

Chapter 1

Antarctic Yeasts: Biodiversity and Potential Applications

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Abstract This review is an attempt in cataloguing the diversity of yeasts in Antarctica, highlight their biotechnological potential and understand the basis of adaptation to low temperature. As of now several psychrophilic and psychrotolerant yeasts from Antarctic soils and marine waters have been characterized with respect to their growth characteristics, ecological distribution and taxonomic significance. Interestingly most of these species belonged to basidiomycetous yeasts which as a group are known for their ability to circumvent and survive under stress conditions. Simultaneously their possible role as work horses in the biotechnological industry was recognized due to their ability to produce novel enzymes and biomolecules such as agents for the breakdown of xenobiotics, and novel pharmaceutical chemicals. The high activity of psychrophilic enzymes at low and moderate temperatures offers potential economic benefits. As of now lipases from *Pseudozyma antarctica* have been extensively studied to understand their unique thermal stability at 90°C and also because of its use in the pharmaceutical, agriculture, food, cosmetics and chemical industry. A few of the other enzymes which have been studied include

extracellular alpha-amylase and glucoamylase from the yeast *Pseudozyma antarctica* (*Candida antarctica*), an extra-cellular protease from *Cryptococcus humicola*, an aspartyl proteinase from *Cryptococcus humicola*, a novel extracellular subtilase from *Leucosporidium antarcticum*, and a xylanase from *Cryptococcus adeliensis*

The ability of these yeasts to adapt to the low temperature conditions has also led to investigations directed towards characterizations of cold stress proteins and heat shock proteins so as to understand the role of these stress protein with respect to adaptation. Antarctic yeasts have also been used as model system to study the inter-relationship among free radicals, antioxidants and UV-induced cell damage.

Keywords Biodiversity, yeast, Antarctica, enzymes, lipase, psychrophilic

1.1 Introduction

The continent of Antarctica which occupies an area of 14 million square kilometers, is a major cold habitat, of which about 99% is covered by ice and snow (Holdgate, 1977). Apart from being very cold, this continent is considered to be a very extreme habitat due to the fact that it is also the driest (Vincent, 1988; Claridge and Campbell, 1977; Campbell and Claridge, 2000), windiest and iciest of all known habitats of the world with high solar radiation at least during the summer season (Smith et al., 1992). Despite these extreme conditions, Antarctica is host to a number of life forms demonstrated by the presence of bacteria, yeasts, fungi, lichens, small invertebrates, many species of birds and mammals (Cameron et al., 1970; Vishniac and Mainzer, 1972; Vincent, 1988; Wynn-Williams, 1990). All these life forms have evolved special mechanisms to overcome the influence of low temperature, high salinity and high radiation.

The microorganisms that thrive in the extreme environment of Antarctica are cold loving and are referred to as psychrophiles. Psychrophilic (cold-loving) organisms differ from the psychrotolerant (cold-tolerant) organisms, on the basis of their cardinal temperatures. Psychrophilic yeasts have an optimum temperature for growth at about 15°C or lower, a maximum up to 25°C but are still capable of growing at 0°C or below (Morita, 1975; Arthur and Watson, 1976); in contrast the psychrotolerant microorganisms are those that are capable of growing at 5°C and below, regardless of whether the optimum temperature was about 15°C or more (van Uden, 1984; Vishniac, 1987). Psychrophiles are unable to grow above 20°C and are widely prevalent in permanently cold habitats, such as in polar regions, at high altitudes or in the deep sea. In contrast, the psychrotolerant which grow over a wider range of temperature and show better growth rates above 20°C are predominant in environments with periodic low temperatures. In Antarctica, psychrophilic and psychrotolerant microorganisms are believed to play key roles in the biodegradation of organic matter and the cycling of essential nutrients (Russell, 1990).

Psychrophilic bacteria, yeasts and other microorganisms define the lower limits of temperature for the survival of life forms. In this context the psychrophilic bacteria and yeasts of Antarctica could serve as excellent model systems to understand the molecular basis of survival at low temperatures. As yet, biological studies in Antarctica have mostly focused on the diversity of bacteria (Shivaji, 2005; Shivaji et al., 2005a; Prabakaran et al., 2006), their taxonomic position (Shivaji et al., 2004; Shivaji et al., 2005b, 2005c), their biotechnological potential (Cavicchioli et al., 2002) and as model systems to understand adaptation of microorganisms to the low temperature (Shivaji et al., 2007; Chintalapati et al., 2006, 2007; Kiran et al., 2004, 2005; Jagannadham et al., 1991, 2000; Chattopadhyay et al., 1997; Ray et al., 1994a, b, c). However, similar studies on yeasts are very limited. This review focuses primarily on the diversity and cataloging of yeasts from Antarctica and their biotechnological potential.

1.2 Diversity of Yeasts in Antarctica

Yeasts are a versatile group of eukaryotic microorganisms which are heterogeneous in their nutritional abilities and are capable of surviving in a range of habitats (Lachance and Starmar, 1988) such as in deep sea (Seiburth, 1979; Fell, 1976), moist and uneven surfaces including polluted waters (Hagler and Ahearn, 1987), on dry substrates and in the presence of high concentrations of salt and sugar (Ingram, 1958). Turkiewicz et al. (2003) suggested that yeasts may be better adapted to low temperatures than bacteria. Therefore, it is not surprising that yeasts belonging to genera such as *Bullera*, *Candida*, *Cryptococcus*, *Cystofilobasidium*, *Debaryomyces*, *Kondoa*, *Leucosporidium*, *Metschnikowia*, *Mrakia*, *Pseudozyma*, *Rhodotorula*, *Sakaguchia*, *Sporopachydermia*, *Symptodiomyces* and *Trichosporon* have been identified in various habitats of Antarctica.

