 2009; Jayani et al. 2005; Margesin et al. 2007). Few cold-active pectinases have been identified from the psychrophilic microorganisms (Brigisson et al. 2003); however, experimental evidence supporting the use of native pectinases within the food industry is scarce, except some heterologously expressed pectinolytic enzymes that have excellent potential for juice making and winemaking industries (Pan et al. 2014). Cold-active pectinases can be applied in juice clarification at a lower temperature, useful in the prevention of heat-sensitive phytonutrients. Cold-active pectinases have excellent potential in the food industry, and microorganisms from cold regions should be screened for secretion of pectinases, functional at cold temperature and acidic pH.

Other enzymes

Besides the major class of cold-active enzymes increasingly gaining the attention of the food industry, few other enzymes have little been explored; however, they could hold remarkable potential for use in the food industry. Cold-active tannases have been identified in many psychrophilic bacteria and yeasts. They can be used in several ways as their thermophilic counterparts, including the manufacturing of instant tea, fruit juices, beer, and wines (Kasieczka-Burnecka et al. 2007; Yao et al. 2014). β-Mannanases, possessing optimal



activity at low temperature, have been reported from some microbial psychrophiles and can be applied to produce manno-oligosaccharides (Nguyen et al. 2019; Dawood and Ma 2020). β -Mannanases can find application as food and feed additives due to their health improving properties. Psychrophilic microbes are also reported as producers of invertases, an important enzyme used in sucrose hydrolysis that yields an equimolar mixture of glucose and fructose. Cold-active invertases can be used for making confectionery, syrup, infant milk, condensed milk, and beverages (Turkiewicz et al. 2005; Madhusudhan and Raghavarao 2011). There are opportunities for exploring cold-active enzymes from new niches for application in the food processing industries.

Cloning, expression and protein engineering of cold-active enzymes

The elegant specificity of enzymes to catalyze varied sets of reactions makes them essential for biochemical transformation beneficial to humanity. In the beginning, most of the commercial enzymes having industrial applications were produced using native microorganisms. This has restricted the application of enzymes from the native cultivable microorganisms, which are generally produced in low yield (Sarmiento et al. 2015; Santiago et al. 2016). Moreover, the natural enzymes display several complexities at the structural and functional level, limiting their prospective in many biotechnological applications (Huston 2008). Therefore, recombinant expression of these enzymes in heterologous hosts has been preferred as a conventional approach to obtain a high yield for desired enzymes. This has also improved the catalytic efficiency, stereo-selectivity and enzyme stability (Santiago et al. 2016; Duarte et al. 2018). Mesophilic hosts are generally favoured for such recombinant strategies; however, the folding temperature of cold-active enzymes required for their structural and functional integrity may not be compatible with the expressed host (Bjerga et al. 2016; Longwell et al. 2017). To overcome this challenge, incubation temperature is lowered after induction of the desired gene in the host (Feller et al. 1998; Santiago et al. 2016). Besides, the expression of such enzymes from eukaryotic microorganisms such as fungi in bacterial hosts can be cumbersome and substandard, which may be because they lack the ability to secrete extracellular proteins (Duarte et al. 2018). In addition, the post-translational modification mechanism is different from those of prokaryotes, and therefore, eukaryotic microbial hosts must be chosen for heterologous expression of eukaryotic cold-active enzymes (Duarte et al. 2018).

Numerous cold-active enzymes identified from psychrophiles have been successfully expressed in heterologous hosts and hold enormous potential for use in the food industry (Wierzbicka-Wos et al. 2011; Pan et al. 2014). Escherichia coli represents an ideal host for the cold-active enzymes sourced from prokaryotes (Krishna 2002). Quite a few β -galactosidases from cold adaptive bacteria such as Arthrobacter sp. and Paracoccus sp. were cloned and expressed in E. coli, making them exceptional candidates for industrial removal of lactose (Białkowska et al. 2009; Wierzbicka-Wos et al. 2011; Pawlak-Szukalska et al. 2014). Similarly, the gene coding for a cold-active and acidic pectin methyl esterase, isolated from an Antarctic fungus Penicillium chrysogenum PE8F46, was expressed in Pichia pastoris. This enzyme was found to improve the firmness of the pineapple dices, representing its significance in the fruit and vegetable industry (Pan et al. 2014). The DNA segment encoding a novel a-amylase, identified from the Antarctic psychrotolerant fungi Geomyces pannorum, was overexpressed in P. pastoris (Gao et al. 2016). It resulted in high glucose yielding ability, which can be a potential application in the syrup industry. In the same way, a cold-active aspartic protease from G. pannorum was cloned and expressed in Aspergillus oryzae displaying itself a promising biocatalyst for cheese making (Gao et al. 2018).

