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Iron and glucose-regulated protein 78: substantial components in the coinfection of mucormycosis and COVID-19

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Trabalho de Conclusão de Curso apresentado como requisito parcial à obtenção do grau farmacêutico pela Universidade Federal do Rio Grande do Sul.

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SUMÁRIO

Title: Iron and glucose-regulated protein 78: substantial components in the coinfection of mucormycosis and COVID-19 **Short title:** Iron and GRP78 role in mucormycosis and COVID-19

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ABSTRACT

The viral outbreak of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) emerged in China by the end of 2019 and was declared a global pandemic in early 2020. Along with the growing number of fatalities and a lack of specific treatment, the increasing incidence of mucormycosis worried world health agencies, as it ran the risk of even more threatening outcomes for patients with coronavirus disease 2019 (COVID-19). In this context, this review aims to assemble case reports of mucormycosis and COVID-19 coinfection and discuss the virulence and the host factors involved in the progress of these infections – key aspects that might unveil potential biological targets and pharmacological approaches to treat these infectious diseases. Recently, elevated serum iron levels during SARS-CoV-2 infection have been reported in the literature. Besides being a clinical characteristic of diabetic patients, iron overload is described as a virulence factor for *Rhizopus oryzae*. Furthermore, the increased expression of human heat-shock protein GRP78 during iron overload and coronavirus infection display a crucial role as a mediator in Mucorales invasion and, likewise, in SARS-CoV-2. These remarkable mechanisms might explain the high incidence of mucormycosis in diabetic and COVID-19 patients and, therefore, suggest regulation of GRP78 levels, management of glucocorticoid treatment and glycemic control as potential therapeutic targets of this severe coinfection.

Keywords: SARS-CoV-2. COVID-19. Mucormycosis. Iron. GRP78. Glucocorticoids. Virulence factors. Diabetes.

1 INTRODUCTION

The outbreak of the coronavirus disease 2019 (COVID-19) in China at the end of 2019 became a global pandemic threat in early 2020. Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) emerged as a remarkable issue of concern due to the rapid increase in mortality rates and the lack of any specific pharmacological therapy. ¹ Mucormycosis is a life-threatening fungal infection, which was shown to be a secondary complication of COVID-19, particularly in immunocompromised hosts.^{2, 3} The global mortality from mucormycosis is 46% and underlying medical conditions could lead to a 93% fatality rate.^{4,5}

Uncontrolled diabetes mellitus, corticosteroid therapy, organ transplantation, hematologic diseases and neutropenia are all relevant risk factors that support the most serious outcomes in mucormycosis infection. ⁶ However, these medical conditions can also trigger lethal coinfection with COVID-19. Major clinical complications include acute respiratory distress syndrome and rhinocerebral impairment, which establish a potential risk to the hosts of these pathogens.^{1,2}

Virulence factors and the host response have an important role in the pathogenesis of infections.⁷ High affinity iron permease (FTR1) upregulation by Mucorales in hyperglycemic mice and COVID-19 patients with an elevated expression of glucose-regulated protein 78 (GRP78), which binds with the spore coat (CotH) protein of Mucorales, suggests that these mechanisms could be crucial components of the coinfection.^{8,9,10}

The lethal alliance of mucormycosis and COVID-19 has a diagnosis that relies substantially on histopathological findings and molecular techniques. In addition, the few pharmacological alternatives available, often lead to nephrotoxicity and surgical debridement.^{11,1} Understanding the association between the virulence factors and the host characteristics can reveal a key aspect of the infection's progression and could provide a more efficient therapeutic strategy.

2 RESEARCH STRATEGY

The study was carried out in the form of scientific research in the ScienceDirect and PubMed databases, using original studies, case reports and reviews published between 2019 and July of 2021. The keywords used in the search were combined into

three groups: ["SARS-CoV-2" or "COVID-19], ["mucormycosis" or "mucor" or "rhizomucor" or "rhizopus" or "mucorales'' or "zygomycosis"] and ["ferritin" or "iron" or "GRP78"]. No language restriction was applied. Antifungal susceptibility studies, chapters of books and clinical trials were excluded (Fig. 1).