1.2.1 Distribution of Yeasts in Antarctica

The entire Antarctic region is cold and therefore the distribution of yeasts in Antarctica if dependent only on temperature one should be able to see yeasts uniformly distributed. But this is not so. The most northern and southern sampling sites in Antarctica are separated by 10° of latitude. Despite this separation, psychrophiles appeared to be random in their distribution and did not increase with latitude (di Menna, 1960, 1966a). The possible reasons for not obtaining yeasts from some Antarctic samples could be due to the fact that the isolation methods were unsuitable, the incubation temperatures being too high or too low, the incubation time too short, or the medium is too acidic or because of too low osmotic pressure (di Menna, 1966a). Yeasts were usually found in substrates which are acidic rather than alkaline, but inspection of the results showed that high pH values were

not in themselves inhibitory. It was also observed that the yeasts found in Antarctic soils appeared to be dependent on plants. Babyeva and Golubev (1969) isolated more yeasts at 5°C than at higher temperature and showed that forty percent of their 63 isolates were “obligate psychrophiles”, failing to grow above 20°C. Vishniac (1996) concluded that the biodiversity of yeasts and filamentous fungi in terrestrial Antarctic ecosystems increases with the availability of water and energy. Further it was also suggested that yeasts predominate in continental Antarctica compared to maritime and sub-Antarctic habitats (Vishniac, 1996).

1.2.2 Survival of Yeasts in Antarctica

Over the years attempts have been made to understand as to how psychrophilic yeasts survive at low temperatures (<20°C) (Inniss, 1975; Larkin and Stokes, 1968). On the basis of melting points of major fatty acids present in yeasts, it was proposed that the psychrophilic yeasts would be able to grow at temperatures as low as -10°C. Further thermotolerance to temperatures > 20°C may be attributed to the capacity of these yeasts to synthesize heat shock proteins (hsp) and (or) trehalose accumulation as in *Mrakia frigida*, *Leucosporidium fellii* and *L. scottii* but not in *L. antarcticum* (Deegenars and Watson, 1997, 1998). In fact based on these studies it was speculated that hsp 110 may play a role in stress tolerance in psychrophilic yeasts, similar to that of hsp 104 in mesophilic species.

1.2.3 Lipid Composition of the Membranes and Psychrophily

Several studies have clearly indicated that the ability to modulate membrane fluidity by regulating the synthesis of fatty acids is very crucial for low temperature adaptation (Shivaji et al., 2007; Chintalapati et al., 2005). As a thumb rule, low growth temperature increases the proportion of unsaturated fatty acids compared to the saturated fatty acids. This phenomenon applies to bacteria or yeasts (Shivaji et al., 2007; Chintalapati et al., 2005; Sato and Murata, 1980; Sato et al., 1979; Murata et al., 1992; Wada and Murata, 1990; Arthur and Watson, 1976) and in several species of psychrophilic yeasts the unsaturated fatty acids, constituted 50–90% of the total fatty acid composition as in species of *Mrakia*, *Candida*, *Torulopsi*, *Leucosporidium*, and *Cryptococcus* (Watson, 1987; Thomas-Hall and Watson, 2002). Sabri et al. (2001) showed that the inability of *Rhodotorula aurantiaca* to grow at temperatures close to 20°C was due to high accumulation of myristoyl-CoA (C₁₄-CoA), (28-fold higher than in cells cultivated at 0°C temperature). Silver et al. (1977), observed that the cessation of growth at temperatures above 20°C in the psychrophilic yeast *Leucosporidium stokesii* is due to the inability of the yeast to complete an event(s) associated with nuclear division

such as DNA synthesis and normal cell division cycle (Silver and Sinclair, 1979). Meyer et al. (1975) observed that the psychrophilic yeasts are more sensitive to freeze-thaw cycles compared to mesophilic yeasts.

1.2.4 Yeasts of the Genus *Cryptococcus*

The abundance of yeast in Antarctica varies depending on the habitat. In fact even in the same habitat, such as soil, the number varied from total absence to as many as 100,000 yeasts per gram of soil (di Menna, 1966a). *Cryptococcus* is the most predominant group of yeasts in the Antarctic. In this genus *C. laurentii* and *C. albidus* are more predominant compared to *C. luteolus* and *C. diffluens* (di Menna, 1966a). Several new species of *Cryptococcus* have been reported from various habitats in Antarctic such as *Cryptococcus friedmannii* from an Antarctic cryptoendolithic community (Vishniac, 1985a); *Cryptococcus vishniacii* (Vishniac and Hempfling, 1979a, b; Vishniac and Baharaeen, 1982), *Cryptococcus antarcticus* (Vishniac and Kurtzman, 1992; Vishniac and Onofri, 2003), *Cryptococcus albidosimilis* (Vishniac and Kurtzman, 1992), *Cryptococcus socialis* (Vishniac, 1985b), and *Cryptococcus consortionis* (Vishniac, 1985b) from Arctic soils; *Cryptococcus victoriae* (Montes et al., 1999), *Cryptococcus adeliensis*, *Cryptococcus albidus*, *C. laurentii* and *Candida oleophila* (Scorzetti et al., 2000; Pavlova et al., 2001) from mosses and lichens; *Cryptococcus nyarrowii* and *Cryptococcus statzelliae* from soil and snow samples (Thomas-Hall et al., 2002). Some strains of yeasts belonging to the same species appeared to be very different morphologically. Interestingly *Cryptococcus nyarrowii* was represented by two different coloured strains CBS 8804^T (pink colonies) and CBS 8805 (yellow colonies). Other yeast strains (CBS 8908, CBS 8915 and CBS 8920) such as *Cryptococcus victoriae*, *Cryptococcus waticus* sp. nov. (CBS 9496^T) were also isolated from samples collected from the Vestfold Hills, Davis Base (Guffogg et al., 2004).

Cryptococcus laurentii and *C. albidus* are considered as ubiquitous, and are reported by almost all investigators from Antarctica. This could be due to incorrect delineation of these species, as several tests used for identifying them are variable (Fell and Statzell-Tallman, 1998; Barnett et al., 2000; Takashima et al., 2003; Fonseca et al., 2000; Sugita et al., 2000). Sequence analysis of D1/D2 domain of the large subunit rRNA gene and the ITS region has resulted in description of several new species of *Cryptococcus* which were earlier thought to be either *C. albidus* or *C. laurentii*, based on phenotypic methods (Takashima et al., 2003; Middelhoven, 2005). It is also difficult to discriminate *Cryptococcus laurentii* from *C. cellulolyticus*, *C. flavus*, *C. humicola* and *C. hungaricus* based on physiological characters (Barnett et al., 2000). Sugita et al. (2000) reported genetic diversity in the ITS and D1/D2 regions among the clinical isolates of *C. laurentii* and 10 isolates examined in that study were found to belong to seven different species. Similarly, Fonseca et al. (2000) examined several strains of "*Cryptococcus albidus*", using sequence

analysis of the D1/D2 domain of large subunit rRNA gene and established eight new species.

