Review

Candida species: current epidemiology, pathogenicity, biofilm formation, natural antifungal products and new therapeutic options


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The incidence of fungal infections has increased significantly, so contributing to morbidity and mortality. This is caused by an increase in antimicrobial resistance and the restricted number of antifungal drugs, which retain many side effects. Candida species are major human fungal pathogens that cause both mucosal and deep tissue infections. Recent evidence suggests that the majority of infections produced by this pathogen are associated with biofilm growth. Biofilms are biological communities with a high degree of organization, in which micro-organisms form structured, coordinated and functional communities. These biological communities are embedded in a self-created extracellular matrix. Biofilm production is also associated with a high level of antimicrobial resistance of the associated organisms. The ability of Candida species to form drug-resistant biofilms is an important factor in their contribution to human disease. The study of plants as an alternative to other forms of drug discovery has attracted great attention because, according to the World Health Organization, these would be the best sources for obtaining a wide variety of drugs and could benefit a large population. Furthermore, silver nanoparticles, antibodies and photodynamic inactivation have also been used with good results. This article presents a brief review of the literature regarding the epidemiology of Candida species, as well as their pathogenicity and ability to form biofilms, the antifungal activity of natural products and other therapeutic options.

Introduction

The incidence and prevalence of invasive fungal infections have increased since the 1980s, especially in the large population of immunocompromised patients and/or those hospitalized with serious underlying diseases (Arendrup et al., 2005; Espinel-Ingroff et al., 2009). Candida species belong to the normal microbiota of an individual’s mucosal oral cavity, gastrointestinal tract and vagina (Shao et al., 2007), and are responsible for various clinical manifestations from mucocutaneous overgrowth to bloodstream infections (Eggimann et al., 2003). These yeasts are commensal in healthy humans and may cause systemic infection in immunocompromised situations due to their great adaptability to different host niches. The genus is composed of a heterogeneous group of organisms, and more than 17 different Candida species are known to be aetiological agents of human infection; however, more than 90% of invasive infections are caused by Candida albicans, Candida glabrata, Candida parapsilosis, Candida tropicalis and Candida krusei (Pfaller et al., 2007). The expanding population of immunocompromised patients that use intravenous catheters, total parenteral nutrition, invasive procedures and the increasing use of broad-spectrum antibiotics, cytotoxic chemotherapies and transplantation are factors that contribute to the increase of these infections (Ortega et al., 2011). The pathogenicity of Candida species is attributed to certain virulence factors, such as the ability to evade host defences, adherence, biofilm formation (on host tissue and on medical devices) and the production of tissue-damaging hydrolytic enzymes such as proteases, phospholipases and haemolysin (Silva et al., 2011b).

Currently, an increase in the number of yeasts that are resistant to antifungal drugs is recognized worldwide; therefore, the use of in vitro laboratory tests may aid the doctor in choosing an appropriate therapy (Ingham et al., 2012). The ability of Candida species to form drug-resistant biofilms is an important factor in their contribution to human disease. As in the vast majority of microbial biofilms (Rajendran et al., 2010), sessile cells within C. albicans biofilms are less susceptible to antimicrobial agents than are planktonic cells (Kuhn & Ghannoum, 2004). The progression of drug resistance within Candida biofilms has been associated with a parallel increase in the maturation process.
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(Sardi et al., 2011). Furthermore, some studies have also shown that biofilms of Candida develop statically in the presence of a minimal matrix and exhibit the same level of resistance to drugs (fluconazole and amphotericin B) as cells grown in a shaker and exhibiting large amounts of matrix (Seneviratne et al., 2008; Sardi et al., 2011). The increase in resistant strains necessitates a search for new targets for new antifungal agents (J. M. White et al., 1998; Sardi et al., 2011). In the present report, the literature on the current epidemiology, pathogenicity, biofilm formation and resistance of Candida species, as well as new therapies, including natural antifungal agents and nanoparticles, is reviewed.

Epidemiology

Several Candida species are commensal and colonize the skin and mucosal surfaces of humans. Critically ill or otherwise immunocompromised patients are more prone to develop both superficial and life-threatening Candida infections (Hasan et al., 2009). Candida infections also constitute the most common fungal infections in AIDS patients (Fidel, 2006; Hasan et al., 2009). These patients predominantly develop oropharyngeal candidiasis, which can lead to malnutrition and interfere with the absorption of medication.

C. albicans is the predominant cause of invasive fungal infections (Horn et al., 2009) and represents a serious public health challenge with increasing medical and economic importance due to the high mortality rates and increased costs of care and duration of hospitalization (Amirante et al., 2005; Lai et al., 2012).

Although C. albicans is the most prevalent species involved in invasive fungal infections, the incidence of infections due to non-albicans species is increasing. In a study with 2019 patients at major North American medical centres, a predominance of non-albicans species was observed; although C. albicans was the most frequently isolated species, it was followed by C. glabrata and other non-C. albicans species. This change in epidemiology could be associated with severe immunosuppression or illness, premature, exposure to broad-spectrum antibiotics and older patients (Horn et al., 2009). In European countries, an analysis showed that more than half of the cases of candidaemia were caused by C. albicans, and the incidence rates for non-albicans candidaemia infections were 14% each for C. glabrata and C. parapsilosis, 7% for C. tropicalis and 2% for C. krusei (Tortorano et al., 2006). Changes in the epidemiology have also been observed in Latin American countries. In Chile, the prevalence of C. albicans has changed, and a progressive increase of non-albicans infection has been observed; C. parapsilosis was the most frequent species, followed by C. tropicalis and C. glabrata. All isolates were susceptible to amphotericin B; however, 50% of the C. glabrata isolates were resistant to fluconazole (Ajenjo H et al., 2011). According to the Brazilian Network Candidaemia Study, C. albicans accounted for 40.9% of cases in Brazil, followed by C. tropicalis (20.9%), C. parapsilosis (20.5%) and C. glabrata (4.9%) (Colombo et al., 2006; Nucci et al., 2010).

