

Fungal Pathogens in CF Airways: Leave or Treat?

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Abstract Chronic airway infection plays an essential role in the progress of cystic fibrosis (CF) lung disease. In the past decades, mainly bacterial pathogens, such as *Pseudomonas aeruginosa*, have been the focus of researchers and clinicians. However, fungi are frequently detected in CF airways and there is an increasing body of evidence that fungal pathogens might play a role in CF lung disease. Several studies have shown an association of fungi, particularly *Aspergillus fumigatus* and *Candida albicans*, with the course of lung disease in CF patients. Mechanistically, in vitro and in vivo studies suggest that an impaired immune response to fungal pathogens in CF airways renders them more susceptible to fungi. However, it remains elusive whether fungi are actively involved in CF lung disease pathologies or whether they rather reflect a dysregulated airway colonization and act as microbial bystanders. A key issue for dissecting the role of fungi in CF lung disease is the distinction of dynamic fungal–host interaction

entities, namely colonization, sensitization or infection. This review summarizes key findings on pathophysiological mechanisms and the clinical impact of fungi in CF lung disease.

Keywords Cystic fibrosis · Fungal colonization · *Aspergillus fumigatus* · *Candida albicans* · *Pneumocystis jirovecii* · *Exophiala dermatitidis*

Introduction

Cystic fibrosis (CF) is the most common lethal inherited disease in Caucasians. Although CF is a multiorgan disease, the lung disease remains the main cause for morbidity and mortality in CF patients [1, 2]. Lung disease develops very early in life and is characterized by chronic progressive inflammation and infection [3–5]. In the past decades, marked effort has been put into the development of antibacterial drugs, especially those directed against *Pseudomonas aeruginosa*, the most common pathogen in CF airways besides *Staphylococcus aureus* [6–15], which resulted in a decrease in the prevalence of *P. aeruginosa* in CF patients [16–18]. At the same time, the prevalence of other pathogens such as fungi has increased over the past decades [19–21], although there was one study from the UK reporting a decrease in the prevalence of *Aspergillus* species between 1985 and 2005 [22]. Some hypotheses have been proposed to explain the increase in fungal pathogen recovery from the CF respiratory secretions:

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(a) prolonged use of inhaled antibiotics [23–27] or inhaled corticosteroids [28] and (b) increasing age of the patients [26, 29]. A simple and very important explanation might be the significant improvement of sample processing and microbiological detection methods over the past decades, using improved culture methods [30–33] or non-culture methods such as PCR [20, 34], genotyping [35] or oligonucleotide arrays [36]. The detection methods will be discussed in detail in separate reviews of this thematic special issue (see “Challenges in laboratory detection of fungal pathogens in the airways of cystic fibrosis patients” and “Towards the standardization of mycological examination of sputum samples in cystic fibrosis—results from a French multicenter study”). In this review, we will focus on the clinical impact of fungal pathogens and the cellular pathomechanisms in CF lung disease.

Although the clinical relevance of fungi in CF lung disease is yet a matter of further investigations, lately, an increasingly appreciated role of fungal pathogens has been attributed especially to the most commonly isolated fungi in CF, the *Aspergillus* and *Candida* species [37, 38]. The airways are constantly challenged by fungal spores, and the human immune system has evolved a plethora of defense mechanisms to clear fungal pathogens effectively [39–43]. CF airways, however, might be prone to fungal colonization/infection because of several factors: (a) The underlying CFTR defect leads to defective mucociliary clearance and airway mucus thickening resulting in persistence of pathogens and recurrent infections [2, 44]; (b) CF patients display a local immune dysfunction within the airways either primary due to the CFTR defect [45–47] or secondary due to the chronic overwhelming inflammatory and proteolytic processes [38, 48–51]; (c) certain modifier genes were found to be associated with increased risk of fungal colonization in CF, especially by *A. fumigatus* [52–54]; (d) CF patients with low lung function, representing an advanced stage of the CF lung disease with, e.g., significant bronchiectasis, have greater risk of colonization by *A. fumigatus* [19, 55].

Epidemiology of Fungal Pathogens in the CF Airways

The most commonly isolated fungi include the filamentous molds of *Aspergillus* species, particularly *A.*

fumigatus, and the yeast fungi of *Candida* species, mainly *C. albicans*. The prevalence among CF patients given in past studies differs significantly, ranging from 6 to 60% for *A. fumigatus* [30, 33, 56–61], 35–93% for *C. albicans* [27, 30, 57, 58, 61–64], 3–17% for *Scedosporium* spp. [58, 61, 65–68], 1–19% for *Exophiala dermatitidis* [30, 58, 61, 69–71] and 7 to 38% for *Pneumocystis jirovecii* [72–75]. The variety of fungal prevalence in these studies might be due to: (a) different detection methods used; (b) geographical differences [58]; or (c) microbiome changes due to the application of inhaler in addition to systemic antibiotics over years. Nagano et al. [20] showed that fungi were detected in only 18% of the patients using conventional microbiological culture, while 78% of the patients were tested positive using fungal-specific culture media and 100% using a molecular method. In another study, Borman et al. [58] identified significant differences between CF centers throughout Europe regarding their microbiological methods and detection rates of fungi and concluded that standardization in microbiological laboratories is indispensable. A recent monocentric study from a German reference center for fungal infections examined the prevalence of airway fungi in CF patients over a 5-year period between 2009 and 2013. The most common pathogens were *C. albicans*, *C. dubliniensis* and *C. glabrata* among the yeasts (38, 12 and 9%, respectively), and *A. fumigatus* and *S. apiospermum* (29 and 2%, respectively) as well as the yeast-like *E. dermatitidis* (3%) among the filamentous fungi [61].

