REVIEW ARTICLE



COVID-19 associated mucormycosis: evolving technologies for early and rapid diagnosis

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Abstract

The post-coronavirus disease (COVID-19) mucormycosis is a deadly addition to the pandemic spectrum. Although it's a rare, aggressive, and opportunistic disease, the associated morbidity and mortality are significant. The complex interplay of factors aggravating CAM is uncontrolled diabetes, irrational and excessive use of antibiotics, steroids, and an impaired immune system. Recently, India has been witnessing a rapid surge in the cases of coronavirus disease-associated mucor-mycosis (CAM), since the second wave of COVID-19. The devastating and lethal implications of CAM had now become a matter of global attention. A delayed diagnosis is often associated with a poor prognosis. Therefore, the rapid and early diagnosis of infection would be life-saving. Prevention and effective management of mucormycosis depend upon its early and accurate diagnosis followed by a multimodal therapeutic approach. The current review summarizes an array of detection methods and highlights certain evolving technologies for early and rapid diagnosis of CAM. Furthermore, several potential management strategies have also been discussed, which would aid in tackling the neglected yet fatal crisis of mucormycosis associated with COVID-19.

Keywords Mucormycosis · COVID-19 · Diagnostics · Biosensors · Probiotics

Introduction

COVID-19 topples down the immune system and the fungal infections reap its benefit. Since 2019, the pandemic of severe acute respiratory syndrome coronavirus-2 (SARS-CoV-2) has caused global havoc. Viral respiratory tract infections are usually accompanied by invasive fungal infections (mycoses) that are largely unrecognized but have a significant mortality rate of more than 50% (Raut and Huy 2021). Recently, critically ill patients with coronavirus disease-19 (COVID-19) were reported vulnerable to secondary infections by bacteria and fungi (Rawson et al. 2021). However, the emerging cases among COVID-19 patients have presented invasive fungal infections, mainly by species of

Mahesh Dharne ms.dharne@ncl.res.in Aspergillus, Candida, Rhizopus, Cryptococcus, and Pneumocystis, causing severe complications in the disease outcome (Hoenigl 2020). Being a new virus, the pathogenicity and associated immunological responses with SARS-CoV-2 are incompletely understood. However, the invasion of this virus triggers the release of danger-associated molecular patterns (DAMPs), that are fundamental in the pathogenesis of fungal diseases (Tolle and Standiford 2013). Besides, immune impairment of COVID-19 affected patients due to treatment with broad-spectrum antibiotics, invasive or noninvasive ventilation, immunosuppressive/corticosteroid, establish a favorable niche for the opportunistic infections by fungal pathogens (Segrelles-Calvo et al. 2020). These invasive fungal infections are notoriously challenging, progressive, and have fatal consequences upon delay in their diagnosis or treatment.

Mucormycosis is a rare but serious fungal infection of the skin, sinuses, lungs, and brain, caused by a group of ubiquitously present molds called mucormycetes. Although its occurrence is sporadic, the manifestation can be life-threatening under several medical conditions and the associated risk factors (Cornely et al. 2019). The major fungi associated with mucormycosis at the global scale are *Rhizopus*



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arrhizus, R. microsporous, R. homothallicus, Mucor spp. Lichtheimia spp., and Apophysomyces spp. (Prakash and Chakrabarti 2021). Mucormycosis is characterized by vessel thrombosis and subsequent tissue necrosis due to extensive angioinvasion by the fungal hyphae (Spellberg et al. 2005). Apart from aspergillosis and candidiasis being reported as the major fungal superinfections with SARS-CoV-2, a recent upsurge of cases with COVID-associated mucormycosis (CAM) has been reported worldwide at an alarming rate (Singh et al. 2021). Although COVID-19 is a global peril, CAM is more rampant in India. A recent review by John et al., has summarized reported cases of patients with CAM. It was observed that more than 71% of the cases were from India, and over 94% of patients had uncontrolled diabetes mellitus (DM) as an underlying condition (John et al. 2021). Typically, during the second wave of COVID-19 infection by the B.1.617.2 (Delta) variant, a sudden increase in the cases of CAM has devastated the prognosis, making it an epidemic in several states (DH Webdesk 2021; Raut and Huy 2021). India is the world capital of diabetes, with around 70 million people with DM (Pandey and Sharma 2018) and a COVID-19 caseload of over 31 million (Mallapaty 2021; The Lancet 2021), answers its prevalence in India. On the contrary, patients with hematological malignancies (HM), solid-organ transplantations are most susceptible to mucormycosis in developed countries (Jeong et al. 2019; Reid et al. 2020). Other predisposing factors associated with CAM are chronic kidney disease, steroid/antibiotic therapy, pulmonary tuberculosis, and chronic obstructive pulmonary disease (Prakash and Chakrabarti 2021; Patel et al. 2021). The precise mechanisms of CAM are yet to be elucidated. Meanwhile, rapid detection of the CAM followed by an appropriate line of treatment has become the need of the hour. To address this issue, the current review provides a comprehensive background on CAM, existing and emerging diagnostic/detection methods, and various management strategies to efficiently tackle mucormycosis.