Other species have been isolated in healthy people and patients. Candida dubliniensis was usually found in combination with other yeast species, especially C. albicans (Sullivan et al., 2004). A high prevalence of C. dubliniensis in the oral cavities of HIV-infected and AIDS patients has also been reported (Tintelnot et al., 2000; Lasker et al., 2001). Since the first description of C. dubliniensis from the oral cavities of HIV-positive patients from Ireland (Z. Khan et al., 2012), subsequent epidemiological studies have revealed that this species is prevalent globally in association with human (Loreto et al., 2010; Z. Khan et al., 2012) and non-human habitats with a possibility of inter-host transmission (Nunn et al., 2007). The species has now been reported from other body sites/specimens, such as the vagina, urine, skin and gastrointestinal tract of both HIV-positive and HIV-negative patients (Loreto et al., 2010; Mokaddas et al., 2011; Z. Khan et al., 2012). Rarely do patients colonized with this species develop candidaemia (Loreto et al., 2010). The reasons for this limited ability of C. dubliniensis to cause invasive disease have been the focus of recent studies (Jackson et al., 2009). It has been shown that the C. dubliniensis genome lacks important hypha-related virulence genes and that it has a limited ability to undergo yeast-to-hyphal transformation (Moran et al., 2012), which in turn may decrease its potential to invade deeper tissue.

C. parapsilosis has emerged as a significant nosocomial pathogen with clinical manifestations that include endophthalmitis, endocarditis, septic arthritis, peritonitis and fungaemia, usually associated with invasive procedures or prosthetic devices (Cantón et al., 2011), and with neonatal infections in the northern hemisphere, although this species is found in patients of all ages in Latin America (Amirante et al., 2005; Nucci et al., 2010). Pires et al. (2011a) isolated 100 strains of C. parapsilosis from a haemodialysis unit; using molecular analysis, 53% were found to be C. parapsilosis and 47% corresponded to Candida orthopsilosis. Tavanti et al. (2005) suggested that the C. parapsilosis complex could be replaced by three different but related species named C. parapsilosis sensu stricto, C. orthopsilosis and Candida metapsilosis on the basis of differences observed for randomly amplified polymorphic DNA and DNA sequencing of different genes. In the FUNGEMYCA study realized by Cantón et al. (2011), 400 out of 1356 isolates were identified as C. parapsilosis sensu lato (29.5%), and this species was isolated the second most frequently from blood after C. albicans in Spain. Of these 400 isolates, 364 were identified by molecular methods; C. parapsilosis sensu stricto represented 90.7% of isolates, C. orthopsilosis 8.2% and C. metapsilosis 1.1%. Candidaemia due to C. tropicalis has been associated with cancer, especially in patients with leukaemia or neutropenia (Colombo et al., 2007). Candidaemia due to C. glabrata has been reported to be related to the use of fluconazole (Nucci et al., 2010). Candida guilliermondii and Candida rugosa were previously uncommon agents; however, the incidence of these is increasing (Pfaller et al., 2009;
Nucci et al., 2010). C. rugosa (1.1 %) has been described in the oral cavity of diabetic patients (Pires-Gonçalves et al., 2007).

Concern is rising about the high incidence of infections caused by non-\textit{albicans} species and the emergence of antifungal resistance (Pereira et al., 2010). Among the non-\textit{albicans} species, \textit{C. tropicalis} and \textit{C. parapsilosis} are both generally susceptible to azoles; however, \textit{C. tropicalis} is less susceptible to fluconazole than is \textit{C. albicans}. \textit{C. glabrata} is intrinsically more resistant to antifungal agents, particularly to fluconazole. \textit{C. krusei} is intrinsically resistant to fluconazole, and infections caused by this species are strongly associated with prior fluconazole prophylaxis and neutropenia. \textit{Candida lusitaniae}, which accounts for 1–2 % of all candidaemias, is susceptible to azoles but has a higher intrinsic resistance to amphotericin B (Cruciani & Serpelloni, 2008).

There has been a gradual increase in the number of antifungal compounds and classes discovered since the 1990s; these include polyenes, azoles, echinocandins and purine analogues. Due to the increased availability of antifungal drugs, selection has occurred with consequent resistance of these micro-organisms. Doctors retain the option of giving the drugs for prophylaxis, empiric therapy, preventive treatment or while waiting for the disease to be diagnosed, so there is a degree of excessive exposure to these agents (Rodriguez-Tudela et al., 2007).