According to Vincent (1988), the *Cryptococcus* yeasts recovered in Antarctic lakes were clearly the result of wash-in from adjacent soils. Moreover, polar soil yeasts, which occur in significant numbers, were found mostly in soil samples that also contain moss, lichen or microalgal material. Vishniac (1995) demonstrated that *Cryptococcus albidus*, a dominant soil organism, was capable of rapid growth when introduced into autoclaved soil, following which viability was retained for 2 months. It was suggested that sterilization altered the nutritional value of the soil in a manner similar to natural weathering factors. Consistent with this, the growth of indigenous soil yeasts would be a function of the frequency and intensity of disturbances of the soil.

1.2.5 Yeasts of Other Genera

Yeasts belonging to the genus *Candida* appear to be quite common in Antarctica but not as predominant as the *Cryptococcus* yeasts. Several strains of *Candida* spp. such as *Candida nivalis*, *Candida gelida* and *Candida frigida*, presently known as *Mrakia frigida* (di Menna, 1966b), *Candida humicola*, *Candida famata*, *Candida ingeniosa* and *Candida auriculariae* (Ray et al., 1989) and *Candida oleophila*. (Pavlova et al., 2001) have been isolated from soil and moss. *Candida (Torulopsis) austromarina* (Fell and Hunter, 1974) has been reclassified as *Candida sake* on the basis of identity of the D1/D2 regions of rDNA. (Kurtzman and Robnett, 1998). All these yeasts were found to be psychrophilic. *Candida* isolates were also identified in various other habitats of Antarctica such as in water, associated with algae, penguin dung etc. (Goto et al., 1969). Other yeasts isolated from Antarctica include *Leucosporidium* (Fell et al., 1969), *Debaryomyces hansenii* (Biswas et al., unpublished results), *Rhodotorula rubra*, (Ray et al., 1989), *Rhodotorula minuta* (Pavlova et al., 2001), *Rhodotorula mucilaginosa* (Pavlova et al., 2001), *Bullera alba* (Ray et al., 1989), *Mrakia frigida* (Biswas et al., unpublished results) and *Mrakia psychrophila* closely related to *Mrakia frigida* (Xin and Zhou, 2007).

1.3 Antarctic Yeasts in Culture Collections

It is interesting to note that about 90% of the yeasts isolated from Antarctica are of basidiomycetous origin (Table 1.1).

The Centraalbureau voor Schimmecultures (CBS), Utrecht, Netherlands has 125 Antarctic yeast strains and the American Type Culture Collection (ATCC), USA has 18 Antarctic yeast cultures, including type strains of nine species of *Cryptococcus* (Table 1.1). Based on the sequence analysis of D1/D2 domain of 26S rRNA gene (Fell et al., 2000) and ITS regions (Scorzetti et al., 2002) these nine type strains

Table 1.1 Antarctic yeast strains available at The Centraalbureau voor Schimmecultures, Utrecht, the Netherlands

Accepted scientific name	CBS accession number	Habitat	Site of collection
<i>Candida davisi</i> Guffogg et al.	CBS 9495	Soil	Antarctica, Davis base, Vestfold Hills, Moss Cirque
<i>Candida parapsilosis</i> group II	CBS 8548	-	Antarctica
<i>Candida psychrophila</i> (S. Goto et al.) S.A. Meyer & Yarrow	CBS 5956	Dung of penguin	Antarctica, Ross Island, Cape Royds
<i>Candida sake</i> (Saito & Oda) van Uden & H.R. Buckley	CBS 5957	Stream water	Antarctica, Lake Bonney
<i>Cryptococcus adeliensis</i> Scorzetti et al.	CBS 8351	Decayed algae	Antarctica, Dumont d'Urville base
<i>Cryptococcus albidosimilis</i> Vishniac & Kurtzman	CBS 7711	Soil	Antarctica, South Victoria Land, Wright Valley, Linnaeus Terrace
<i>Cryptococcus albidus</i> (Saito) C.E. Skinner et al. var. <i>albidus</i>	CBS 9809	Soil	Antarctica, Victoria Land, Edmonson Point
<i>Cryptococcus antarcticus</i> Vishniac & Kurtzman var. <i>antarcticus</i> Vishniac & Kurtzman	CBS 7687	Soil	Antarctica, University Valley
<i>Cryptococcus antarcticus</i> Vishniac & Kurtzman var. <i>circumpolaris</i> Vishniac & Onofri	CBS 7689	Soil	Antarctica, University Valley
<i>Cryptococcus consortionis</i> Vishniac	CBS 7159	Soil	Antarctica, South Victoria Land, Linnaeus Terrace
<i>Cryptococcus friedmannii</i> Vishniac	CBS 7160	Soil	Antarctica, Ross Desert
<i>Cryptococcus humicola</i> (Daszewska) Golubev	CBS 5958	Water	Antarctica, Lake Vanda
<i>Cryptococcus mycelialis</i> Golubev, V.I. & Golubev, N.V	CBS 7712	Soil	Antarctica, East Falkland Island
<i>Cryptococcus nyarrowii</i> Thomas-Hall & Watson	CBS 8805	Soil and lichen	Antarctica, Lichen Valley, Vestfold Hills, Davis base
<i>Cryptococcus nyarrowii</i> Thomas-Hall & Watson	CBS 8804	Bird	Antarctic, Lichen Valley, Vestfold Hills, Davis base
<i>Cryptococcus socialis</i> Vishniac	CBS 7158	Soil	Antarctica, South Victoria Land, Linnaeus Terrace
<i>Cryptococcus victoriae</i> Montes et al.	CBS 8685	Soil	Antarctica, Victoria Land
<i>Cryptococcus vishniacii</i> Vishniac & Hempfling var. <i>vishniacii</i>	CBS 6808	Soil	Antarctica, Mount Baldr

(continued)

Table 1.1 (continued)