Pathogenesis of mucormycosis and CAM

A: patho-mechanism of mucormycosis in healthy individuals

The pathogenesis of mucormycosis, typically by species of *Rhizopus* involves the entry of fungal spores in the susceptible host through inhalation, ingestion of food-soiled with spores, and through an abraded skin (Baldin and Ibrahim 2017). A spore cot protein cotH is present exclusively in all Mucorales and plays crucial role pathogenesis of mucormycosis. Following the entry, the attachment of fungal spore coat protein CotH3 to glucose-regulated protein 78 (GRP78) on the surface of endothelial cells occurs (Gebremariam



et al. 2019). Upon contact of CotH to the epithelial cells of humans, increased signaling of platelet-derived growth factor receptor B (PDGFRB) occurs which result in subsequent damage of the epithelial cells. The increased PDGFRB provide suitable growth factors for successful germination, development, and growth of Mucorales hyphae. Further, CotH3 plays an important role as invasins and mediate disruption and damage of immune cells, thereby promoting infection (Gebremariam et al. 2014). The mature hyphae produces Mucorales specific T cells, that in turn produce pro-inflammatory cytokines such as interleukins (IL-4, IL-10 and IL-17) and interferon gamma (IFN- γ) which then stimulate CD4⁺T cells and cause further damage the host cells. Further, the mature fungal hyphae suppress the release of immunomodulatory molecules such as regulated upon activation, normal T-cell expressed and secreted (RANTES), and IFN- γ from the natural killer (NK) cells (Schmidt et al. 2013, 2016). Once the fungal hyphae has invaded the immune system, it intrudes into deep tissues, causing tissue infarction, necrosis, and thrombosis (Brunke et al. 2016). Clinical conditions such as hyperglycemia, diabetes mellitus (DM), and diabetic ketoacidosis (DKA) enhance the expression of GRP78 and CotH3, increasing the susceptibility of the host to mucormycosis (Hassan and Voigt 2019). As the amount of free serum iron is abundant in these individuals, it creates a favorable niche for the growth of Mucorales. These fungi employ their high affinity iron permease to transport and use the available iron for their growth and development inside the human body (Artis et al. 1982; Roilides et al. 2012). Besides, reduced chemotactic factors in DM, DKA, and hyperglycemia conditions, damage the neutrophils, making it facile for the rapid spread of mucormycosis. Details of mucormycosis causing Mucorales, their virulence factors and clinical manifestations have been summarized in Table 1.

B: COVID-19 and mucormycosis: an intertwined association

The spread of SARS-CoV-2 occurs mainly through droplets, and is characterized by the binding of the virion particle to the angiotensin-converting enzyme 2 (ACE2) receptors of humans, via its spike protein. The infection with SARS-CoV-2 negatively influences the inflammatory response resulting into an uncontrolled immune response, called the cytokine storm. Further, the cellmediated immune response by the CD 4+ and CD 8+ T-cells, decreases in COVID-19 patients. Moreover, SARS-COV-2 mediated pneumonia results into atelectasis, which results into COVID-19 associated silent hypoxia. Under normal conditions, the hypoxia-inducible factor or HIF-1 remain inactive. However, during hypoxic conditions of COVID-19, the HIF-1α transcription factor

Table 1 A comprehensic	Lable 1 A comprehension of the Mucorales group causing mucormycosis in humans	ising mucormycosis in huma	IIIS			
Pathogens	Route of transmission	Site of infection	Virulence factors	Susceptible hosts and associated risk factors	Disease manifestations	References
Rhizopus spp.	Respiratory, oral, gas- trointestinal (GI), and percutaneous	Sinuses, brain, lungs, skin, and GI tract. Disseminated infection involves the central nervous system, heart, liver, kidney, and spleen	Thermotolerance, Ergot alkaloids (agroclavine, ergosine, and ergot- amine), Mycotoxin like rhizonin A, hydroxa- mate siderophores, glycosidic, lipolytic and proteolytic enzymes, antigenic extracellular polysaccharides, alka- line <i>Rhizopus</i> protease enzyme (Arp) ketone reductase system	Patients with predis- posed illness, diabetes, compromised immune system, steroid or broad-spectrum anti- biotics, iron overload, hyperglycemia,	Angioinvasion, infarc- tion, thrombosis, necrosis, and less often acute inflammation	(Morace and Borghi 2012; Binder et al. 2014; Walther et al. 2019)
Mucor spp.	Respiratory, oral, gastro- intestinal (GI), cutane- ous and subcutaneous	Skin, nails, ear, sinuses, brain, eyes, liver, kid- ney, and heart	Calcineurin (CaN), ADP- ribosylation factor (Arf), rodlet hydro- phobins, ferroxidases	Immunosuppressed individuals, patients with leukemia, aplastic anemia, bone/organ transplants, diabetes mellitus, asthma, burns, corticosteroid treat- ment, Hepatitis/HIV infection	Cutaneous infections with necrotic and hem- orrhagic lesions with typical black eschars or gangrenous cellulitis and erythema, nail infections with punctate erosions, invasive organ damage in immuno- compromised hosts. It is also responsible for mastitis in cattle	(Petrikkos et al. 2012; Binder et al. 2014; Petrikkos and Tsioutis 2018; Hassan and Voigt 2019)
Rhizomucor spp.	Respiratory, percutaneous	Skin, lungs, or dis- seminated disease. Very rarely heart is involved	Thermotolerance, acid- proteases, lipases, ethanol, acids, and mycotoxins	Patients with immuno- suppressive treatment like antibiotics and steroids, profound neutropenia, aplastic anemia, myelofibrosis, diabetes	Mostly associated with animal disease and rare in humans with pulmo- nary or disseminated disease	(Wickline et al. 1989; Severo et al. 1991; St-Germain et al. 1993; Weitzman et al. 1995)