**Pathogenicity of \textit{Candida} species**

An infection caused by \textit{Candida} is termed candidiasis or candidosis. Mycoses caused by these fungi show a wide spectrum of clinical presentations and can be classified as superficial, as with cutaneous and mucosal infections, to deep, widespread and of high severity, as is the case with invasive candidiasis. According to Colombo & Guimarães (2003), the main transmission mechanism is through endogenous candidaemia, in which \textit{Candida} species that constitute the microbiota of various anatomical sites under conditions of host weakness behave as opportunistic pathogens. Another mechanism for transmission is exogenous, and this occurs mainly through the hands of health professionals who care for patients. Also indicated in the spread of infection are health-care materials, such as contaminated catheters and intravenous solutions (Ingham et al., 2012). \textit{Candida} species are considered important pathogens due to their versatility and ability to survive in various anatomical sites (Calderone & Fonzi, 2001). It was believed decades ago that yeasts passively participated in the process of pathogenesis in the establishment of fungal infection. Thus, organic weakness or an immunocompromised host was considered the only mechanism responsible for the establishment of opportunistic infection. Today, this concept has been modified. The current consensus is that these organisms actively participate in the pathophysiology of the disease process using mechanisms of aggression called virulence factors (Tamura et al., 2007). \textit{Candida} species are eukaryotic opportunistic pathogens that reside on the mucosa of the gastrointestinal tract as well as the mouth, oesophagus and vagina (Kim & Sudbery, 2011; Lim et al., 2012). Although this commensal organism normally colonizes mucosal surfaces in an asymptomatic manner, it can become one of the most significant causes of a disabling and lethal infection (Wisplinghoff et al., 2006; Vincent et al., 2009). In the early 1980s, fungi emerged as major causes of nosocomial infections, mainly affecting immunocompromised patients or those who were hospitalized for long periods due to serious underlying diseases (Vidigal & Svidzinski, 2009). \textit{Candida} species belonging to the microbiota of healthy individuals can be found scattered in the environment. It is believed that most people usually have a single strain of \textit{Candida} in different places in the body for a long period. However, some individuals have more than one strain or species at the same time, and this is commonly observed among hospitalized patients (Klotz et al., 2007b). Moreover, the potential for \textit{Candida} species to become pathogenic should be appreciated. A crucial component of this versatility is the fact that these organisms survive as commensals in diverse and distinct anatomical sites, each with its particular environmental stresses (Vidigal & Svidzinski, 2009). Although \textit{Candida} species can infect different anatomical sites of the human host, evidence exists that immune protection is site-specific for each type. Moreover, cutaneous candidiasis and vaginal infections are more likely to be associated with a phagocytic response involving neutrophils and mononuclear phagocytes (Vidigal & Svidzinski, 2009). The urinary tract is the anatomical site most conducive to the development of infections in hospitalized patients, although this remains a problem of questionable significance (Guler et al., 2006; Vidigal & Svidzinski, 2009). Although most of these infections are of bacterial origin, it is estimated that at least 10 % have fungi as the principal aetiological agent and that \textit{Candida} species are the most frequently isolated (Sobel et al., 2000). Data show the isolation of \textit{Candida} species in 22 % of urine samples from patients admitted to intensive care units (Alvarez-Lerma et al., 2003). The colonization of the respiratory tract by \textit{Candida} species is common in patients receiving mechanical ventilation for periods of longer than 2 days. This occurs due to haematogenous spread or pulmonary aspiration of the contents of colonies of oropharyngeal or gastric origin (Vidigal & Svidzinski, 2009). Prolonged intensive care unit stay and hospitalization are also important. Xie et al. (2008) evaluated the impact of invasive fungal infection in surgical patients with severe sepsis. The authors found that 28.3 % of patients exhibited invasive fungal infection; \textit{C. albicans} was most frequently isolated (58 %), followed by \textit{C. tropicalis} (17 %) and \textit{C. glabrata} (15 %). In addition, the organ most affected by invasive fungal infection was the lung. \textit{Candida} pathogenicity is facilitated by a number of virulence factors, the most important of which are those for adherence to host tissues and medical devices, biofilm formation and secretion of hydrolytic enzymes (e.g.
proteases, phospholipases and haemolysins). Furthermore, although there has been extensive research to identify pathogenic factors in fungi, particularly in *C. albicans*, relatively little is known about non-*albicans* species (Silva *et al.*, 2011a).

Virulence in *C. albicans* and other pathogens includes host recognition, which enables the pathogen to bind to host cells and proteins. Additionally, degradative enzymes play a special role in virulence. Fungal invasion is facilitated more by the transition between yeast cells and filamentous growth than by yeast growth (Cullen & Sprague, 2012).

The primary factor in the fungal colonization of human tissues is adherence to host surfaces; this process is controlled and induced by several cell-signalling cascades in both the fungus and the environment. In addition, *Candida* species can adhere to the surfaces of medical devices and form biofilms. The initial attachment of *Candida* cells is mediated by non-specific factors (hydrophobicity and electrostatic forces) and promoted by specific adhesins present on the surface of fungal cells that recognize ligands such as proteins, fibrinogen and fibronectin (Li *et al.*, 2003). The phenomenon of adhesion is exhibited by specialized surface proteins, called adhesins, that specifically bind to amino acids and sugars on the surface of other cells or promote adherence to abiotic surfaces (Verstrepen & Klis, 2006).

As *Candida* species are part of the human microbiota, they are often found on biomaterials, implants and various types of catheters (Kojic & Darouiche, 2004). Biofilms cause clinical problems of concern because they increase resistance to antifungal therapy; the mechanism of biofilm resistance to antimicrobial agents is not fully known. One hypothesis is that the presence of the matrix restricts the penetration of drugs through the formation of a diffusion barrier (Kojic & Darouiche, 2004), and only the most superficial layers are in contact with lethal doses of antibiotics.

Extracellular hydrolytic enzymes appear to play an important role in adherence, tissue penetration, invasion and the destruction of host tissues (Silva *et al.*, 2011b). The most important hydrolytic enzymes are proteases and phospholipases. Ten aspartic proteinase (Sap) isoenzymes are responsible for the proteinase activity of *C. albicans*. These proteins have molecular masses between 35 and 50 kDa and are encoded by the *SAPI–6* genes. The role of the *SAPI–6* genes is related to adherence, tissue damage and changes in the immune response. The functionality of *SAP7* has not been fully discovered, but reports exist that Sap9 and Sap10 are not secreted proteinases; instead, they are used in preserving the regulatory surface integrity of the yeast cells (Naglik *et al.*, 2003). Sap proteins have also been described in *C. tropicalis*, *C. parapsilosis* and *C. guilliermondii* (Zaugg *et al.*, 2001). Several studies have demonstrated a relationship between an increase in the synthesis and activity of extracellular hydrolytic enzymes and an increase in the pathogenic potential of the yeasts, leading to clinical signs of severe candidiasis (Bramono *et al.*, 2006; Ingham *et al.*, 2012).

Putative roles of microbial extracellular lipases include the digestion of lipids for nutrient acquisition, adhesion to host cells and tissues, unspecific initiation of inflammatory processes by affecting immune cells and self-defence by lysing the competing microflora (Stehr *et al.*, 2004; Gácer *et al.*, 2007). Gácer *et al.* (2007) have shown that lipase inhibitors significantly reduce tissue damage during infection of reconstituted human tissues. Furthermore, biofilm formation was inhibited in lipase-negative *C. parapsilosis* mutants, and their growth was significantly reduced in lipid-rich media. Additionally, lipase-negative mutants were significantly less virulent in infection models.

Additionally, the production of haemolysin plays an important role in virulence. This protein is essential for survival and is related to the acquisition of iron (Vaughn & Weinberg, 1978). Haemolysins are proteins produced by micro-organisms to destroy red blood cells. Iron, an inorganic element, is essential for the development of micro-organisms, including yeasts, and the ability to obtain this element is essential for the establishment of an infectious process (Manns *et al.*, 1994).