Accepted scientific name	CBS accession number	Habitat	Site of collection
<i>Cryptococcus waticus</i> Guffogg et al.	CBS 9496	Soil	Antarctic, Davis base, Vestfold Hills, Watts Lake
<i>Cystofilobasidium bisporidii</i> (Fell et al.) Oberwinkler & Bandoni	CBS 6346	Sea water	Antarctic Ocean
<i>Cystofilobasidium capitatum</i> (Fell et al.) Oberwinkler & Bandoni	CBS 6358	Zooplankton	Antarctic Ocean
<i>Cystofilobasidium infirmominiatum</i> (Fell et al.) Hamamoto et al	CBS 6350	Zooplankton	Antarctic Ocean
<i>Kondoa malvinella</i> (Fell & Hunter) Y. Yamada et al.	CBS 6082	Sea water	Antarctica
<i>Leucosporidium antarcticum</i> Fell et al.	CBS 5942	Sea water	Antarctica, Weddell Sea off Joinville Island
<i>Leucosporidium scottii</i> Fell et al.	CBS 5930	Sea water	Antarctica
<i>Metschnikowia australis</i> (Fell & Hunter) Mendonça-Hagler et al.	CBS 5847	Sea water	Antarctic Ocean
<i>Metschnikowia koreensis</i> Hong et al.	CBS 9068	-	Antarctica
<i>Mrakia frigida</i> (Fell et al.) Y. Yamada & Komagata	CBS 5266	Soil	Antarctica, Scott Base
<i>Rhodospiridium sphaerocarpum</i> S.Y. Newell & Fell	CBS 5939	Sea water	Antarctica, Marguerite Bay
<i>Rhodotorula minuta</i> (Saito) F.C. Harrison var. <i>minuta</i>	CBS 9810	Soil	Antarctica, Victoria Land, Edmonson Point
<i>Rhodotorula</i> sp. F.C. Harrison	CBS 8940	Water	Antarctica, Chelnok lake
<i>Sakaguchia dacryoidea</i> (Fell et al.) Y. Yamada et al.	CBS 6353	Sea water	Antarctic Ocean
<i>Sporopachydermia lactativora</i> Rodrigues de Miranda	CBS 5771	Sea water	Antarctic Ocean
<i>Sympodiomyces parvus</i> Fell and Statzell-Tallman	CBS 6147	Sea water	Antarctic Ocean
<i>Trichosporon pullulans</i> (Lindner) Diddens & Lodder	CBS 5108	Soil	Antarctica

were identified as being synonyms to *Cryptococcus vishniacii* var. *vishniacii*. The Microbial Type Culture Collection and Gene Bank (MTCC) in India has 25 Antarctic yeasts in its collection, isolated from the Schirmacher Oasis region of Antarctica (Ray et al., 1989).

1.4 Are Antarctic Yeasts Endemic?

The larger question in microbial ecology is whether microbes are endemic? The continent of Antarctica due to its remoteness and isolation from the remaining landmass of the earth for millions of years should be amongst the first places to look for endemic organisms and also to examine the evolutionary processes that can give rise to microbial speciation. *Cryptococcus antarcticus* and *C. vishniacii* occur in Antarctica and as of now are unknown outside Antarctica (Vishniac, 1999). But this may not be sufficient evidence in support of endemism since many other yeasts are widely distributed. *Candida antarctica* (reclassified as *Pseudozyma antarctica*), was first isolated from Antarctica (Goto et al., 1969); but later it was identified from Japanese natural samples and from flowers in India (Saluja and Prasad, unpublished observations). Similarly, *Cryptococcus victoriae*, first reported from Antarctica (Montes et al., 1999) is also found in flower and soil samples in India (Saluja and Prasad, unpublished observations). The yeast genus *Leucosporidium* originally isolated from Antarctica was later isolated from temperate climates (Summerbell, 1983). However, the species of the genus *Mrakia* seems to be confined to cold habitats. Besides, Antarctica it has been reported from other cold habitats such as European Alps (Margesin et al., 2005), Hokkaido, Japan (Nakagawa et al., 2004), glacial and subglacial waters of northwest Patagonia, Argentina (Brizzio et al., 2007), Western Siberia (Poliakova et al., 2001) and Tinto river in southwestern Spain (Lopez-Archilla et al., 2004). A new species *Mrakia curviuscula* was isolated from forest substrates collected in the central part of European Russia (Bab'eva et al., 2002). It appears that organisms are extremely versatile in their adaptive capabilities and therefore would break the shackles of endemism and attain an ubiquitous distribution.

1.5 Biotechnological Potential of Antarctic Yeasts

1.5.1 Enzymes from Antarctic Yeasts

Bioprospecting for biomolecules such as enzymes, pigments, polyunsaturated fatty acids etc. from psychrophilic yeasts has gained momentum with the realization that these yeasts due to their unique ability to survive and grow at low temperatures

Table 1.2 Enzymes produced by Antarctic yeasts

Enzyme	Yeast	Reference
Proteinase	<i>Cryptococcus friedmannii</i>	Vishniac, 1985
Serine proteinase	<i>Leucosporidium antarcticum</i>	Turkiewicz et al., 2003
Aspartyl proteinase	<i>Candida humicola</i>	Ray et al., 1992
Xylanase	<i>Cryptococcus adeliensis</i>	Gomes et al., 2000; Petrescu et al., 2000
Xylanase	<i>Cryptococcus albidosimilis</i> (<i>Cryptococcus albidus</i> TAE85)	Amoresano et al., 2000
Lipase	<i>Leucosporidium antarcticum</i>	Turkiewicz et al., 2003
Lipases A and B	<i>Pseudozyma antarctica</i> (<i>Candida antarctica</i>)	
α -Glucosidase	<i>Leucosporidium antarcticum</i>	Turkiewicz et al., 2003
α -Amylase	<i>Candida antarctica</i>	De Mot and Verachtert, 1987
Glucosamylase	<i>Candida antarctica</i>	De Mot and Verachtert, 1987
Acid phosphatase	<i>Leucosporidium antarcticum</i>	Turkiewicz et al., 2003
Alkaline phosphatase	<i>Leucosporidium antarcticum</i>	Turkiewicz et al., 2003
Beta-fructofuranosidase	<i>Leucosporidium antarcticum</i>	Turkiewicz et al., 2003

would be producing enzymes which are cold active and also other biomolecules so as to facilitate their survival at low temperatures. Cold active enzymes may provide interesting clues that would add to our understanding of the relationship between structure, stability and activity of enzymes at low temperatures (Gerday et al., 1997). Most biological systems show 2–3 times reduced reaction rate when the temperature is decreased by 10°C. Enzymes from psychrophilic microorganisms are thought to have evolved a more flexible structure when compared to their mesophilic and thermophilic counterparts. This character probably originates from weakening of intramolecular interactions and is supposed to be responsible for the increased catalytic efficiency and the low thermal stability of psychrophilic enzymes in general (Feller and Gerday, 1997). Several different types of enzymes have been characterized from psychrophilic yeasts (Table 1.2).