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Pathogens	Route of transmission	Site of infection	Virulence factors	Susceptible hosts and associated risk factors	Disease manifestations	References
Lichtheimia corymbif- era (formerly Absidia corymbifera)	Respiratory, oral, gastro- intestinal, percutaneous	Sinuses, brain, GI tract, lungs, and skin	Ability to survive in extreme environmental conditions, dihydroli- poyl dehydrogenase, serine and aspartate proteases, and sidero- phores,	Immunocompromised individuals, patients with organ transplants and tissue grafts, diabe- tes, and cystic fibrosis Coinfection may occur with cytomegalovirus (CMV), HIV, hepatitis C virus (HCV), and pulmonary tubercu- losis. Risk factors include Hyperglycemia, Hyperferritinemia, low pH, and decreased phagocytic defense	Cutaneous infections with grey-black plaques, extensive necrosis. Rhino-cere- bral infections involve typical black necrotic nasal turbinate, swell- ing, inflammation, and rarely direct trauma	(Scalise et al. 1999; Ribes et al. 2000; Rickerts et al. 2006; Blomberg et al. 2007)
Apophysomyces elegans	Respiratory, Percutane- ous, postoperative surgical wounds	Skin, sinuses, bone, muscles, fat, kidney, bladder, orbital tissue, central nervous tissue, and less frequently dis- seminated infections	Thermotolerance, pro- teases, siderophores	Soil-contaminated wounds, immunocom- promised (severe burns and organ transplants) and diabetic individu- als,	Necrotizing cellulitis, tissue, and angioinva- sion, painful swell- ing, edema, extensive thrombosis, black hemorrhagic or white and friable lesions	(Cooter et al. 1990; Neblett Fanfair et al. 2012; Egge et al. 2018; Rashid et al. 2021)
Saksenaea vasiformis	Open, soil contaminated cutaneous wounds and/ or lesions,	Skin, sinuses	No specific virulence factors	Individuals with open wounds together with trauma, immunocom- promised hosts due to antibiotics, steroid treatment or with an in- dwelling catheter, leu- kemia with neutropenia, diabetes, thalassemia, and solene-fromy	Cutaneous lesions with painful red blisters and peeling skin with purulent eschar, less frequent evidence of cheesy necrosis or fri- able tissue Abscess formation in sinuses	(Saksena 1953; Gomes et al. 2011; Pilch et al. 2017; Lumyongsatien et al. 2020)

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Pathogens	Route of transmission	Site of infection	Virulence factors	Susceptible hosts and associated risk factors	Disease manifestations	References
Cuminghamella berthol- letiae	Upper respiratory tract, GI tract, percutaneous, cutaneous, and subcu- taneous	Eyes, cars, lungs, skin, sinuses, brain, joints, liver, and kidneys	Thermotolerance	Individuals with a compromised immune system, hemato- logic malignancies, neutropenia, diabetes, acidosis, nonmalignant hematologic condi- tions, hyperglycemia, iron overload, treatment with broad-spectrum antibiotics, predisposed infections of HIV, CMV, and HCV	Angioinvasion, hemopty- sis, multiple cavities in lower and upper lobes of lungs	(Zeilender et al. 1990; Pat- rick et al. 1993; Darrisaw et al. 2000; Rickerts et al. 2000; Liu et al. 2019)
Cokeromyces recurvatus	Unknown	Vagina and cervix are the major sites followed by the bladder and the G tract	Extracellular mycotoxins	Patients with compro- mised or dysfunctional immune system, antibiotic or steroid treatment, and diabetes	No visible sign of fungal invasion but acute and chronic inflammation in case of bladder infec- tion and mucus-laden, watery diarrhoea in the GI tract	(Rippon and Dolan 1979; Kemna et al. 1994; Tsai et al. 1997; Odronic et al. 2012)
Syncephalastrum rac- emosum	Contamination of open wounds with air-borne or soil-borne fungal spores (cutaneous), res- piratory tract, GI tract	Skin, nails, sinuses, eyes, GI tract, lungs	Mycotoxins, thermotoler- ance	Individuals with a previous history of lung abscess, diabetic ketosis, immunocom- promised, post-oper- ative trauma, chronic HBV infection, and nail injury for several months	Intracavitary fungal balls in lungs, arteritis, and osteomyelitis, onycho- mycosis, and multiple discharging lesions	(Schlebusch and Looke 2005; Gulati et al. 2021)
Mortierella wolfii	Unknown but presumed to be respiratory, through infected semen, GI tract, conjunctiva, skin wounds	Skin, lungs, eyes,	Thermotolerance, A water-soluble, heat- and trypsin-stable toxin (nephrotoxic)	Immunocompromised individuals, patients with organ transplants, and grafting	Chronic granuloma- tous disease, multiple hypodense lesions in the hepato-splenic parenchyma, keratitis	(Davey et al. 1973; Layios et al. 2014; Therese et al. 2020)

Table 1 (continued)

subunit has potent role over the expression of ACE-2 gene, hypoxia induced endothelial cell damage, and upregulation of innate and adaptive immune cells (Herrmann et al. 2020; Rahman et al. 2021). The overall impact of HIF-1 results into increased localized inflammation and tissue damage in COVID-19 patients. Besides, COVID-19 infection has also been shown to change iron metabolism. Several reports have suggested hyperferritinemic syndrome following COVID-19 infection. High levels of free iron result into tissue damage through production of free oxygen radicals (Vargas-Vargas and Cortés-Rojo 2020; Shoenfeld 2020; Perricone et al. 2020; Gómez-Pastora et al. 2020; Sonnweber et al. 2020). Furthermore, during the treatment of COVID-19, use of immunosuppressant's and corticosteroids are warranted to reduce the inflammation in the lungs and restrain the damage caused by cytokine storm. Steroids assist to curb the inflammation but it also suppresses the immune system, and could also trigger the uncontrolled release of sugar. Moreover, to overcome COVID-19 mediated hypoxia, use of oxygen masks and ventilators is essential. These potential factors make COVID-19 infected person more vulnerable to mucormycosis.

C. possible patho-mechanism of CAM

The main symptoms of COVID-19 such as hypoxia, hyperglycemia, hyperferritinemia, increased body temperature, and altered osmolarity, constitute an ideal environment for growth and development of mucormycosis causing fungi (Fig.1). Besides, if the oxygen masks or cylinders used in ventilators are contaminated with the spores of Mucorales, they serve as an entry pass of the spores into the COVID-19 infected patients. As steroid treatment results in immunosuppression, the spores easily enter the hosts system and germinate. The prevailing conditions in COVID-19 patients such as hyperglycemia and hyperferritinemia aid in rapid growth and multiplication of the fungal hyphae (Mahalaxmi et al. 2021).