Seven phospholipase genes have been identified (*PLA*, *PLB1*, *PLB2*, *PLC1*, *PLC2*, *PLC3* and *PLD1*); however, the role of the enzymes encoded by these remains unclear (Samaranayake *et al.*, 2006). *PLB1* has been described as having a virulence role in animal models of candidiasis. Plb1p is an 84 kDa glycoprotein present at hyphal tips during tissue invasion and has hydrolase and lyso-phospholipase-transacylase activity (Ghannoum, 2000).

*C. albicans* reversibly converts from unicellular yeast cells to either pseudohyphal or hyphal growth, a morphogenesis phenomenon (creating a transition between unicellular yeast cells and a filamentous growth form). *C. albicans* and *C. dubliniensis* form both types of filamentous growth, indicating that these yeasts are capable of growing isotropically (yeast) or apically (hyphal and pseudohyphal).

The growth of hyphae, a virulence mechanism, plays an important function in tissue invasion and resistance to phagocytosis (Jayatilake *et al.*, 2006).

**Candida biofilm and disease**

Biofilm formation by fungi may play an important role in pathogenesis (Martinez & Fries, 2010). Recent evidence suggests that the majority of disease produced by *C. albicans* is associated with biofilm growth (Ramage & López-Ribot, 2005). Studies have shown that the micro-organisms are almost non-existent in their planktonic free form in the tissues of the host, but are grouped together, forming a multicellular community, both in tissues and on prostheses, catheters and other surfaces (Soll, 2008). Adhesion between *Candida* species cells and materials or host cells has been implicated as an early step in biofilm formation. Attachment of *Candida* species cells to materials is mediated by
non-specific interactions, such as hydrophobic and electrostatic forces, as well as by specific adhesin-ligand bonds (Ramage & López-Ribot, 2005; Chaffin et al., 1998; Chandra et al., 2005). Cell hydrophobicity and charge also depend on cell growth morphology and cell surface structure (Kriznik et al., 2005).

Biofilms are specific and organized communities of cells under the control of signalling molecules, rather than random accumulations of cells resulting from cell division (Fig. 1). These biological communities can be embedded in an extracellular matrix that is self-produced. Silva et al. (2009) showed that C. parapsilosis, C. tropicalis and C. glabrata are also capable of producing biofilms. In addition, the extracellular matrix contains different amounts of protein and carbohydrate for different species. Villar-Vidal et al. (2011), comparing biofilm formation by C. albicans and C. dubliniensis, found that biofilm formation by C. albicans isolates was higher than that by C. dubliniensis. In general, the biofilm matrix comprises carbohydrates, proteins, phosphorus and hexosamines; however, environmental conditions such as medium composition, pH and oxygen as well as the fungal strain and species affect biofilm formation and matrix composition. For example, C. parapsilosis biofilms contain large amounts of carbohydrates, and the protein content is low when compared with biofilms of C. glabrata and C. tropicalis (Silva et al., 2009, 2011b; Baillie & Douglas, 1999).

Biofilms can grow on any biotic or abiotic moist surface. The association of organisms in biofilms is a form of protection for their development, encouraging symbiotic relationships and allowing survival in hostile environments (Davey & O’toole, 2000). Biofilms may help maintain the role of fungi as pathogenic by evading host immune mechanisms, resisting antifungal treatment and withstanding competitive pressure from other organisms. Consequently, biofilm-related infections are difficult to treat. Biofilm production is also associated with a high level of antimicrobial resistance of the associated organisms (Ozkan et al., 2005). The first crucial event in the process of biofilm formation is microbial attachment, which is followed by the maturation process of the biofilm (Ramage et al., 2001). After the yeast associates with the substrate, many physical and chemical events enable initial adhesion of the micro-organism to the surface. After this accession stage, the biofilm goes through a maturation process. Hawser & Douglas (1994) reported that isolates of C. parapsilosis and C. glabrata were significantly less likely to produce biofilms than the more pathogenic C. albicans. Most infections result from pathogenic biofilms; therefore, the biomedical significance of biofilms is substantial (Ganguly & Mitchell, 2011; Harriott et al., 2010). Transplantation procedures, immunosuppression, the use of indwelling devices and prolonged intensive care unit stays have increased the prevalence of fungal disease (Mukherjee & Chandra, 2004). Availability to the inside of medical devices, such as central and peripheral catheters, haemodialysis and peritoneal dialysis units and intracardiac prosthetic valves, facilitates biofilm formation (Mukherjee & Chandra, 2004; Chandra et al., 2001; Ramage et al., 2005). Although C. albicans is the fungal species isolated most often, the incidence of other species has recently increased (Medrano et al., 2006), and these have also been described as the aetiological agent in nosocomial infections (Bruder-Nascimento et al., 2010). The ability of C. albicans to form biofilms on medical devices has a profound effect on its capacity to cause human disease (Hawser & Douglas, 1994). Catheter-related infections are the major cause of morbidity and mortality among hospitalized patients, and catheter microbial biofilms are associated with 90% of these infections. Catheter-associated Candida biofilms can lead to bloodstream infections with an approximate incidence of one episode per 100 hospital admissions (DiDone et al., 2011). C. albicans abiotic-surface biofilms consist of two general types of cells: yeast cells and filamentous cells (Finkel & Mitchell, 2011; Ramage et al., 2006). In vitro, the basal biofilm layer is composed of yeast cells from which filamentous cells emanate. These yeast and filamentous cells are embedded in a dense layer of extracellular matrix. The primary component of biofilm matrix is β-glucan (Finkel & Mitchell, 2011; Nett et al., 2007).