3 DISCUSSION

3.1 Virulence factors

3.1.1 Iron uptake and FTR1 expression

Zygomycosis is a fungal infection caused by *Zygomycetes*, a class of fungi that encompasses the orders Mucorales and Entomophthorales. Diverging from entomophthoromycosis, rare infections that usually affect immunocompetent individuals, fungi of the order Mucorales cause worldwide life-threatening infections in immunocompromised hosts, which are known as mucormycosis.² In 1885, the first case of mucormycosis in humans was reported and identified as "Mycosis Mucorina". 12 The detailed information described in Paltauf's et al. ¹² paper strongly suggested that *Absidia corymbifera*, a mold that belongs to the order Mucorales, was the etiologic agent of the disseminated infection referred to in a cancer patient.

The majority of isolated species in patients with mucormycosis belong to the *Mucoraceae* family, especially *Rhizopus oryzae*. ² This ubiquitous mold is the most frequent *Rhizopus* species in the environment and has been identified all around the world, 13,14 which might explain *why R. oryzae* is the most common cause of mucormycosis. The expression of virulence factors by this pathogen may justify its high incidence in the environment and of infections. Ibrahim et al., ¹⁵ revealed that *R. oryzae* hyphae and spores have the ability to adhere to endothelial cells in vitro and damage them.

Furthermore, *R. oryzae* can obtain iron from the host through high affinity iron permease (FTR1), a component of a cytoplasmic membrane reductive-oxidase complex. Iron is an essential micronutrient for human life that has an important role in cell growth and virulence of several pathogens. However, increased iron serum levels can result in toxicity, mucosal damage and impaired immune response due to oxidative cell injury, enhancing the risk of fungal infections. 16,17 Iron is stripped by the FTR1

system from heme, transferrin, ferritin and chelating agents, such as deferoxamine, which is used to treat diabetic patients with iron overload (Fig. 2). The ferric iron is reduced into its more soluble form, carried through the fungus cell membrane into the cytoplasm and oxidated into ferric iron, that enhances the virulence of the pathogen, over a mechanism that remains unclear.¹⁴ Ibrahim et al., ¹⁰ study revealed that FTR1 expression by *Rhizopus oryzae* is upregulated in a murine model of hyperferritinemia. However, Ibrahim further showed that FTR1 inhibition decreased the virulence and passive anti-FTR1 immunotherapy in diabetic ketoacidosis (DKA) mice protected against *R. oryzae* infection, which suggests that iron is a crucial component in the progression of mucormycosis. 10

First described in *Saccharomyces cerevisiae*, FTR1 is a highly conserved gene in fungi that, later, was found to be expressed in *Candida albicans* as well as in green algae. 18,19,20 However, in these organisms, the FTR1 upregulation occurs in low iron concentrations, which constrasts with *R. oryzae* upregulated FTR1 during iron overload – a condition that can occur in diabetic and COVID-19 patients.¹⁹⁻²² Since the use of deferoxamine increases the susceptibility to mucormycosis in patients with DKA, recent data has suggested using lactoferrin to manage iron serum levels. This potential iron-chelator shows that its fungistatic and immune regulator activity can be a helpful tool in the adjuvant therapy of mucormycosis.²³

In a similar way to Mucorales infection, COVID-19 modifies the iron metabolism of the host. Increased ferritin levels and iron overload associated with coronavirus disease have been described in the literature. ²⁴ During SARS-CoV-2 infection, the inflammatory process releases cytokines that increase intracellular iron, leaking the metal into the bloodstream and establishing a higher risk of mucormycosis.²⁵ In this context, ferritin - an iron storage protein and marker for acute inflammation, might be an abundant source of iron for Mucorales – additionally to transferrin, chelating agents and red blood cells previously described by Ibrahim et al., 14 , 26 . Since ferritin levels increase in diabetic patients and during SARS-CoV-2 infection,^{27, 41} this globular protein could offer Mucorales easy access to iron, increasing its virulence mechanisms, facilitating the fungus to establish itself in the host and providing more fatal outcomes. In this review, only five case reports presented patients with hyperferritinemia^{28,29,30,31,32} (Table 1), although it is known that most case reports did not have described serum iron and ferritin levels and it is likely that more patients may have had an iron overload that potentiated the progression of mucormycosis.