Besides, several other important factors such as diabetes mellitus (DM), diabetic ketoacidosis (DKA), immune impairment, neutropenia, metabolic dysfunction, treatment condition with iron elevators, thrombocytopenia, treatment with voriconazole render the individuals more susceptible to the infection of mucormycosis (Ibrahim et al. 2012; Petrikkos and Tsioutis 2018). As this fungal infection is life threatening, and opportunistic, vigorous studies with experimental evidences are warranted to understand its pathogenesis especially in the individuals affected with COVID-19. A future

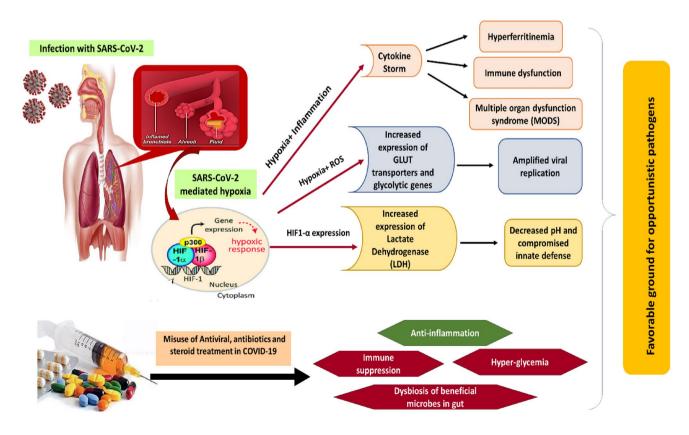


Fig. 1 SARS-CoV-2 infection and the associated outcomes laying the foundation for secondary infections to pitch in. Hypoxia-inducible factor- 1α (HIF- 1α); reactive oxygen species (ROS), glucose transporters (GLUT)



research on CAM would be instrumental in early diagnosis and management of this tangled duo.

Rapid detection of mucormycosis

The joint clinical guideline from the European Society for Clinical Microbiology and Infectious Diseases (ESCMID) and the European Confederation of Medical Mycology (ECMM) emphasizes the diagnosis and management of mucormycosis. The most widely used methods for detecting mucormycosis have been summarized here.

A. Diagnostic imaging

The most common clinical presentation of mucormycosis is rhino-cerebral, pulmonary, soft tissue, and disseminated disease. Although tissue necrosis is the hallmark of mucormycosis, its detection and early diagnosis lack sensitivity and specificity (Skiada et al. 2018). Computed tomography (CT) findings have been crucial in the early detection of pulmonary or spinal lesions, which includes halo signs, reverse halo signs, cavities with pleural effusion, nodules, and wedge-shaped infiltrates. These are considered signatures of angio-invasive fungi (Walsh et al. 2012). The presence of RHS along with more than ten pulmonary nodules, and pleural effusion are the most distinguishing characteristics of mucormycosis from aspergillosis (Skiada et al. 2018). Although CT scan provides a detailed picture of the organs/ tissue under study, combining it with positron emission tomography (PET) scan is recommended due to its sensitivity and higher accuracy to precisely point out anatomic abnormality (Liu et al. 2020). Magnetic resonance imaging is useful for assessing the extent of fungal disease in soft tissues (Yasmin et al. 2021). In addition to anatomic imaging modalities, functional and metabolic imaging using (PET/CT) coupled with [18F]-fluorodeoxyglucose (FDG) has been considered a valuable tool in the diagnosis and management of mucormycosis. Several reports have demonstrated an increased FDG uptake at the site of lesion upon diagnosis with FDG PET/CT during mucormycosis, and after antifungal therapy, the FDG uptake was reduced at the infection site (Dang et al. 2012; Liu et al. 2013; G Díaz et al. 2019; Song 2019; Gallo et al. 2019). Currently there are no reports supporting use of any particular technique in diagnosis of CAM. However, certain diagnostic imaging techniques based on the case reports have used high resolution computed tomography (HRCT), PET/CT, and 18F- FDG-PET/CT to evaluate structure of lungs in case of COVID-19 associated pulmonary mucormycosis (Dilek et al. 2021). (MRI) and 18F- FDG-PET/CT has been used for diagnosis of COVID-19 associated rhino-sinusal or rhino-orbitocerebral mucormycosis. While, for patients with COVID-19 associated rhino-facial mucormycosis, CT scan was used for diagnostic imaging (Garg et al. 2021; Mohammadi et al. 2021). Besides, recognition of patient's risk factors and careful assessment of clinical manifestations by use of imaging techniques would contribute significantly to the early diagnosis of mucormycosis. Further, a definitive diagnosis with a culture-based and molecular approach could be considered for initiating the appropriate antifungal therapy.

B. Microscopy and culture-based examination

The traditional detection methods in mycology include microscopy, histology, fungal culture, and serology. Direct mycological inspection of wide, non-septate, ribbon-like, hyaline hyphal elements in black chlorazol solution is the key feature related to Zygomycetes, which could be further supported by fungal culture (Machouart et al. 2006). For culture-based examination of mucormycosis-related clinical specimens, the use of fungal culture media, such as Sabouraud agar and potato dextrose agar is considered followed by its incubation 37 °C. All Mucorales grow on these media in 2-7 days but for some species, a microaerophilic environment improves culture yield (Lass-Flörl and Mayr 2009). However, the culture-based approach is not completely reliable because even when fungal hyphae are seen in histopathological analysis, fungal cultures are only positive in 50% of cases (Walsh et al. 2012). This is usually attributed to the fragile nature of fungal hyphae, which is prone to damage during tissue grinding or homogenization (Kontoviannis et al. 2007). Histopathological examination of the infected tissue would allow a keen differentiation between the hyphae of Mucorales and closely related fungi such as Aspergillus or Penicillium (Song et al. 2020). The typical hyphae of Mucorales genera are non-pigmented, wide (5-20 µm), thinwalled, ribbon-like, pauciseptate hyphae with right-angle branching. As mucormycosis is characterized by angioinvasion, perineural invasion, and prominent infarcts, an accurate examination would be critical for the choice of treatment. A rapid presumptive diagnosis of mucormycosis could be done through direct histopathological examination of the tissue through fluorescence microscopy using fluorescent brighteners such as Blankophor and Calcofluor White, which bind to chitin and cellulose in the fungal cell wall and fluoresce under ultraviolet (UV) light (Cornely et al. 2014).