Device-associated Candida species infections cause mortality rates as high as 30% (Viudes et al., 2002; Finkel & Mitchell, 2011), and the annual cost of antifungal therapies in the United States alone is estimated at US$2.6 billion (Finkel & Mitchell, 2011). Like the biofilms formed by bacterial pathogens, Candida species biofilms are resistant to many antimicrobial agents; therefore, treatment can require surgical removal and later replacement of the infected device (Finkel & Mitchell, 2011). C. albicans biofilms consist of two main types of cells: small oval yeast-form cells (blastospores) and long tubular hyphal cells. C. albicans biofilms grown in vitro often have a foundation of yeast cells from which a hyphal layer emanates (Douglas, 2003). Extracellular matrix material is also clearly evident and is bound to both yeast and hyphal cells. The matrix is typically interspersed throughout the biofilm.

![Fig. 1. Scanning electron microscopy of a C. albicans biofilm.](image)
Genetic analyses indicate that both yeast cells and hyphae are crucial for biofilm formation, which suggests that each cell type has a unique role in the process (Finkel & Mitchell, 2011; Al-Fattani & Douglas, 2006; Douglas, 2003). C. albicans has the capacity to switch from yeast morphology to hyphal morphology, one of its major virulence determinants (Lo et al., 1997; Ramage et al., 2002a). In Candida, the dimorphic transition from the yeast to the hyphal phase is a crucial step in the formation of biofilms, and mutant strains that are defective in the ability to germinate are unable to form dense and homogeneous biofilms (Baillie & Douglas, 1999; Ramage et al., 2002a; Kruppa et al., 2004; Jabra-Rizk et al., 2006; Brown et al., 1999). The morphological transition from the yeast to the mycelial form (dimorphic switching) is induced by many environmental factors, such as serum, high temperatures (<37 °C) and neutral pH (Brown et al., 1999). Farnesol, which is associated with mycelial development, may be an important regulatory (quorum-sensing) molecule in C. albicans biofilm formation. Ramage et al., (2002c) found that C. albicans biofilm density and morphology were drastically altered by high concentrations of farnesol, most likely as a direct consequence of the inhibitory effect of farnesol on the morphogenetic process. As the concentration of farnesol was decreased, the morphology of the biofilms changed in a dose-dependent manner to a typical hyphal morphology. Microarray analysis of biofilms exposed to farnesol revealed that genes related to drug resistance, cell wall maintenance, cell surface hydrophilicity, iron transport and heat-shock proteins were influenced in addition to genes associated with hyphae formation (Cao et al., 2005; Albuquerque & Casadevall, 2012).

Other studies have shown the same result with other Candida species (C. dubliniensis and C. tropicalis) (Kruppa et al., 2004).

Yi et al. (2011) observed that although a majority (approx. 90%) of C. albicans strains form traditional biofilms that are impermeable to molecules of low and high molecular mass and impenetrable to white blood cells, a minority (approx. 10%) form biofilms that are both permeable and penetrable. The formation of minority-type alternative biofilms is dictated by a change at a single genetic locus, the mating type locus. Homozygous a/a or x/x cells are mating-competent, whereas the heterozygous a/x cells are mating-incompetent. Cells of the mating-incompetent a/x genotype form impermeable, traditional biofilms whereas cells of the mating-competent a/a or x/x genotype form permeable biofilms. The characteristics of a/a and x/x biofilms are consistent with a suggested role in mating by facilitating the transfer of hormone signals through the permeable biofilm. The two types of biofilm are also regulated by different signal transduction pathways: the a/x form is regulated by the Ras1/cAMP pathway, and the a/a or x/x form is regulated by the MAP kinase pathway.

Nobile et al., (2012) combined ‘classical’ genetics, genome-wide approaches, RNA deep sequencing technology and two in vivo animal models to comprehensively map the transcriptional circuitry controlling biofilm formation in C. albicans. The elucidation of this circuit has led to many new predictions about the genes involved in biofilm formation, and a set of these predictions has been validated by confirming the roles of several of these genes in biofilm development.

Many studies have focused on the formation of C. albicans biofilms on implanted vascular catheters because this is a major source of infection (Finkel & Mitchell, 2011). However, biofilms can also be formed on many other devices. A rat denture biofilm model that recapitulates features of denture stomatitis has been described (Nett et al., 2010). Microscopic and microbiological analyses showed the signature features of biofilm development: adherent cells, the presence of extracellular matrix material and high-level drug resistance. The biofilms in another recently described rat model involving subcutaneously implanted catheters also contained characteristic matrix material and an abundant hyphal population (Ricicova et al., 2010). The closely related cell wall proteins Als1 and Als3 might also function in biofilm surface attachment (Finkel & Mitchell, 2011). The expression of Als1 or Als3 in S. cerevisiae promotes binding to several different protein-coated substrates that may resemble the conditioned surface of an implanted device. In addition, a C. albicans mutant lacking both ALS1 and ALS3 is defective in biofilm formation in vitro and in vivo (Nobile et al., 2006). One or more genes in the ALS family are upregulated in biofilm-associated cells when compared with planktonic cells (Chandra et al., 2001). ALS3 expression was shown to be necessary for biofilm formation on silicone substrates in vitro (Zhao et al., 2006), whereas ALS3 expression was not required for biofilm formation in vivo. Raad et al., (2008) demonstrated that Eap1p is a glucan-cross-linked, cell wall-localized protein. In addition, Eap1 mutants exhibited reduced adhesion to plastic surfaces and epithelial cells whereas Eap1p was able to mediate adhesion to yeast cells. EAPI expression was also required for biofilm formation under shear flow in both in vitro and in vivo central venous catheter biofilm models. Biofilms are the most common form of microbial growth in nature, and the National Institutes of Health estimate that biofilms cause the majority of infections in humans (Douglas, 2003; Nett et al., 2010). Frequently, an implanted device, such as an intravascular catheter, is associated with biofilm infections (Ramage et al., 2002b). C. albicans is the fourth and third leading cause of hospital-acquired bloodstream and urinary tract infections, respectively. Up to 70–80% of Candida bloodstream infections are associated with central venous catheters, and the majority of Candida urinary tract infections are associated with bladder catheters (Nett et al., 2007).