Recent mycobiome studies in airway samples from CF patients reported a high variability of fungal pathogens, and the results differ from studies utilizing conventional media-based culture methods [76–79]. These studies showed that fungal microbiota in the lower airways of CF patients are more diverse and complex than previously thought, especially when it comes to cross-kingdom synergy and interactions [76, 80]. This might necessitate novel methods to identify the fungal and bacterial colonization/infection in CF airways as well as different strategies for antimicrobial treatment in CF patients in future.

The epidemiology of fungal pathogens and the microbiome of CF lung disease are discussed in detail in separate reviews within this thematic special issue (see “Organization of patient management and epidemiology of fungal infections in cystic fibrosis” and “The airway mycobiome in cystic fibrosis”).

***Aspergillus* Species and Their Role in CF Lung Disease**

Colonization of the Respiratory Tract by *Aspergillus* Species

The most common filamentous fungi isolated from respiratory secretions from CF patients are *Aspergillus* species, particularly *A. fumigatus*. *Aspergillus* species are saprophytic spore-forming fungi that are ubiquitously found in the environment. The spores (also called conidia) which are very small (2–4 µm in diameter) are inhaled by hundreds every day and directly enter the small airways [81]. In healthy airways, they are instantly cleared without provoking inflammatory responses. However, in CF airways, the spores are able to persist and germinate (hyphae), thus eliciting an immune response with leukocyte infiltration and mucus accumulation [82]. A number of studies have reported a negative association of fungi with the lung function in CF patients. In a cross-sectional study with 7010 CF patients from the European Epidemiologic Registry of Cystic Fibrosis (ERCF), colonization by *Aspergillus* species was associated with slightly impaired lung function [5–10% forced expiratory volume in one second (FEV₁)] [83]. Amin et al. [59] conducted a retrospective study in 230 Canadian children with CF between 1999 and 2006 and found significantly lower FEV₁ in children chronically infected by *A. fumigatus* compared to uninfected individuals. At the same time, these patients were at higher risk of pulmonary exacerbations requiring hospitalization. In another Canadian prospective study between 2005 and 2008 with 446 adult CF patients, the prevalence of *Aspergillus* species was twice as high in patients with frequent exacerbations per year compared to those with less than one exacerbation per year [84]. Noni et al. [55] performed a prospective case–control study in Greece with 80 pediatric CF patients and found a significant impact of chronic colonization by *A. fumigatus* on lung function decline over a 7-year period. Interestingly, vice versa, lower FEV₁ at baseline 4 years before study enrollment was the only factor associated with acquisition of chronic *A. fumigatus* colonization [55]. In a retrospective study with 45 CF children from UK, Saunders et al. [29] concluded that chronic colonization by *A. fumigatus* might be associated with worse lung function,

especially in children with severe lung disease. Importantly, the Australian Respiratory Early Surveillance Team for Cystic Fibrosis (AREST CF) study with 56 CF newborns from Australia highlighted the significant impact of early airway infection within the first 2 years of life as predictors of lung function and found *Aspergillus* species among the harmful pathogens [5]. In our retrospective study comprising 770 CF patients from three CF centers in Germany and Austria, *A. fumigatus* colonization was associated with a more rapid decline in FEV₁ when compared to CF patients who had never had positive culture for *A. fumigatus*. In this study, *A. fumigatus* was also the only risk factor for acquisition of chronic *P. aeruginosa* infection in these patients [85]. Consistently, in a past study, Amin et al. [59] found a significant interaction between *A. fumigatus* and *P. aeruginosa* on the course of lung function. An Irish study by McMahon et al. [86] in 36 adolescent and adult CF patients reported significant changes in the severity of bronchiectasis using computed tomography in the *Aspergillus*-colonized subgroup compared to non-colonized individuals, although Bhalla scores and other radiological parameters as well as FEV₁ were not different. A very interesting study from the AREST CF consortium in Australia with 215 pediatric CF patients found increased inflammatory responses, namely increased neutrophil cell count, neutrophil elastase and interleukin (IL)-8 levels, in bronchoalveolar lavage samples from *Aspergillus*-positive CF children compared to never-infected CF children, while colonization by *Candida* species had no effect [87]. However, other studies could not show any association between colonization by *A. fumigatus* and lung function or radiological abnormalities. In a study from the Netherlands in 259 adult CF patients, de Vrankrijker et al. [26] did not find an independent association between chronic *Aspergillus* colonization and lung function or radiological changes over a 5-year period. Similarly, colonization by *Aspergillus* species did not correlate with lung function or computed tomography imaging in a cross-sectional study with 104 adult CF patients [24].