C. Serological methods

Currently, there are no routine serological tests used for the diagnosis of mucormycosis. However, based on the specific antigens, techniques such as immunohistochemistry, Enzyme-linked immunosorbent assays, immunoblots, and immunodiffusion tests could be further evaluated for commercial use (Wysong and Waldorf 1987; Cornely et al.



2014). Additionally, a study by Potena et al., demonstrated Mucorales-specific T cells in a group of 28 hematologic patients with invasive mucormycosis using enzyme-linked immunospot (ELISpot) assay. Interestingly, none of the control group patients showed Mucorales-specific T cells, indicating its potential use as a reliable marker for the diagnosis of invasive mucormycosis (Potenza et al. 2011). Serum tests involving the use of 1,3-beta-D-glucan assay yield confounding results for the Mucorales group (Kontoyiannis 2016). This is because, as compared to other fungal pathogens, the dearth in understanding the cell wall structure and components of Mucorales. However, certain polysaccharides such as mucoran, mucoric acid, chitin, chitosan, and mannose have been reported as major components of the cell wall, while the presence of glucan has been demonstrated only in the spore cell wall of Mucor circinelloides, M. ramannianus, and Rhizopus oryzae (Lecointe et al. 2019).

D. Molecular methods

Molecular techniques are promising for rapid, sensitive, and accurate identification of pathogenic fungi causing mucormycosis (Voigt et al. 1999; Wu et al. 2003). This includes the use of polymerase chain reaction (PCR), restriction fragment length polymorphism analyses (RFLP), DNA sequencing of internal transcribed spacer ITS or 18S rRNA gene region for detection or identification of Mucorales (Lackner et al. 2014). A study by Machouart et al., demonstrated the use of a rapid and sensitive direct PCR-RFLP technique on a histopathological sample, for the identification of Mucorales in human pathology. This involved selective amplification of genomic DNA from molds followed by digestion step with restriction enzymes for species-level identification and further antifungal susceptibility testing (Machouart et al. 2006). In another study, the 18S rDNA and the D1–D2 domains of the 28S rDNA were identified for 42 isolates of Zygomycetes and 13 taxon-specific PCR primer pairs were designed for rapid and accurate identification of etiological agents of mucormycosis (Voigt et al. 1999). Scherer et al., demonstrated the use of, real-time, quantitative PCR (qPCR) assays, with 28S rDNA of Mucorales from bronchoalveolar lavage fluid, for rapid and accurate detection of invasive pulmonary mucormycosis (Scherer et al. 2018). Thus, given the increase in the number of cases with mucormycosis, the use of these molecular tools can be escalated for rapid and accurate identification of Mucorales from a pure culture.

E. Advances in fungal detection and emerging methods

Recent advancement in fungal diagnostics is the application of Matrix-assisted laser desorption ionization-time of flight mass spectrometry of (MALDI-TOF MS) that identifies



species-specific fungal peptides The Food and Drug Administration (FDA) approved commercially available MALDI-TOF MS platforms, are Bruker Biotyper (Germany) and Vitek MS (France). The advantage of this technique lies in the rapid identification of fungal isolates or direct clinical specimens in ~ 30 min (Yan et al. 2011; Spanu et al. 2012; Lau et al. 2019). Another FDA-approved commercial platform is 2 Magnetic Resonance (T2MR) (T2 Biosystems). This technique has been used for detecting pathogenic Can*dida* spp. Here, the clustering/complex between particles coated with target-specific agents is observed with the target pathogen with the altered microenvironment of the surrounding water molecules, which is detected and quantified by T2MR. Although individual sample costs are high, this technique can detect low levels of Candida cells within 5 h (Mylonakis et al. 2015; Zacharioudakis et al. 2018). Therefore, considering the current scenario, this technique could be a valuable tool for early (as it can detect low levels of fungal cells) and rapid detection of mucormycosis causing pathogens by escalating development of particles coated with Mucorales-specific agents. Additionally, a study by Koshy et al., suggested the use of breath-based biosensors for the diagnosis of IM. In this study, the volatile sesquiterpene metabolite profiles from the breath of murine models were analyzed using thermal desorption gas chromatography/tandem mass spectrometry (GC-MS/MS), and it was found that each Mucorales species produced a consistent profile of sesquiterpene secondary metabolite, which can be distinguished from aspergillosis (Koshy et al. 2017).

Biosensor technologies have been used for the direct detection of fungal pathogens. Biosensors are analytical devices that detect the concentration of bio-analyte and convert the physical or chemical signal into electrochemical or optical signals. Based on the technology, there are four types of biosensors: optical, electrochemical, thermal, and piezoelectric (Samson et al. 2020). Cai et al., has demonstrated optical detection of cell surface mannan of Candida albicans binding to hydrogel Con-A using a photonic crystal sensor (Cai et al. 2015). In a study by Bhatnagar et al., a sensitive electrochemical nanobiosensor fabricated with 1,6-Hexanedithiol and chitosan-stabilized gold nanoparticles was designed for rapid detection of Aspergillus fumigatus (Bhatnagar et al. 2018). Another study by Asghar et al., developed an immuno-based microfluidic chip that can rapidly detect and capture C. albicans from phosphate-buffered saline (PBS) and human whole blood (Asghar et al. 2019). Since biosensors have not been much explored for detecting fungal pathogens, it could be an active area for current interdisciplinary research. Likewise, given the current epidemic situation, the development of biosensors would be of immense importance for the rapid and early detection of mucormycosis, and would also aid in monitoring treatment response based on specific marker analytes (Fig. 2). A wide range of fungal biomarkers such as mannan, galactomannan,