3.1.2 GRP78 expression and COTH3 interaction

GRP78 is a class of stress-responsive heat-shock proteins involved with the unfolded protein response process in the endoplasmic reticulum lumen. Under normal conditions, GRP78 directs misfolded proteins to refolding and degradation mechanisms. However, under stress conditions, GRP78 can break out of the endoplasmic reticulum and settle in the cell membrane as an exposed protein, characteristic of different types of cancer and an enabler of various viral infections, such as Dengue Virus, Zika Virus, Middle-East Respiratory Syndrome Coronavirus and, most recently, SARS-CoV-2.⁴⁷⁻⁴⁹

 Curiously, GRP78 was revealed as a mediator in host cell invasion by Mucorales as well.⁵⁰ In 2014, Gebremariam et al., ⁵⁰ identified a spore coating protein family in *Rhizopus oryzae,* CotH3, which mediates the attachment to GRP78 and promotes fungal endocytosis, invading host endothelial cells. They also demonstrated that anti-CotH antibodies (Abs) therapy protected a murine model of DKA from mucormycosis and showed that a mutant of *R. oryzae* with downregulated CotH expression presented decreased virulence. Furthermore, GRP78 expression is upregulated in hyperglycemic mice with iron overload and anti-GRP78 Abs therapy protected the murine model from mucormycosis.⁸ In this context, the remarkable ability of the fungus to invade and damage host cells challenges the immunity of patients with DKA which, among its many complications, expresses a ligand – GRP78 that facilitates the fungal infection. Considering that until this moment no records have been found in the literature that GRP78 binds to any other fungi, it is possible that this interaction may be unique to these eukaryotes, which requires further research to prove the supposed exclusivity.

Moreover, recent data has shown that there is increased GRP78 serum levels in COVID-19 patients, which supports Ibrahim et al.,⁴⁸ findings that indicated GRP78 was a receptor for SARS-CoV-2.⁵¹ However, the impact of increased expression of GRP78 on the prognosis of COVID-19 patients is not fully determined. Although GRP78 levels have not been measured in any case report, is expected that its expression would be increased, since all patients tested positive for COVID-19 and most were diabetic (Table 1), which provides potentially aggravating components for the development of mucormycosis and its fatal outcomes.

3.2 Host factors

3.2.1 Systemic corticosteroid therapy

Corticosteroids are among the most commonly prescribed medications available and these drugs are used quite broadly due to their potent anti-inflammatory and immunosuppressive properties. ⁵² The use of glucocorticoids (GC) during the COVID-19 pandemic as an adjuvant therapy has significantly increased. The World Health Organization has initially not recommended its use outside of clinical trials.⁵³ However, in September 2020, new guidance based on moderate certainty evidence was proposed with a strong recommendation about using steroid therapy to treat patients with severe COVID-19, in order to control the cytokine storm caused by the infection. 54,55 The effectiveness of corticosteroid therapy on SARS-CoV-2 infections remains controversial. COVID-19 patients demonstrated delayed viral clearance times and there has been shown to be a not significantly statistical improvement in survival rates when steroid treatment was been used. 56

According to the selected case reports in this review arranged in Table 1, patients who were treated for COVID-19 with corticosteroids developed mucormycosis. These findings corroborate Rickerts et al.,⁶ study, which suggested systemic corticosteroid therapy was a risk factor for the fungal infection and supports Lionakis et al., ⁵⁷ data, which proposed that high dose steroid treatment and its long-term use increased the susceptibility to opportunistic fungal infections – as invasive aspergillosis, candidaemia, cryptococcosis and zygomycosis, due to GC immunosuppressive effects. In this context, the risk of infection is directly affected by underlying medical conditions that determine the duration of corticosteroid therapy and the dosage. Moreover, viral, bacterial and parasitic infections can be predisposed by GC as well. 58