There are estimated to be more than 45 million medical devices implanted every year in the United States. Infection of these devices occurs in 60% of patients, and Candida species are responsible for up to 20% of these infections.
antifungal mechanism of the respective drug/drug class or a fungal pathogen, involving either interference with the resistance are classified as either primary or secondary and kill or inhibit the growth of fungal pathogens (Pfaller, 2012; Each of the antifungal classes utilizes a different means to eliminate a multiplicity of species regarding immunity and drug evasion and manipulation of immune responses will most likely aid in the search for novel therapies that are directed to the overexpression of efflux genes, CDRI, CDR2 (the family of ABC membrane transport proteins – the ATP-binding cassette) (Sardi et al., 2011) and MDRI (the Major Facilitator protein family) as well as amino acid substitutions in the enzyme Erg11p (lanosterol 14-α-demethylase), encoded by the gene ERG11. CDR1, CDR2 and other genes are often coregulated and are overexpressed simultaneously; therefore, it is believed that genes regulating this expression can be mutated (Staib et al., 2000).

Given the increase in fungal infections, two triazoles (voriconazole and posaconazole) and three echinocandins (anidulafungin, caspofungin and micafungin) have been approved to treat and prevent these infections (Mattiuazzi & Giles, 2005) (Table 1). Of the three classes of antifungal agents currently in clinical use, only amphotericin B and the echinocandins, e.g. caspofungin, have demonstrated consistent in vitro activity against C. albicans biofilms (Kuhn et al., 2002). Even with these two agents, Candida biofilm-related infections are extremely difficult, if not impossible, to eradicate, and infected medical devices typically must be removed (Montejo, 2011). One of the barriers to the development of new antifungal agents that are active against biofilms is that very few assays are available for either the characterization or identification of new molecules with activity towards biofilms (Ramage et al., 2009). Candida is the fourth most common cause of bloodstream infections in hospitalized patients (Nucci et al., 1998). Up to environmental factors lead to colonization or replacement of a susceptible species with a resistant species. The antifungal effects of polyene and azole antifungals are due to their actions on the fungal cell membrane, whereas echinocandins act by disrupting the fungal cell wall (Pfaller, 2012; Pemán et al., 2009; T. C. White et al., 1998).

The ability of Candida to form drug-resistant biofilms is an important factor in its contribution to human disease. Like the vast majority of microbial biofilms (Rajendran et al., 2010), sessile cells within a C. albicans biofilm are less susceptible to antimicrobial agents than are planktonic cells (Kuhn & Ghannoum, 2004; Rautemaa & Ramage, 2011). The formation of biofilms causes clinical problems of concern because they increase resistance to antifungal therapies, and the mechanism of biofilm resistance to antimicrobial agents is not fully known. One hypothesis for this resistance is the presence of the matrix, which restricts the penetration of drugs by formation of a diffusion barrier (Nett et al., 2011); hence, only the most superficial layers are in contact with lethal doses of antibiotics. Planktonic cells generally rely on irreversible genetic changes to maintain a resistant phenotype, whereas biofilms are able to persist due to their physical presence and the density of the population, which provides an almost inducible resistant phenotype irrespective of defined genetic alterations (Ramage et al., 2012).

Several molecular mechanisms of resistance to antifungal agents in C. albicans have been described. In particular, these include the increased efflux of antifungal agents due to the overexpression of efflux genes, CDRI, CDR2 (the family of ABC membrane transport proteins – the ATP-binding cassette) (Sardi et al., 2011) and MDRI (the Major Facilitator protein family) as well as amino acid substitutions in the enzyme Erg11p (lanosterol 14-α-demethylase), encoded by the gene ERG11. CDR1, CDR2 and other genes are often coregulated and are overexpressed simultaneously; therefore, it is believed that genes regulating this expression can be mutated (Staib et al., 2000).

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40% of patients with intravenous catheters from which Candida has been isolated have underlying fungaemia (Kuhn et al., 2002), and the mortality rate for patients with catheter-related candidaemia approaches 40% (Kuhn et al., 2002).

Although C. albicans is the most commonly isolated fungal species, other species are being isolated with increasing frequency (Kuhn et al., 2002). C. parapsilosis has become the second most commonly isolated fungal organism in several studies (Kuhn et al., 2002). Pires et al. (2011a) isolated 17 environmental isolates of C. parapsilosis from the hydraulic circuit of a haemodialysis centre. This species is of particular concern in critically ill neonates, where it is known to be associated with central lines and parenteral nutrition (Kuhn et al., 2002). Antifungal therapy alone is insufficient for curing these infections, and affected devices generally need to be removed (Nucci et al., 1998; Rex et al., 2000). Removal of these devices has serious implications in the case of infected heart valves, joint prostheses and central nervous system shunts. The reason why device removal is necessary has been, until recently, a mystery. However, several studies have reported a near-total resistance to antifungal agents exhibited by biofilm-associated Candida (Chandra et al., 2001; Nucci et al., 1998). C. albicans biofilms in vivo (Martinez & Fries, 2010; Mukherjee & Chandra, 2004) and in vitro demonstrated higher susceptibility when treated with a combination of amphotericin B and posaconazole than when treated with single drugs (Martinez & Fries, 2010; Tobudic et al., 2010). Combining the use of echinocandins with other drugs that have antifungal activity is becoming an important alternative form of therapy in mycoses caused by fungi that are resistant to standard antifungal monotherapy or in certain biofilm-associated diseases.

Persistor cells are an important mechanism of resistance in chronic infections (Fauvart et al., 2011; Ramage et al., 2012), and these have gathered some attention recently in the context of fungal biofilms (Lewis, 2008; Bink et al., 2011). In C. albicans biofilms, a small subset of yeast cells have been described that are highly resistant to amphotericin B following adhesion, which is independent of the upregulation of efflux pumps and cell membrane composition (LaFleur et al., 2006). The first study described persistor cells in fungi as a subpopulation of highly tolerant cells. In this study, C. albicans persisters were detected only in biofilms and not in various planktonic populations (LaFleur et al., 2006; Ramage et al., 2012). When a biofilm was killed with amphotericin B and reincubated with cells that survived, a new biofilm was produced with a new subpopulation of persisters; this suggests that these cells were not mutants but phenotypic variants of the wild-type. In this clinically relevant scenario, prolonged and inefficient antifungal treatment may be beneficial to the biofilm population, and this effect may be responsible for antimicrobial drug failure and relapsing infections.