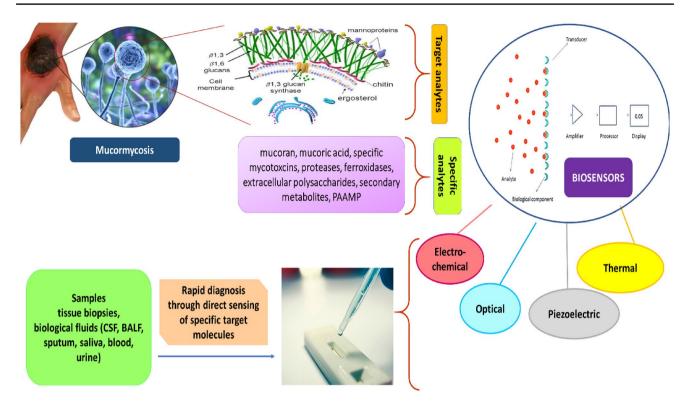


Fig. 2 Schematic illustration of target and specific analytes for development of biosensors for mucormycosis causing fungal pathogens

mannan, beta-glucan and cryptococcal antigen, have been explored for the development of diagnostic tests. In case of mucormycosis, there are no specific biomarkers for rapid identification. However, to the best of our knowledge only one study by Orne et al., where lateral flow immunoassay (LIFA) technique was developed by using monoclonal antibody (mAb 2DA6) against cell wall fucomannan of Mucorales. The monoclonal antibody mAb 2DA6 is extremely sensitive for fucomannan due to its apparent low-level of side chain substitution. This was then used as a biomarker for rapid diagnosis of invasive murine mucormycosis (Orne et al. 2018). However, no further studies were carried out with this biomarker to develop advanced techniques for rapid detection of mucormycosis. Therefore, considering its sensitivity, mAb 2DA6 based biosensors could be developed and employed for early diagnosis and rapid detection of CAM. Based on all the above-described detection methods, the use of a multidisciplinary approach for rapid detection and appropriate treatment of mucormycosis in patients could be instrumental in reducing the mortality rate of these infections.

Management of mucormycosis

Effective management of mucormycosis is a multimodal approach involving early, rapid detection and prompt treatment for effective prognosis. An optimal management relies upon recognizing the various patterns of the disease along with convenient diagnostic and therapeutic choices.

A. Reversal of the underlying condition

It is of pivotal importance to reverse or prevent the underlying disease/medical condition/immune dysfunction in the patients while treating them for mucormycosis. The use of immunosuppressant, in particular corticosteroids should be minimized or terminated if possible (Spellberg and Maertens 1991). Moreover, to reduce the severity of mucormycosis infections, administration of iron and blood transfusions should be avoided. Similarly, use of DFO should be minimized as it is known to aggravate mucormycosis. Besides, for patients with ketoacidosis and DM, a rapid restoration of euglycemia, and normal acid base status is crucial (Spellberg et al. 2009).

B. Surgical management

As per the global guidelines by The European Confederation of Medical Mycology (ECMM) jointly with the International Society for Human and Animal Mycology (ISHAM), and ECMM together with the Mycoses Study Group Education & Research Consortium (MSG ERC), an early and meticulous surgical resection and debridement of the tissue affected with mucormycosis is of pivotal



importance in localizing the spread of mucormycosis (Cornely et al. 2019). As the surgical treatment vastly depends on the site and extent of the disease, it is a preferred choice of treatment for rhino-orbito-cerebral and soft tissue mucormycosis. Additionally, for patients with pulmonary or disseminated mucormycosis, surgery could be considered on case-by-case basis (Skiada et al. 2013). A recent report by Claustre et al., suggested curative surgery to be the key survival factor for mucormycosis in patients with hematological malignancies (Claustre et al. 2020). In another study involving evidence-based mapping of CAM patients, a higher percentage of survival (64.96%) was observed for those who underwent antifungal therapy in adjunct to surgical debridement (Hussain et al. 2021). Thus, prompt surgical intervention is crucial in ameliorating the fatal outcomes of CAM. However, the subsequent prognosis would essentially depend on the infection site, spectrum, and etiology of the fungus, and the associated comorbidities. In cases of disseminated mucormycosis or absence of removal of infected focus, surgical management becomes challenging, and the antifungal therapy singly is seldom curative (Chibucos et al. 2016). Hence, surgical limitations drive the necessity to seek additional alternative strategies for prevention and management of invasive mucormycosis.

C. Antifungal therapy

Treatment of mucormycosis encompasses the use of antifungal agents along with surgical debridement. As per the recommendations by ECMM and MSG- ERC, the primary prophylaxis involves treatment of mucormycosis patients having comorbidities such as neutropenia or bone marrow transplant with moderate strength posaconazole tablets or its marginal strength oral suspension. The secondary prophylaxis includes treatment of mucormycosis affected immunosuppressed patients with surgical resection together with continuation of the last drug effective in that patient (Cornely et al. 2014).

C1: Primary antifungal therapy

Primary therapy includes use of polyenes. The only licensed antifungal for mucormycosis is Amphotericin B deoxycholate. However, to overcome its substantial nephrotoxicity and poor CNS penetration, lipid formulation of amphotericin B (5 mg/kg-10mg/kg), is administered as a safer and efficacious alternative for the treatment of mucormycosis (Walsh et al. 2008; Lewis et al. 2010). Besides, moderate strength isavuconazole, and posaconazole are also recommended as first-line antifungal monotherapy (Kyvernitakis et al. 2016).