1.5.2 Lipases

Two lipases from *Pseudozyma antarctica* (*Candida antarctica*) namely CAL-A and CAL-B have been patented and used for various processes such as preparation of optically active amines, acid ethyl esters, triglycerides, alkyl ester derivatives of restaurant grease (Hsu et al., 2003), hydrolysis of fats, hydrolysis of water insoluble esters of fats, hydrolysis of a mixture of (chloromethyl-dimethylsilyl)-2-propenyl acetate (Rubio et al., 2001), synthesis of polyesters etc. which are useful to the detergent, food, pharmaceutical and other industries (UNEP report on Antarctic bioprospecting, 2004). Thus both these lipases have extensive applications (de Maria et al., 2005) and CAL-A is considered as the most thermostable lipase known, being able to work efficiently even at above 90°C

(Anderson et al., 1998; Kirk et al., 2002). The biotechnological applications of *Candida antarctica* lipase has been reviewed (de Maria et al., 2005) and detail properties related to the catalytic properties (Passicos et al., 2004) substrate specificity (Raza et al., 2001; Larios et al., 2004; Arsan and Parkin, 2000) and thermostability have been studied (Anderson et al., 1998; Kirk and Christensen, 2002). The immobilized form of CAL-B is thermostable even under non-aqueous conditions (Arroyo et al., 1998; Koops et al., 1999). Recently, DNA shuffling was used to create chimeric CAL-B with improved activity toward the hydrolysis of diethyl 3-(3',4'-dichlorophenyl) glutarate (DDG) (Suen et al., 2004). Three variants of the *Candida antarctica* CAL-B lipase have been constructed and it was found that the variant containing the T103G mutation, that introduces the consensus sequence G-X-S-X-G found in most other known lipases, showed increased thermostability but retained only half the specific activity of the native enzyme (Patkar et al., 1998).

1.5.3 Xylanases

Antarctic yeast *Cryptococcus adeliensis* produces a cold-adapted xylanase (Scorzetti et al., 2000; Gomes et al., 2000). In addition to xylanase, this strain also showed activities of endoglucanase, β -mannanase, β -xylosidase, β -glucosidase, and α -L-arabinofuranosidase enzymes. The authors observed that the broad pH and temperature ranges suggest that the xylanase of *C. adeliensis* exists in multiple forms, but could not determine the isoenzymic composition of the crude xylanase. They predicted that the very low thermal stability of the *C. adeliensis* xylanase is most probably the result of increased protein flexibility. Petrescu et al. (2000) studied a xylanase of *Cryptococcus adeliensis* which shared 84% identity with its mesophilic counterpart from *Cryptococcus albidus*, but was less thermostable than its mesophilic homologue. The cold-adapted xylanase displayed a lower activation energy and a higher catalytic efficiency in the range of 0–20°C. These observations suggested a less compact, more flexible molecular structure. Molecular modeling indicated that the adaptation to cold consists of discrete changes in the three-dimensional structure that may lead to a less compact hydrophobic packing, to the loss of one salt bridge, and destabilization of the helices. The structural characterization of the xylanase from the psychrophilic antarctic yeast *Cryptococcus albidosimilis* (*C. albidus* TAE85), showed that it is a glycoprotein made up of 338 amino acids (Amoresano et al., 2000) and has both the N- and O-linked glycans and suggested that the glycosylation system in cold-adapted organisms might have similarities as well as differences with respect to mesophilic and thermophilic yeasts. Xylanases may be suitable for applications such as digestion of industrial or sewage wastes and decomposition of agricultural residues at low or ambient temperatures. The yeast *C. adeliensis*, with its ability to produce xylanase and other enzymes, may find application as a probiotic inclusion and a therapeutic agent in food.

1.5.4 *Proteases and Other Enzymes*

Ray et al. (1992) examined the extracellular protease from a psychrotolerant dimorphic yeast *Candida humicola*, isolated from Antarctic soil. Secretion of the enzyme was greater during exponential growth and low temperatures than during growth at higher temperatures. The enzyme was active from 0 to 45°C, with optimum activity at 37°C. Turkiewicz et al. (2003) reported an extracellular serine proteinase, lap2, from the psychrophilic antarctic yeast *Leucosporidium antarcticum* 171. This enzyme was a glycoprotein, and was most active at temperatures between 20 to 30°C with an optimum at 25°C. Partial activity of the enzyme was retained at zero (20 to 25% activity) and subzero temperatures (18% activity at -10°C). The proteinase lap2 is the first psychrophilic subtilase in this family.

An α -amylase and a glucoamylase were purified to homogeneity from the culture fluid of β -cyclodextrin-grown *Candida antarctica* CBS 6678 (De Mot and Verachtert, 1987). α -Amylase was active on cyclodextrins, whereas debranching activity was demonstrated for glucoamylase. List of enzymes reported from Antarctic yeast species is given in Table 1.2.

1.6 Other Applications of Antarctic Yeasts

Candida sp. which was isolated from the upper layer of Lake Vanda in the McMurdo Dry Valleys, Antarctica was used for removing nitrogen and nitrate from water samples at low temperature (5°C) (Katayama-Hirayama, et al., 2003). Some yeasts from the Antarctic have been used as model systems to study the effects of UV-radiation (Tsimako et al., 2002) and these attempts could form the basis of future studies to establish inter-relationship among free radicals, antioxidants and UV-induced cell damage. In a recent paper, Libkind et al. (2006) suggested that in Patagonian freshwater yeasts there is an apparent relationship between the ability to produce photoprotective compounds, their tolerance to UV exposure and their success in colonizing habitats highly exposed to UV. Similar mechanism may be in operation in the yeast strains isolated from Antarctica.