Signs of progress in recombinant technology have further revolutionized the calibration, development and cost-effective production of customized enzymes that possess some industrial relevance. Cold-active enzymes isolated from psychrophilic hosts can be tailored to convene the process specifications by introducing mutations in the protein in a controlled manner (Bornscheuer et al. 2012). Enzyme engineering can be achieved either by site-directed mutagenesis or by directed evolution strategy. Site-directed mutagenesis is one of the earliest and widely used enzyme engineering techniques based on known structural features of the desired characteristic, such as protein sequence or crystal structures (Coker and Brenchley 2006; Wang et al. 2014a, b). A combined directed and random mutagenesis approach to alter the cold-active β-galactosidase, identified from an Antarctic Arthrobacter sp., has provided interesting mutations leading to increased lactose hydrolysis at low temperature, suggesting its potential use in the dairy industry (Coker and Brenchley 2006). A cold-active endo-1,5-α-L-arabinanase (pectinase) from Paenibacillus polymyxa was engineered using site-directed mutagenesis, which shifted its optimal activity pH towards acidic conditions, making it a promising candidate for pectin extraction from vegetables and fruits, and juice clarification (Wang et al. 2014a, b).

The most successful strategy in engineering novel cold-active enzymes is the directed evolution method that involves random mutagenesis of a gene translating the enzyme of interest. This is chiefly carried out using polymerase chain reaction (PCR) based methods, followed by screening or selection of the protein variants showing desired features from the resulting library of mutants.

Saturation mutagenesis, cassette mutagenesis, error-prone PCR, random-priming recombination, DNA shuffling, and staggered extension process (StEP recombination) are some of the random mutagenesis techniques that have been used so far (Arnold 2001). Alteration in the enzyme properties mainly requires multiple amino acid substitutions simultaneously, producing several protein variants for screening. In addition, present-day high throughput screening methods, including fluorescence-activated cell sorting (Bernath et al. 2004; Becker et al. 2008; Fernandez-Alvaro et al. 2011), permit high-throughput screening and selection of a large number of variants within a short time. Besides, various statistical approaches and bioinformatic methods (e.g., protein structure-activity relationship algorithms) are being used for creating multiple mutations, and at the same time, identifying whether a particular mutation is beneficial or not (Fox et al. 2007). Many cold-active enzymes have been engineered using directed evolution methods, making them relevant for numerous industrial applications (Zhang et al. 2003; Gatti-Lafranconi et al. 2008). Hopefully, these relevant cold-active enzyme variants in the coming days are going to witness a steady growth leading to the manufacturing of value-added nutraceuticals of food and pharmaceutical significance.

Commercial cold-active enzymes used in the food industry

There is a growing demand for processed food products and beverages across the globe, with respect to nutritional excellence as well as favorable taste. This might be because of consumer fondness that shifted their preferences towards healthy food products of high nutritional quality, ultimately promoting the demand for food enzymes. The global market size for food enzymes was estimated at \$1944.8 million in the year 2018 and is predicted to achieve \$3056.9 million by 2025, with an anticipated compound annual growth rate (CAGR) of ~5.5% within the period 2019–2029 (https:// www.persistencemarketresearch.com/market-research/ food-enzymes-market.asp). North American countries such as United States of America, Canada and Mexico are leading consumers of food enzymes. Moreover, Asia-Pacific region is perched to grow with a substantial market share in the food enzyme market globally (https://www.alliedmarketres earch.com/food-enzyme-market). Commercial biocatalysts have several applications in the food industry as they are applied in cheese production, in bread making, maintaining color and clarity of the wine, and reducing its sulfur content (Aehle 2007; Dewan 2014; Sarmiento et al. 2015). The application of enzymes during food processing increases the nutritional quality, flavour, appearance, and taste (Fernandes and Carvalho 2016; Raveendran et al. 2018). Some enzymes



(e.g., lactases, α -amylases, proteases, and lipases) are used as processing aids that act on food components without affecting the nutritional and organoleptic properties of the food. Nevertheless, narrow temperature ranges, pH instability, and side effects, such as allergies associated with a few enzymes, limit food enzymes' market growth.