C2: Primary combination therapy

There are no convincing studies to support the use of combination of antifungals in therapeutic regime of mucormycosis, and therefore, it is not included as major treatment guidelines. However, use of combination of antifungals in step-down therapy is warranted (Honavar 2021). In this combination therapy, use of broad-spectrum azoles such as posaconazole and isavuconazole in the form of oral or parenteral formulation could be used as a step-down therapy or as a salvage therapy for patients who are intolerable or do not respond to amphotericin B (Kauffman 2019). Also, itraconazole and terbinafine could be used depending on different strains of Mucorales (Skiada et al. 2018). A recent retrospective review by Reed et.al, suggested combination of polyene-caspofungin therapy as a promising alternative for ROCM patients treated with polyene monotherapy (Reed et al. 2008). Since the duration of treatment for mucormycosis remain unknown, it should be continued until the indicative findings through imaging have been resolved along with improvement of the hosts' immune system. Additionally, administration of isavuconazole or posaconazole could be considered as maintenance therapy (Kontoyiannis and Lewis 2011).

D: Adjunct therapy

D1: Iron chelators

Iron is fundamental for growth and virulence of microbial pathogens. Patients with elevated an increased serum iron levels are susceptible to mucormycosis. Iron metabolism plays an important role in infections with Mucorales, and therefore, use of iron chelators as an adjunctive therapy is important (Roilides et al. 2003; Bullen et al. 2006). Iron chelators have higher affinity constants for iron and therefore, prevent the accumulation of excess iron and challenge the fungal growth with the iron-depleting environment (Symeonidis 2009). Food and drug administration (FDA) approved iron chelators include oral tablets such as Deferiprone (DFP), Deferasirox (DFX), and injectable Desferrioxamine (DFO) (Scott et al. 2021). DFO received FDA approval for treating iron overload in transfusion-dependent anemia in the year 2005. This led to its unapproved use as an adjunctive therapy in advanced cases of mucormycosis, particularly in patients with underlying diabetic ketoacidosis (DKA). Further, as compared to DFO, DFP and DFX, do not increase susceptibility to mucormycosis. DFP and DFX have been reported to have antifungal activity in-vitro and invivo against Mucorales, and therefore, are used as effective iron chelators in clinical practice (Mahalmani et al. 2021). In CAM, iron is fundamental for the growth and life cycle of the mold. Hence, the use of iron chelators to regulate the iron concentration in the body would be imperative. Till date there are limited scientific evidences with respect to use of iron chelators in CAM. Therefore, a combinatorial approach of non-xenosiderophore iron chelators and antifungal agents could be employed in the treatment regime for their possible synergistic effect. Additionally, comprehensive studies evaluating the role of combined antifungals and iron chelators would be insightful in efficient management of mucormycosis.

D2: Cytokines

Cytokines are an integral part of the viable innate antifungal host defenses. Particularly, interferon- γ (IFN- γ) and granulocyte-macrophage colony-stimulating factor (GM-CSF) are the critical in immunomodulation of invasive fungal infections (Roilides et al. 2003). A study by Gil-Lamaignere et al., showed enhanced antifungal activity of human polymorphonuclear leukocytes (PMNLs) against R. oryzae and R. microsporous, by using pro-inflammatory cytokines IFN-y and GM-CSF, thereby suggesting, use of these cytokines in management of invasive mucormycosis (Gil-Lamaignere et al. 2005). Besides, several case reports have also suggested use of recombinant GM-CSF and granulocyte colony-stimulating (G-CSF) as an adjuvant therapy in management of invasive mucormycosis (Mastroianni 2004; Abzug and Walsh 2004; Spellberg et al. 2009). Additionally, G-CSF in combination with antifungal therapy has been employed as secondary prophylaxis in treatment of mucormycosis in patients with hematological malignancies (Grigull et al. 2006). Another recent study by dos Santos et al., suggested critical role of IFN-γ in enhancing fungal clearance of pulmonary mucormycosis in mice (dos Santos et al. 2021). This is because, IFN- γ induces a T-helper cell type 1 (Th1) cells and the subsequent immunological response enhances the antifungal activities of PMLs' and favors resistance to invasive fungal infections Although, the reported evidences with cytokines therapy for mucormycosis are limited, it would be an important area of future research to determine their role in primary treatment. Also, rigorous clinical trials are essential to warrant the use of GM-CSF, G-CSF, and IFN- γ auxiliary to the conventional antifungal treatment.

D3: Hyperbaric oxygen

The use of hyperbaric oxygen therapy (HBOT) as an adjunctive treatment for mucormycosis was first described in 1970. In this, 100% of oxygen is administered to the patients through masks or endotracheal tubes. The pressure inside the chambers is increased to 250–280 kPa, and the partial pressure of oxygen (PaO₂) is ~1200–2000 mmHg. *In-vitro* studies have shown antifungal activity of HBOT at pressures

<10 atm (Gudewicz et al. 1987; Galiazzo et al. 1987). There are several reports highlighting the direct antifungal effect of hyperbaric oxygen under hyper-oxic conditions due to increased production of oxygen based free radicals (Couch et al. 1988; Ferguson et al. 1988; Arnáiz-García et al. 2009; Ezra et al. 2011; Valente Aguiar et al. 2021). Additionally, HBOT have several indirect antimicrobial effects which includes restoration of phagocytosis, reversal of lactic acidosis, enhanced oxidative burst of PMLs and augmented antifungal action of amphotericin B (Siddiqui et al. 1997; John et al. 2005; Mathieu and Wattel 2006; Bitterman 2007; Tragiannidis and Groll 2009). Data supporting the clinical efficacy of HBOT in invasive mucormycosis is still limited. However, a comprehensive review by John et al., described use of HBOT as an adjunct therapy to treat zygomycosis. Of the total 28 patients, 81% had underlying conditions such as DM (61%), trauma (18%), HM (11%), alcoholic liver disease (7%), treatment with corticosteroids (3%). The remaining 11% had no underlying conditions. The overall survival rate after treatment was 86% (John et al. 2005). Although HBOT has been used to treat zygomycosis, the clinical evidence for its use in mucormycosis remain anecdotal and inconclusive. Therefore, further experimental studies are essential to evaluate the clinical efficacy of HBOT in CAM. This would be critical to determine the outcome and impact of this treatment in management of CAM.