1.7 Conclusions

Culture collections are important repositories of microbial biodiversity and are essential for the long-term availability of authentic cultures. They also serve as key sources of taxonomic expertise and are needed for the long-term preservation of strains and organisms for biotechnological research. Unfortunately, many researchers do not understand and appreciate the importance of depositing their cultures in known culture collections, as a result some cultures isolated from exotic locations are lost forever, once the researcher retires or changes his field of research. For this

reason, of the several hundreds of yeast cultures isolated from Antarctica, very limited numbers are available from the culture collections in the World. Extensive research into the biodiversity of Antarctic yeasts from various habitats of Antarctica is essential to establish yeast species richness, to identify various strategies by which they adapt to low temperatures and to unravel the molecular basis of their adaptation to the extreme conditions in Antarctica like low temperature, low water activity and low nutrient availability.

References

- Amoresano, A., Andolfo, A., Corsaro, M.M., Zocchi, I., Petrescu, I., Gerday, C., and Marino, G. 2000. *Glycobiology*, **10**: 451–458.
- Anderson, E.M., Larsson, K.M., and Kirk, O. 1998. *Biocatal. Biotransform.*, **16**: 181–204.
- Arroyo, M., Sanchez-Montero, J.M., and Sinisterra, J.V. 1998. *Enzyme Microb. Technol.*, **24**: 3–12.
- Arsan, J. and Parkin, K.L. 2000. *J. Agric. Food Chem.*, **48**: 3738–3743.
- Arthur, H. and Watson, K. 1976. *J. Bacteriol.*, **128**: 56–68.
- Bab'eva, I.P., Lisichkina, G.A., Reshetova, I.S and Danilevich, V.N. 2002. *Mikrobiol.*, **71**: 526–532.
- Babyeva, I.P. and Golubev, W.I. 1969. *Microbiology*, **38**: 436–440.
- Barnett, J.A., Payne, R.W., and Yarrow, D. 2000. *Yeasts: Characteristics and Identification*, 3rd edn. Cambridge University Press, Cambridge.
- Biswas, K., Shivaji, S. and Prasad, G.S. (unpublished results)
- Brizzio, S. Turchetti, B. Garcia, V. de Libkind, D. Buzzini, P. and van Broock, M. 2007. *Can. J. Microbiol.*, **53**: 519–525.
- Cameron, R.E. King, J. and David, C.N. 1970. Holdgate, M. (ed.), *Antarctic ecology*, vol. 2 In: Academic Press, New York, pp. 702–716.
- Campbell, I.B. and Claridge, G.G.C. 2000. In: Davidson, W., Howard-Williams, C., and Broady, P., (eds.), *Antarctic Ecosystems: models for wider understanding*, Caxton Press, Christchurch, pp. 2332–2340.
- Cavicchioli, R., Siddiqui, K.S., Andrews, D., and Sowers, K.R. 2002. *Curr. Opin. Biotechnol.*, **13**: 253–261.
- Chattopadhyay, M.K., Jagannadham, M.V., Vairamani, M., and Shivaji, S. 1997. *Biochem. Biophys. Res. Commun.*, **239**: 85–90.
- Chintalapati, S., Kiran, M.D., and Shivaji, S. 2005. *Cell Mol. Biol.*, **50**: 631–642.
- Chintalapati, S., Prakash, J.S.S., Gupta, P., Ohtani, S., Suzuki, I., Sakamoto, T., Murata, N., and Shivaji, S. 2006. *Biochem. J.*, **398**: 207–214.
- Chintalapati, S., Prakash, J.S.S., Singh, A.K., Ohtani, S., Suzuki, I., Murata, N., and Shivaji, S. 2007. *Biochem. Biophys. Res. Commun.*, (In press).
- Claridge, G.G. and Campbell, I.B. 1977. *Soil Sci.*, **123**: 337–384.
- de María, P.D., Carboni-Oerlemans, C., Tuin B., Bargeman, G., van der Meer, A.B. and van Gemert, R. 2005. *J. Mol. Catal. B-Enzym.*, **37**: 36–46.
- De Mot, R. and Verachtert, H. 1987. *Eur. J. Biochem.*, **164**: 643–654.
- Deegenaars, M.L. and Watson, K. 1997. *FEMS Microbiol. Lett.*, **151**: 191–196.
- Deegenaars, M.L. and Watson, K. 1998. *Extremophiles*, **2**: 41–49.
- Di Menna, M.E. 1960. *J. Gen. Microbiol.*, **23**: 295–300.
- Di Menna, M.E. 1966a. *Antonie van Leeuwenhoek* **32**: 29–38.
- Di Menna, M.E. 1966b. *Antonie van Leeuwenhoek* **32**: 25–28.
- Fell, J.W. 1976. In: Jones E.B.G. (ed.), *Recent advances in aquatic mycology*, Elek Science, London, pp. 93–124.
- Fell, J.W. and Hunter, I.L. 1974. *Antonie van Leeuwenhoek* **40**: 307–310.