Cold-active enzymes in food industries are in their infancy and are yet to witness significant growth. The introduction of psychrophilic enzymes in the food processes can be considered as a significant driver that will escalate the growth reconnoitre of food enzymes as they possess unique economic and environmental benefits (Pulicherla et al. 2011; Sarmiento et al. 2015). Therefore, food enzymes are expected to evidence significant adoptions in the coming years, straddling a wide range of materiality and high production efficiency. Nonetheless, very few coldactive enzymes have been developed and commercialized for use in the food market (Table 3). The cold-active xylanase produced by P. haloplanktis, which was reported to be efficient in bread making, is now sold by Puratos (Grand-Bigard, Belgium) (Collins et al. 2002). Selected pectinases that are not considered psychrophilic but display activity at low temperatures are also being used within the food and beverages industries. These enzymes include Novoshape®, Novozymes (pectin methyl esterase from recombinant A. oryzae) (Kitamoto et al. 1999), Pectinase 62L, Biocatalysts (mix of polygalactorunase and pectin lyase from Aspergillus sp.) (Combo et al. 2012) and Lallzyme®, Lallemand (mix of polygalactorunase, pectin esterase and pectin lyase from A. niger) (Zavala-Páramo et al. 2021). Research on potential cold-active enzymes can lead to several commercial biocatalysts in future, having applications in food processing industries.

Challenges and opportunities

Cold-active enzymes are prospective substitutes to their conventional mesophilic and thermophilic matching parts in many ways. They possess high activity at low and moderate temperatures and carry out processes without any apparent loss in their catalytic efficiency, which consequences savings of consumption energy. On the contrary, at low temperatures, sometimes enzyme activity is compromised due to low substrate solubility and decreased substrate specificity (Collins and Margesin 2019; Mangiagalli and Lotti 2021). There are certain challenges associated with their instability at higher temperatures and at alkaline pH, which might impede the potential use of such cold adaptive enzymes. Furthermore, the low diversity of the psychrophilic microorganisms paired with non-specific isolation techniques to study such microbes is a major limiting factor in discovering novel cold-active enzymes and unfolding their biotechnological potential. Nonetheless, with the growing discovery and invention of up-to-the-minute techniques and instrumentation facilities, it is highly doable that the challenges could be conquered. Engineering the active site or the whole enzyme by introducing changes in the amino acid type and its positioning could be considered a potential strategy in enhancing structural and functional stability at ambient temperatures. Similar strategies could also be applied to increase the enzymatic function in the alkaline or acidic pH.

The metagenomic approaches that have been instrumental in identifying novel genes encoding proteins of high stability and activity in wide ranges of temperature and pH would be a valuable loom to clone and characterize the cold adaptive enzymes even from the uncultivable microbial community. Furthermore, undertaking genetic changes at the cellular level in the microorganisms producing the cold-active enzymes would be more hopeful in revitalizing the quality and quantity of the cold-active enzymes to meet the commercial food market's expectations.

Conclusion and future directions

Cold-active enzymes reported so far are distinguished by low stability at high temperatures and demonstrated high catalytic efficiency at low temperatures. Besides their easy inactivation, few additional features, including saving energy

 Table 3
 Selected commercial cold active enzymes used in food industry

	cold detive enzyr	nes used in 1660 industry		
Enzyme name	Brand name	Source microorganism	Enzyme properties	Manufacturing company
Xylanase	Premix X-618	Pseudoalteromonashalo- planktis	Active at temperature range 5–25 °C	Puratos NV, Grand-Bigard, Belgium
Pectin methyl esterase	Novoshape®	Aspergillus oryzae	Active at temperature range 10–60 °C	Novozymes Biopharma, United States
Mixture of polygalactorunase and pectin lyase	Pectinase 62L	Aspergillus sp.	Active at temperature range 10–60 °C	Biocatalysts Ltd, Wales, United Kingdom
Mixture of polygalactorunase, pectin esterase and pectin lyase	Lallzyme®	A. niger	Active at temperature range 5–15 °C	Lallemand, Canada

مدينة الملك عبدالعزيز للعلوم والتقنية KACST and reaction time, and stabilizing thermolabile compounds in the reaction mixture, can be considered exceptional alternates to their thermophilic counterparts for use in many industrial applications. However, a minimal number of coldactive enzymes have been used in real food applications. Their low structural stability and cost constraints associated with isolating and purifying these enzymes have been remaining as a significant holdup. Modern molecular and enzyme engineering techniques have drastically predisposed the quality and productivity of enzymes, though enough efforts have to be undertaken in identifying novel cold-active genes that can be further improved to meet the industrial need. Therefore, extensive investigations are on-demand to explore diverse sources of psychrophilic microorganisms to discover unique and novel cold-active enzymes for diverse applications within the biotechnology industries and potential service to humanity.

Acknowledgements The authors would like to thank Institute of Bioresources and Sustainable Development, Regional Centre, Sikkim and Department of Biotechnology, Govt. of India for the support and encouragement. The manuscript corresponds to manuscript ID IBSD/2020/01/062.

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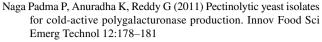
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