Probiotics: an experimental potential future management alternative

Probiotics are live microorganisms that, when expended in appropriate amounts, confer a health benefit to the host (Gohil et al. 2021). Besides having anti-inflammatory activity, probiotics are also known to have anti-fungal and antimycotoxigenic activities (Dalié et al. 2010; Aiko et al. 2016; Russo et al. 2017; Arasu and Al-Dhabi 2017; Azizkhani et al. 2021). A study by Lipinska et al., had shown enhanced antifungal activity of 60 Lactobacillus sp. strains against selected food contaminating fungi such as Alternaria alternata, A. brassicicola, Aspergillus niger, Fusarium latenicum, Geotrichum candidum, Candida vini, and Mucor hiemalis, in presence of xylitol and galactosyl-xylitol (Lipińska et al. 2016). Another study by Karami et al., indicated a significant antifungal effect of two species of Lactobacillus viz L. alimentarius and L. delbrueckii on Penicillium notatum and Aspergillus fulvous (Karami et al. 2017). Lacto*bacillus* spp. has also been demonstrated to have antifungal activity against Candida spp. isolated from the patients with AIDS with HIV (Salari and Ghasemi Nejad Almani 2020), and against planktonic cells, biofilms, and persister cells of C. auris through postbiotic elements (Rossoni et al. 2018).



Details of probiotics and their immunomodulatory effects in fungal infections have been comprehended in Table. 2.

Recent meta-analysis by Hu et al., showed efficacy of probiotic bacteria in prevention and treatment of oral candidiasis in both animal experiments and in clinical trials. The possible mechanism behind prophylactic activity could be competition of probiotic species with the pathogens for nutrients and binding receptors, and release of certain metabolites which inhibits or antagonizes the growth of Candida spp (Hu et al. 2019). Several other studies have also shown in-vivo efficacy and safety of probiotics in treatment of candidiasis and other fungal infections (Brzozowski et al. 2005; Vicariotto et al. 2012; Demirel et al. 2013; Matsubara et al. 2016; Silva et al. 2016). Although the role of probiotics as antifungal agents has been well known, to date there are no experimental evidences supporting or evaluating the role of probiotics in treatment or prophylaxis of CAM. However, there are several reports and experimental studies highlighting the importance of use of probiotics as an adjunctive or prophylactic treatment for COVID-19 (He et al. 2020; Morais et al. 2020; Olaimat et al. 2020; Gohil et al. 2021; Chhibber-Goel et al. 2021). To date, four clinical studies have used probiotics in management of COVID-19 and around six studies are being assessed for the clinical trials (Baindara et al. 2021; Tang et al. 2021; Peng et al. 2021) Therefore, on the basis of the existing fundamental studies, future research could be directed toward assessing the efficacy of probiotics singly or as an adjunct with the existing management strategies in treatment of CAM. Thus, probiotics could be used to tackle the triple threat of COVID-19 i.e.,

Table 2 Antifungal activity of probiotics in fungal infections

as a replenisher of gut dysbiosis, as an anti-inflammatory agent, and as an antimicrobial (antifungal) agent.

Conclusion and prospects

Crisis brings opportunity, and the time to act on mucormycosis is now. The rapid increase in mucormycosis infection and the associated fatality is alarming. The clinical symptoms are elusive and the use of conventional techniques for detection/diagnosis is time-consuming, insensitive, and unspecific (Skiada et al. 2018). Therefore, there is an urgent need for developing methods for rapid detection of mucormycosis and appropriate antifungal management. Currently, combining the culture-based method with molecular methods or advanced techniques such as MALDI-TOF, GC MS/MS, T2MR would significantly improve the diagnosis of mucormycosis. Further, the emerging biosensor platforms could be developed by combining specific molecular biomarkers or species-specific recognition elements to address the rapid detection challenge of mucormycosis. Also, advancement in microchipbased or implantable biosensors for continuous monitoring of fungal analytes during treatment would be crucial to highlight the critical points during therapeutic intervention. For effective management of CAM, antifungal agents with iron chelators could be used. Moreover, the use of probiotics could be beneficial in tackling mucormycosis/ CAM and achieving immune homeostasis.

Sr. No	Probiotic microbes	Immunomodulatory activ- ity of the probiotic	Activity against	Useful in combat- ing disease/condi- tion	References
1	L. fermentum 20.4, L. paracasei 28.4, L rhamnosus, Pediococcus acidilactici. HW01	Biosurfactant produced by the organism reduces adhesion and biofilm formation of the fungus	Candida albicans	Candidiasis	(Rossoni et al. 2018; Kim and Kang 2019)
2	Saccharomyces cerevisiae KTP	Adhesion, filamentation and biofilm formation	Candida krusei	Candidiasis	(Kurrey et al. 2019)
3	L. casei, L. acidophilus, L. paracasei	Decreased pH due to acid production along with anti-microbial agents	Aspergillus niger, A. flavus, A. parasiticus, P. chrys- ogenum	Aspergillosis	(Abbaszadeh et al. 2015)
4	L. plantarum KCC-28	Production of hydrolytic enzymes	Fusarium oxysporum	Keratitis	(Ilavenil et al. 2015; Batista et al. 2020)
5	Lactobacillus harbinen- sis K. V9.3.1Np	No observation	Yarrowia lipolytica, Kluyveromyces marxi- anus, Penicillium brevi- compactum	Food poisoning	(Delavenne et al. 2013)
6	Leuconostoc mesenter- oides	Bioactive compound production	Meyerozyma guilliermon- dii, P. roqueforti, Asper- gillus oryzae	Food poisoning	(Yépez et al. 2017)



Declarations

Conflict of interest The authors declare no conflict of interest.

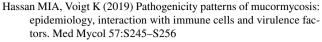
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