- Fell, J., Boekhout, T., Fonseca, A., Scorzetti, G., and Statzell-Tallman, A. 2000. *Int. J. Syst. Evol. Microbiol.*, **50**: 1351–1371.
- Fell, J.W. and Statzell-Tallman, A. In: 1998. Kurtzman, C.P. and Fell, J.W., (eds.), The yeasts, a taxonomic study, 4 th edn. Elsevier, B.V. Amsterdam.
- Fell, J.W., Statzell, A.C., Hunter, I.L., and Phaff, H.J. 1969. *Antonie van Leeuwenhoek* **35**: 433–442.
- Feller, G. and Gerday, C. 1997. Psychrophilic enzymes: molecular basis of cold adaptation. *CMLS Cell Mol. Life Sci.*, **53**: 830–841.
- Fonseca, A., Scorzetti, G., and Fell, J.W. 2000. *Can. J. Microbiol.*, **46**: 7–27.
- Gerday, C., Aittaleb, M., Arpigny, J.L., Baise, E., Chessa, J.P., Garsoux, G., Petrescu, I., and Feller, G., 1997. *Biochim. Biophys. Acta.*, **1342**: 119–131.
- Gomes, J., Gomes, I., and Steiner, W., 2000. *Extremophiles*, **4**: 227–235.
- Goto, S., Sugiyama, J., and Iizuka, H. 1969. *Mycologia*, **61**: 748–774.
- Guffogg, S.P., Thomas-Hall, S., Holloway, P., and Watson, K. 2004. *Int. J. Syst. Evol. Microbiol.*, **54**: 275–277.
- Hagler, A.N. and Ahearn, D.G. 1987. In: Rose, A.H. and Harrison, J.S. (eds.), The Yeasts, vol. 1 Academic Press, London, UK.
- Holdgate, M.V. 1977. *Philos. T. Roy. Soc. B*, **279**: 5–25.
- Hsu, A.F., Jones, K., Foglia, T.A., and Marmer, W.N. 2003. *Biotechnol. Appl. Biochem.*, **36**: 181–186.
- Ingram, M. 1958. In: Cook A.H., (ed.), The chemistry and biology of yeasts, Academic Press, New York, pp. 603–633.
- Inniss, W.E. 1975. *Annu. Rev. Microbiol.*, **29**: 445–465.
- Jagannadham, M.V., Chattopadhyay, M.K., Subbalakshmi, C., Vairamani, M., Narayanan, K., Mohan Rao, Ch., and Shivaji, S. 2000. *Arch. Microbiol.*, **173**: 418–424.
- Jagannadham, M.V., Jayathirtha Rao, V., and Shivaji, S. 1991. *J. Bacteriol.*, **173**: 7911–7917.
- Katayama-Hirayama, K., Koike, Y., Kaneko, H., Kikuo Kobayash, K., and Hirayama, K. 2003. *Polar Biosci.*, **16**: 43–48.
- Kiran, M.D., Annapoorni, S., Suzuki, I., Murata, N., and Shivaji, S. 2005. *Extremophiles*, **9**: 117–125.
- Kiran, M.D., Prakash, J.S.S., Annapoorni, S., Dube, S., Kusano, T., Okuyama, H., Murata, N., and Shivaji, S. 2004. *Extremophiles*, **8**: 401–410.
- Kirk, O., Borchert, T.V., and Fuglsang, C.C. 2002. *Curr. Opin. Biotechnol.*, **13**: 345–351.
- Kirk, O. and Christensen, M.W. 2002. *Org. Process Res. Dev.*, **6**: 446–451.
- Koops, B.C., Papadimou, E., Verheij, H.M., Slotboom, A.J., and Egmond, M.R. 1999. *Appl. Microbiol. Biotechnol.*, **52**: 791–796.
- Kurtzman, C.P. and Robnett, C.J. 1998. *Antonie van Leeuwenhoek* **73**: 331–371.
- Lachance, M.A. and Starmer, W.T. 1988. In: Kurtzman, C.P. and Fell, J.W. (eds.), The yeasts, a taxonomic study, 4th edn. Elsevier, B.V. Amsterdam.
- Larios, A., Garcia, H.S., Oliart, R.M., and Valerio-Alfaro, G. 2004. *Appl. Microbiol. Biotechnol.*, **65**: 373–376.
- Larkin, J.M. and Stokes, J.L. 1968. *Can. J. Microbiol.*, **14**: 97–101.
- Libkind, D., Dieguez, M.C., Moline, M., Perez, P., Zagarese, H.E., and van Broock, M. 2006. *Photochem. Photobiol.*, **82**: 972–980.
- Lopez-Archilla, A.I., Gonzalez, A.E., Terron, M.C., and Amils, R. 2004. *Can. J. Microbiol.*, **50**: 923–934.
- Margesin, R., Fauster, V., and Fonteyne, P.A. 2005. *Lett. Appl. Microbiol.*, **40**: 453–459.
- Meyer, E.D., Sinclair, N.A., and Nagy, B. 1975. *Appl. Microbiol.*, **75**: 739–744.
- Middelhoven, W.J. 2005. *Antonie Van Leeuwenhoek*. **87**:101–108.
- Montes, M.J., Belloch, C., Galiana, M., Garcia, M.D., Andres, C., Ferrer, S., Tores-Rodriguez, J.M., and Guinea, J. 1999. *Syst. Appl. Microbiol.*, **22**: 97–105.
- Morita R.Y. 1975. *Bact. Rev.*, **39**: 144–167.
- Murata, N., Wada, H., and Gombos, Z. 1992. *Plant Cell Physiol.*, **33**: 933–941.
- Nakagawa, T., Nagaoka, T., Taniguchi, S., Miyaji, T., and Tomizuka, N. 2004. *Lett. Appl. Microbiol.*, **38**: 383–387.