Chronic high-dose steroid therapy leads to suppression of the host's immunity due to impairment of monocyte migration, facilitating angioinvasion by Mucorales.⁵⁹ The transcriptional modulation mechanism of glucocorticoids inhibits the c-Jun Nterminal kinase (JNK) pathway and kinase-β (IKKβ) resulting in the repression of nuclear factor-kB (NFκB) and activator protein 1 (AP-1), important transcription factors involved in monocyte-macrophage activation. The inhibition of NFκB and AP1 reduces the macrophage secretion of pro-inflammatory cytokines, like interleukin-1 (IL-1), IL-6,

IL-8 and tumor necrosis factor-α (TNF-α), which are essential to the innate immunity's line of defense against fungi. In addition, steroid therapy reduces the secretion of IL-12, IL-2 and interferon γ, which affects the adaptive immune response (Fig. 3). Furthermore, glucocorticoids cause an imbalance in T-helper (Th) cells, which produces an enhanced Th2 cytokine response that leads to an impaired phagocytic function. Associated with NFκB inhibition by steroids, anti-apoptotic effects are observed in dysfunctional neutrophils and suppression of cell adhesion molecules required for lymphocytic signaling, as CD18 and intercellular adhesion molecule 1 (ICAM-1), were observed as well. 60,57,61

Besides, steroid therapy can develop resistance to neuromuscular blocking drugs – generally used in patients that require mechanical breathing support, hyperglycemia, neutropenia and can also predispose multiple opportunistic infections. ⁶² These features could explain the high incidence of mucormycosis in diabetic patients with COVID-19 that were treated with corticosteroids and reveal that strict monitoring and management of corticosteroid therapy might be a valuable tool to control the progression of SARS-CoV-2 and Mucorales infections.

3.2.2 Hyperglycemia and diabetes mellitus

Similar to the immunosuppressive effects of corticosteroid therapy, hyperglycemia and ketoacidosis can also suppress the immune system by impairing chemotaxis and phagocytic function. Mooradian et al., ⁶³ study revealed that peripheral mononuclear blood cells isolated from diabetic patients had a lower levels of IL-1 secretion compared to healthy controls. Additional data also showed that diabetic patients presented suppression of IL-2, IL-6 and IL-10 cytokine secretion, which affects antibody production and increases the risk of infection. Diabetic patients exhibited greater tumor necrosis factor alpha (TNF-α) and IL-8 levels as well. However, it is has been reported in the literature that high concentrations of TNF-α cause insulin resistance by inhibiting the peroxisome proliferator-activated receptor-gamma, which leads to a worsening of hyperglycemia condition.⁶⁴ Clinical evidence revealed that patients with diabetes and dysfunctional phagocytes are more susceptible to developing fungal infection.⁶⁵ Furthermore, hyperglycemia induces transferrin and ferritin glycosylation, which decreases the iron binding to these proteins and increases the serum and intracellular iron levels, generating reactive oxygen species that lead to

tissue damage (Fig. 2).⁶⁶ The described mechanisms above support the increased risk of mucormycosis in diabetic patients, since cytokine production is suppressed and the amount of serum iron is increased.

Notably, some patients described in this review, who reported a history of diabetes, had episodes of DKA when hospitalized (Table 1). The acidic environment caused by hyperglycemia decreased the transferrin's iron-binding ability, which released the metal into the bloodstream.⁶⁷ High iron serum levels lead to a higher expression of GRP78, that binds to spore coat protein of Mucorales and mediates the fungal invasion.⁵⁰ However, Gebremariam et al.,⁶⁸ study found that sodium bicarbonate reversed the acidosis in ketoacidotic mice, which protected them from mucormycosis and highlighted the importance of pH correction as a therapeutic measure.

Similarly, the spike protein of SARS-CoV-2 also binds to GRP78, which expression is increased during the infection, allowing virus attachment and cell invasion. Furthermore, COVID-19 induces hyperglycemia by affecting the function of pancreatic beta cells, requiring insulin therapy even in euglycemic patients.^{69,47} In these circumstances, hyperglycemia creates a potential gateway for both pathogens and suggests strict glycemic control as a possible target to prevent the progression of mucormycosis and SARS-CoV-2 infection.