### Candida biofilm and new antifungal strategies

The increasing incidence of drug-resistant pathogens and the toxicity of existing antifungal compounds have drawn attention towards the antimicrobial activity of natural products. The small number of drugs available for fungal treatment, most of which are fungistatic, and the emerging resistance to antifungal agents encourage the search for alternative treatments (de Oliveira et al., 2006). Plants are good options for obtaining a wide variety of drugs and have been used in medicine for a long time; in particular, they are extensively used in folk medicine because they represent an economic alternative, are easily accessible and can be applicable to various pathologies (Sardi et al., 2011; Cruz et al., 2007). Plants therefore constitute an excellent source for substances that can be used in the formulation of new antifungal agents (Holetz et al., 2002).

Many studies have focused on biofilms because they are more resistant to antimicrobials and host defences. Rossignol et al. (2011) studied the 18-amino acid cationic, tryptophan-rich, ApoEdpl-W peptide, derived from human ApoE apolipoprotein, which was shown to have antifungal activity against pathogenic yeasts of the Candida genus (except C. glabrata). ApoEdpl-W was active against planktonic cells and early-stage biofilms but less active against mature biofilms, possibly because of its affinity for extracellular matrix β-glucans. Moreover, ApoEdpl-W partially prevented the formation of biofilms on medical devices. Another study by Mandal et al. (2011) demonstrated in vitro growth inhibition of C. tropicalis and disrupted biofilm formation in a concentration-dependent manner with purified Tn-AFP1 (the peptide derived from Trapa natans). That study also demonstrated downregulation of MDR1 and ERG11 gene expression using real-time PCR analysis. However, usnic acid had inhibitory and fungicidal activity against biofilms of C. parapsilosis and C. orthoparapsilosis in an isolated environment (Pires et al., 2011).

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**Table 1. Conventional antifungal drugs and their cellular targets**

Adapted source: Martinez-Rossi et al. (2008).

<table>
<thead>
<tr>
<th>Chemical class</th>
<th>Drug</th>
<th>Target</th>
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<tbody>
<tr>
<td>Azoles</td>
<td>Miconazole</td>
<td>Ergosterol synthesis</td>
</tr>
<tr>
<td></td>
<td>Ketocanazole</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Fluconazole</td>
<td></td>
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<tr>
<td></td>
<td>Itraconazole</td>
<td></td>
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<tr>
<td></td>
<td>Terconazole</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Voriconazole</td>
<td></td>
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<tr>
<td></td>
<td>Posaconazole</td>
<td></td>
</tr>
<tr>
<td>Polynes</td>
<td>Amphotericin B</td>
<td>Ergosterol (membrane function)</td>
</tr>
<tr>
<td>Pyrimidine Echinocandins</td>
<td>Nistatin</td>
<td>DNA and RNA synthesis</td>
</tr>
<tr>
<td></td>
<td>Flucytosine</td>
<td>Glucan synthesis</td>
</tr>
<tr>
<td></td>
<td>Caspofungin</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Micafungin</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Anidulafungin</td>
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</tbody>
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http://jmm.sgmjournals.org
2012). Stanciuc et al. (2011), with the objective to assess the antibacterial and antifungal potential of some Romanian medicinal plants such as Arnica montana, Artemisia absinthium and Urtica dioica, conducted an antimicrobial screening by obtaining plant extracts and testing them on a series of Gram-positive and Gram-negative bacteria and two fungal strains. The extracts presented antimicrobial activity preferentially directed against planktonic fungal and bacterial growth; an effect on biofilm formation and development was shown only against S. aureus and C. albicans. In vitro assays indicated that the studied plant extracts were a significant source of natural alternatives to antimicrobial therapy, thus avoiding antibiotic therapy, the use of which has become excessive in recent years. Several studies have been conducted using natural products to evaluate interference in C. albicans biofilms (Furletti et al., 2011; Taweechaisupapong et al., 2010; Coleman et al., 2010) (Table 2). Pires et al. (2011b) investigated the anticandidal activity of 16 essential oils on planktonic and biofilm cultures of the C. parapsilosis complex. The most active essential oil was cinnamon oil, which showed antifungal activity against C. orthopsilosis and C. parapsilosis in both suspension (MIC=250 and 500 g l⁻¹, respectively) and biofilm [minimum biofilm reduction concentration (MBRC)=1000 and 2000 g l⁻¹, respectively] cultures. Cinnamon oil also inhibited biofilm formation (MBIC) at concentrations above 250 g l⁻¹ for both species tested. Further research is required to evaluate the antifungal potential of various plants against microbial biofilms because the total elimination of biofilms is difficult.

Another promising antifungal strategy is the use of silver nanoparticles (AgNPs). Silver is a well-known antimicrobial agent and has a long history of application in medicine with a well-tolerated tissue response and a low toxicity profile. Silver is more toxic than many other metals against a broad spectrum of sessile bacteria and fungi that colonize plastic surfaces (Nam, 2011).

More recently, the use of AgNPs has been suggested for coating medical titanium implants (Alviano et al., 2005; Huang et al., 2012; Lu et al., 2011; Garcia-Contreras et al., 2011) in the hope of inhibiting biofilm formation and thereby reducing the incidence of microbial infections and rejection.

Silver has been described as being ‘oligo-dynamic’, meaning it has a toxic (bactericidal/fungicidal) effect on living cells even at low concentrations (Dastjerdi & Montazer, 2010). Silver has been shown to interfere with microbial DNA replication within bacteria and fungi. Metal ions such as silver inhibit the multiplication of micro-organisms. Silver ions can also lead to protein denaturation and cell death because of their reaction with nucleophilic amino acid residues in proteins and their attachment to thiol, amino, imidazole, phosphate and carboxyl groups of membrane proteins or enzymes (Percival et al., 2005). AgNPs have also frequently been shown to inhibit yeast growth (Kim et al., 2007; Mastrolorenzo et al., 2000), and their antifungal activity against certain species such as Trichophyton species and Candida species (Kim et al., 2008) is well documented. These studies show evidence for the molecular mechanism of AgNPs ability, whereby AgNPs act on and inhibit a number of oxidative enzymes such as yeast alcohol dehydrogenase through the generation of reactive oxygen species (Kim et al., 2007). This leads to protein inactivation and induces apoptosis via mitochondrial pathways (Hsin et al., 2008). AgNPs have been demonstrated to exhibit a desirable and promising antifungal effect in several studies, with no serious side effects to the host.