- Passicos, E., Santarelli, X., and Coulon, D. 2004. *Biotechnol. Lett.*, **26**: 1073–1076.
- Patkar, S., Vind, J., Kelstrup, E., Christensen, MW., Svendsen, A., Borch, K., and Kirk, O. 1998. *Chem. Phys. Lipids*, **93**: 95–101.
- Pavlova, K., Grigorova, D., Hristozova, T., and Angelov, A. 2001. *Folia Microbiol. (Praha)*, **46**: 397–401.
- Petrescu, I., Lamotte-Brasseur, J., Chessa, J.-P., Claeysens, M., Devreese, B., Marino, G., and Gerday, C. 2000. *Extremophiles*, **4**: 137–144.
- Poliakova, A.V., Chernov, I.Y., and Panikov, N.S. 2001. *Microbiology*, **70**: 617–622.
- Prabakaran, S.R., Manorama, R., Delille, D., and Shivaji, S. 2006. *FEMS Microbiol. Ecol.*, **59**: 342–355.
- Ray, M.K., Devi, K.U., Kumar, G.S., and Shivaji, S. 1992. *Appl. Environ. Microbiol.*, **58**: 1918–1923.
- Ray, M.K., Seshu Kumar, G., and Shivaji, S. 1994a. *Microbiology*, **140**: 3217–3223.
- Ray, M.K., Seshu Kumar, G., and Shivaji, S. 1994b. *J. Bacteriol.*, **176**: 4243–4249.
- Ray, M.K., Sitaramamma, T., Ghandhi, S., and Shivaji, S. 1994c. *FEMS Microbiol. Lett.*, **116**: 55–60.
- Ray, M.K., Shivaji, S., Rao, N.S., and Bhargava, P.M. 1989. *Polar Biol.*, **9**: 305–309.
- Raza, S., Fransson, L., and Hult, K. 2001. *Protein Sci.*, **10**: 329–338.
- Rubio, C., Latxague, L., Deleris, G., and Coulon, D. 2001. *J. Biotechnol.*, **92**: 61–66.
- Sabri, A., Bare, G., Jacques, P., Jabrane, A., Ongena, M., Heugen, J.C., Van Devreese, B., and Thonart, P. 2001. *J. Biol. Chem.*, **276**: 12691–12696.
- Saluja, P. and Prasad, G.S. (unpublished results)
- Sato, N. and Murata, N. 1980. *Biochim. Biophys. Acta.*, **619**: 353–366.
- Sato, N., Murata, N., Miura, Y., and Ueta, N. 1979. *Biochim. Biophys. Acta.*, **572**: 19–28.
- Scorzetti, G., Fell, J. W., Fonseca, A., and Statzell-Tallman, A. 2002. *FEMS Yeast Res.*, **2**: 495–517.
- Scorzetti, G., Petrescu, I., Yarrow, D., and Fell, J.W. 2000. *Antonie van Leeuwenhoek* **77**: 153–157.
- Seiburth, J. McN. 1979. *Sea Microbes*. Oxford University Press, New York.
- Shivaji, S. 2005. In: Satyanarayana T. Johri B.N. (eds.), *Microbial diversity: current perspectives and potential applications*. I.K. International Pvt. Ltd., New Delhi, pp. 3–24.
- Shivaji, S., Gupta, P., Chaturvedi, P., Suresh, K., and Delille, D. 2005a. *Int. J. Syst. Evol. Microbiol.*, **55**: 1083–1088.
- Shivaji, S., Kiran, M.D., and Chintalapati, S. 2007. In: Gerday C. Glansdorff N. (eds.), *Physiology and biochemistry of extremophiles*, ASM Press, Washington, pp. 194–207.
- Shivaji, S., Reddy, G.S.N., Aduri, R.P., Kutty, R., and Ravenschlag, K. 2005b. *Cell Mol. Biol.*, **50**: 525–536.
- Shivaji, S., Reddy, G.S.N., Raghavan, P.U.M., Sarita, N.B., and Delille, D. 2004. *Syst. Appl. Microbiol.*, **27**: 628–635.
- Shivaji, S., Reddy, G.S.N., Suresh, K., Gupta, P., Chintalapati, S., Schumann, P., Stackebrandt, E., and Matsumoto, G. 2005c. *Int. J. Syst. Evol. Microbiol.*, **55**: 757–762.
- Silver, S.A. and Sinclair, N.A. 1979. *Mycopathologia*, **67**: 59–64.
- Silver, S.A., Yall, I., and Sinclair, N.A. 1977. *J. Bacteriol.*, **132**: 676–680.
- Smith, R.C., Prezelin, B.B., and Baker, K.S. et al. 1992. *Science* **255**: 952–959.
- Suen, W.C., Zhang, N., Xiao, L., Madison, V., and Zaks, A. 2004. *Protein Eng. Des. Sel.*, **17**: 133–140.
- Sugita, T., Takashima, M., Ikeda, R., Nakase, T., and Shinoda, T. 2000. *J. Clin. Microbiol.*, **38**: 1468–1471.
- Summerbell, R.C. 1983. *Can. J. Bot.*, **61**: 1402–1410.
- Takashima, M., Sugita, T., Shinoda, T., and Nakase, T. 2003. *Int. J. Syst. Evol. Microbiol.*, **53**: 1187–1194.
- Thomas-Hall, S. and Watson, K. 2002. *Int. J. Syst. Evol. Microbiol.*, **52**: 1033–1038.
- Thomas-Hall, S., Watson, K., Scorzetti, G. 2002. *Int. J. Syst. Evol. Microbiol.*, **52**: 2303–2308.
- Tsimako, M., Guffogg, S., Thomas-Hall, S., and Watson, K. 2002. *Redox Rep.*, **7**: 312–314.

- Turkiewicz, M., Pazgier, M., Kalinowska, H., and Bielecki, S. 2003. *Extremophiles*, **7**: 435–442.
- United Nations Environment Programme 2004. Industry involvement in Antarctic bioprospecting. Prepared by United Nations University Institute of Advanced Studies, Tokyo, Japan.
- van Uden, N. 1984. *Adv. Microb. Physiol.*, **25**: 195–251.
- Vincent, C.F. 1988. Microbial ecosystems of Antarctica. Cambridge University Press, Cambridge: p. 303.
- Vishniac, H.S. 1987. In: de Hoog G.S., Smith M.T., Weijman A.C.M. (eds.), Proceedings of an international symposium on the perspectives of taxonomy, ecology and phylogeny of yeasts and yeast-like fungi. CBS, Delft; Elsevier Science Publishers, Amsterdam.
- Vishniac, H.S. 1985a. *Int. J. Syst. Bacteriol.*, **35**: 119–122.
- Vishniac, H.S. 1985b. *Mycologia*, **77**: 149–153.
- Vishniac, H.S. 1995. *Microbial Ecol.*, **30**: 309–320.
- Vishniac, H.S. 1996. *Biodivers. Conserv.*, **5**: 1365–1378.
- Vishniac, H.S. 1999. In: Seckbach J. (ed.), Enigmatic microorganisms and life in extreme environments, Kluwer Academic Publishers, The Netherlands. pp. 317–324.
- Vishniac, H.S. and Baharaeen, S. 1982. **32**: 437–445.
- Vishniac, H.S. and Hempfling, W.P. 1979a. *Int. J. System. Bacteriol.*, **29**: 153–158.
- Vishniac, H.S. and Hempfling, W.P. 1979b. *J. Gen. Microbiol.*, **112**: 301–314.
- Vishniac, H.S. and Kurtzman, C.P. 1992. *Int. J. Syst. Bacteriol.*, **42**: 547–553.
- Vishniac, H.S., and Onofri, S. 2003. *Antonie Van Leeuwenhoek*, **83**: 231–233.
- Vishniac, V.W. and Mainzer, S.E. 1972. *Antarct J. US*, **7**: 88–89.
- Wada, H. and Murata, N. 1990. *Plant Physiol.*, **92**: 1062–1069.
- Watson, K. 1987. Rose, A.H. and Harrison, J.S. (eds.), The yeasts, In: 2nd edn., vol.2, Academic Press, London, UK. pp. 41–47.
- Wynn-Williams, D.D. 1990. *Adv. Microbial. Ecol.*, **11**: 71–146.
- Xin, M.X. and Zhou, P.J. 2007. *J. Zhejiang Univ. Sci. B*, **8**: 260–265.