4 CONCLUSIONS

Mucormycosis and COVID-19 coinfection demonstrates outstanding strategies to invade the human organism and survive against the immune response. These actions include taking advantage of a weakened immune system with a reduced response, which given the virulence factors of these pathogens, is very limited. The expression of ligands that interact with human receptors, such as the Coth3-GRP78 and spike protein-GRP78 interaction, enable Mucorales and SARS-CoV-2, respectively, to enter the host. Highlighting the opportunism and invasion skills of these microorganisms, host factors as diabetes, iron overload and systemic corticosteroid therapy facilitate these infections. However, understanding the mechanisms Mucorales and SARS-CoV-2 operate to accomplish the infection reveals a key point to, possibly, an effective therapy. Strict glycemic control, as well as adequate management of iron stores, might destabilize the main point of the infectious process, which relies on iron and glucose to establish itself. Without these components, the virulence of Mucorales is affected and the host's immune response can operate more effectively. In addition, GRP78 might be a therapeutic target to prevent the entry of both pathogens. Anti-GRP78 Abs therapy, as well as anti-CotH and anti-SARS-CoV-2 allied to supportive care approaches, could prevent the invasion of Mucorales and Coronavirus and improve its fatal outcomes.

5 CONFLICT OF INTEREST

No conflict of interest declared.

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APPENDICES

Figure 1: Flowchart of research evidence and criteria for inclusion and exclusion of studies.

Figure 2: Mechanisms of Mucorales FTR1 complex involved in iron sequestration during mucormycosis. **A** – Hyperglycemic state of diabetic patients leads to a glycosylation (Glc) of transferrin and ferritin, releasing ferric iron (Fe+3) into the blood stream, which is carried through the fungal cell membrane by the FTR1 complex. **B** – Ferric iron contained in red blood cells is stripped from heme and transported intracellularly by the high affinity iron permease system. **C** – Chelating agents used as therapeutic measures during iron overload in diabetic patients also provide source of a ferric iron, which is carried across the fungal cell membrane by the FTR1 system.

Table 1: Characteristics of the case reports of mucormycosis and COVID-19 coinfection

Table 1: Continued

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Note: Caspofugine: CAS; Amphotericin B: AMB; Voriconazole: VRC; Amoxicillin-clavulanate: AMC; Hydroxychloroquine: HCQ; Meropenem: MEM; Azithromycin: AZM; Remdesivir: RDV; Dexamethasone: DEX; Cefepime: FEP; Ceftriaxone: CRO; Prednisolone: PRDL; Isavuconazole: ISA; Fluconazole: FLC; Posaconazole: POS; Vancomycin: VAN; Methylprednisolone: mPRED; Tocilizumab: TCZ; Prednisone: PRED; Piperacillin-Tazobactam: TZP; Lopinavir-Ritonavir: LPV/r; Interferon alfa: IFNα; Cyclosporine: CYA; Tigecycline: TGC; Teicoplanin: TEC; Metronidazole: MTZ; Tobramycin: TOB; Oseltamivir: OTV; Sitagliptin: SG; Metformin: MF; Omeprazole: OMEP; Linezolide: LZD; Imipenem: IPM; Hydrocortisone: HC

Figure 3: Mechanisms of the host immune response mediated by glucocorticoids. **A** – Diverse stimuli can mobilize the JNK and IKKβ pathway leading to the activation of transcriptional factors AP-1 and NFκB, which promotes expression of inflammatory genes that provide an effective immune response. **B** – Glucocorticoids enter the cell and bind to glucocorticoids receptor, inhibiting the JNK and IKKβ pathway, which represses the AP-1 and NFκB transcriptional factors – leading to a dysfunctional immune response.

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Book:

2. Voet D, Voet JG. Biochemistry. New York: John Wiley & Sons; 1990. 1223 p. Chapter from a book:

3. White TJ, Bruns TD, Lee SB, Taylor JW. Amplification and Sequencing of Fungal Ribosomal RNA Genes for Phylogenetics, In PCR Protocols and Applications: A Laboratory Manual. New York, NY: Academic Press; 1990:315-322.

Internet document:

4. FORWARD Act for antifungals. New US initiative: Finding Orphan-disease Remedies with Antifungal Research & Development. 2018. https://www.ecmm.info/wp-content/uploads/FORWARD-BILLS-115hr6562ih.pdf Accessed March 3, 2003

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