Anti-Candida antibodies can reduce the binding of Candida to various surfaces (Rodier et al., 2003; Elguezabal et al., 2004; López-Ribot et al., 2004; Bujdáková et al., 2008). Studies with antibodies have been performed by various authors to test their effects on various fungal and bacterial organisms.

### Table 2. Antifungal activity of natural products against a variety of biofilms

<table>
<thead>
<tr>
<th>Natural products</th>
<th>Biofilm</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>Peptide (18-amino acid cationic ApoEdpL-W)</td>
<td>All species of Candida except C. glabrata</td>
<td>Rossignol et al. (2011)</td>
</tr>
<tr>
<td>Peptide (Trapa natans)</td>
<td>C. tropicalis</td>
<td>Mandal et al. (2011)</td>
</tr>
<tr>
<td>Romanian plant extracts</td>
<td>C. albicans</td>
<td>Stanciuc et al. (2011)</td>
</tr>
<tr>
<td>Coriandrum sativum L. (oil)</td>
<td>C. albicans</td>
<td>Furletti et al. (2011)</td>
</tr>
<tr>
<td>Cinnamon oil</td>
<td>C. parapsilosis</td>
<td>Pires et al. (2011b)</td>
</tr>
<tr>
<td>Usnic acid</td>
<td>C. parapsilosis and C. orthopsilosis</td>
<td>Pires et al. (2012)</td>
</tr>
<tr>
<td>Boesenbergia pandurata and Piper sarmentosum</td>
<td>C. albicans</td>
<td>Taweechaisupapong et al. (2010)</td>
</tr>
<tr>
<td>Saponins</td>
<td>C. albicans</td>
<td>Coleman et al. (2010)</td>
</tr>
<tr>
<td>Polyphenols (green tea)</td>
<td>C. albicans</td>
<td>Evensen &amp; Braun (2009)</td>
</tr>
<tr>
<td>Ocimum americanum L. essential oil</td>
<td>C. albicans</td>
<td>Thaweboon &amp; Thaweboon (2009)</td>
</tr>
<tr>
<td>Caesalpinia ferrea Martius fruits</td>
<td>C. albicans</td>
<td>Sampaio et al. (2009)</td>
</tr>
<tr>
<td>Cassia spectabilis extract</td>
<td>C. albicans</td>
<td>Sangetha et al. (2009)</td>
</tr>
<tr>
<td>Croton cajucara Benth linalool-rich oil</td>
<td>C. albicans</td>
<td>Alviano et al. (2005)</td>
</tr>
<tr>
<td>Garlic extract</td>
<td>C. albicans</td>
<td>Shuford et al. (2005)</td>
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</tbody>
</table>
Fujibayashi et al. (2009) prepared a polyclonal anti-C. albicans antibody in chicken egg yolk (anti-C. albicans IgY) and investigated its in vitro effectiveness in preventing C. albicans adherence and biofilm formation. The results demonstrated a significant reduction in the adherence of C. albicans SC 5314 to human oral epithelial cells in a dose-dependent manner. The same effect was also observed in other Candida species including serotypes A and B. Furthermore, IgY inhibited C. albicans biofilm formation in serum-free medium, but the inhibition was slightly restored in medium conditioned with 10% serum. This demonstrates that anti-C. albicans IgY cross-reacts with various Candida species and may have a protective effect against oral candidiasis and reduce the dissemination of Candida species. This effect may be due to the blocking of the binding of Candida species to host cells. However, this blocking did not play a role when Candida formed a germ tube in the presence of serum. Therefore, anti-C. albicans IgY could be considered for prophylactic immunotherapy under limited conditions.

The application of photodynamic therapy has been investigated regarding its inactivation of micro-organisms that are pathogenic to the human host. Several authors have associated light emitting diodes (LEDs) with other substances (Ribeiro et al., 2012; Chen et al., 2012; Junqueira et al., 2012). Coleman et al. (2010) observed that saponins may preferentially bind to fungal ergosterol compared to cholesterol, and the fungus displayed increased susceptibility to photodynamic inactivation when used in combination with photosensitizer compounds due to the ability of saponins to increase cell permeability, thereby facilitating penetration of the photosensitizers. This result suggests that it may be a suitable chemical scaffold for a new generation of antifungal compounds.

Dovigo et al. (2011) described the combined use of curcumin with LEDs for the inactivation of C. albicans in both planktonic and biofilm forms. Suspensions of Candida were treated with nine curcumin concentrations and exposed to LEDs at different fluences. The photodynamic effect was greatly increased in the presence of curcumin.

S. Khan et al. (2012) studied gold nanoparticle-enhanced photodynamic therapy including the use of methylene blue against recalcitrant pathogenic C. albicans biofilms. Antibiofilm assays and microscopy studies showed significant reduction of biofilm; therefore, gold nanoparticle conjugate-mediated photodynamic therapy may be used against nosocomially acquired refractory C. albicans biofilms.

Conclusions

The increased incidence of systemic mycoses caused by Candida species in hospitalized patients is an important cause of morbidity and mortality worldwide, especially in critically ill patients. Biofilms play an important role in the perpetuation of these infections primarily with respect to their ability to adhere to various medical devices. The interest in natural products has increased regardless of whether they are associated with other therapies. In addition, other promising strategies such as the use of nanoparticles, antibodies and, more recently, photodynamic inactivation for treating fungal infections have been studied extensively because of their promising therapeutic potential. There is a need to search for new products with effective antifungal abilities due to the adverse side effects of existing medications, increasing emergence of strains that are resistant to conventional antifungal agents, and the formation of biofilms in medical devices and tissues.

References


Candida spp. and new therapeutic options