Other *Aspergillus* species such as the transient pathogens *A. flavus*, *A. niger*, *A. nidulans*, *A. tubingenensis* and the chronic colonizer *A. terreus* are frequently detected in CF airways, yet their clinical impact remains elusive [88–90].

Aspergillus Bronchitis

Aspergillus bronchitis was first described by Shoseyov et al. [91] in 2006, when six CF patients experienced respiratory exacerbations without clinical improvement despite appropriate antibiotic treatment, had positive sputum cultures for *Aspergillus* species and responded to antifungal treatment. Importantly, these patients did not fulfill the criteria for allergic bronchopulmonary aspergillosis (ABPA). In 2013, Baxter et al. [92] proposed a novel classification for *Aspergillus*-related lung disease in CF using *Aspergillus*-specific PCR and measurement of sputum galactomannan alongside with detection of total IgE and specific anti-*A. fumigatus* IgE and IgG serum antibodies (Table 1). By this way, the enrolled 130 adult CF patients were subdivided into four groups: (1) non-diseased (37.7%)—either positive or negative PCR, no specific anti-*A. fumigatus* antibodies, negative galactomannan; (2) ABPA (17.7%)—positive PCR, elevated total IgE and specific anti-*A. fumigatus* IgE and IgG, positive galactomannan; (3) *Aspergillus* sensitized (AS) (14.6%)—either positive or negative PCR, elevated specific anti-*A. fumigatus* IgE (but not IgG), negative galactomannan; and (4) *Aspergillus* bronchitis (30%)—positive PCR, elevated specific anti-*A. fumigatus* IgG (but not IgE), positive galactomannan. Notably, 30% of these patients fell into the group with *Aspergillus* bronchitis. Furthermore, in this 2-year prospective study, FEV₁ dropped significantly faster in the CF patients with ABPA, AS and *Aspergillus* bronchitis compared to the non-diseased CF patients [92]. This classification might be helpful to discriminate *Aspergillus* colonization from *Aspergillus* bronchitis, and AS from ABPA, and, more importantly, might be a useful tool to identify those patients who would benefit from antifungal therapy.

Aspergillus Sensitization

The prevalence of AS in CF varies from 20 to 65%. In a recent review, Maturu and Agarwal [93] performed a meta-analysis on 41 studies and calculated the pooled prevalence to be 39%. Interestingly, studies using cutaneous testing showed higher prevalence (43%) compared to studies using serological testing (33%) [93]. AS is defined by T helper cell type 2 (Th2) immune response and a subsequent increase in IgE-mediated immune response to *Aspergillus* species without fulfilling the diagnostic criteria of ABPA. The role of AS in CF lung disease and its association with poorer lung function are well established by a number of studies [60, 92, 94–98]. In the above-mentioned study by Baxter et al. [92], CF patients with AS (elevated IgE, normal IgG, negative galactomannan, PCR positive or negative) experienced a significantly higher FEV₁ decline over the course of 2 years compared to the group of non-diseased patients. Furthermore, Baxter et al. [60] reported that sensitization to *Aspergillus*, but not to *Candida*, was associated with greater FEV₁ decline over a 2-year period and with increased numbers of pulmonary exacerbations in a cohort of 55 CF adults. Interestingly, there was no association between *Aspergillus* or *Candida* colonization and the development of serological sensitization. In another retrospective study over two 5-year periods from 1996 to 2000 (19 sensitized and 19 non-diseased CF children) and 2001–2005 (24 sensitized and 23 non-diseased CF children), AS was associated with changes in pulmonary function only in the first period. The AS patients in the second cohort had significantly higher baseline FEV₁ values compared to those from the first cohort, and this was associated with significantly more oral antifungal treatment in the second cohort [98].

Table 1 Classification of *Aspergillus*-related lung disease in CF, modified from Baxter et al. [92]

	<i>Aspergillus</i> colonization	ABPA	<i>Aspergillus</i> sensitization	<i>Aspergillus</i> bronchitis
<i>Aspergillus</i> -specific PCR	±	+	±	+
Sputum galactomannan	–	+	–	+
Total IgE	–	+	+	–
Specific anti- <i>A. fumigatus</i> IgE	–	+	+	–
Specific anti- <i>A. fumigatus</i> IgG	–	+	–	+

ABPA allergic bronchopulmonary aspergillosis

Allergic Bronchopulmonary Aspergillosis (ABPA)

ABPA is the most well-characterized and well-recognized *Aspergillus*-related disease in CF. The given prevalence varies significantly among the studies, ranging from 3 to 25% with a pooled prevalence of 9% in the meta-analysis of Maturu and Agarwal [93], being higher in adults (10.1%) compared to children (8.9%). ABPA is caused by an exaggerated hypersensitivity reaction to antigens of various fungi, mostly *A. fumigatus*, in individuals with chronic lung diseases such as asthma and CF. Th2-associated immune response with elevated cytokine levels for IL-4, IL-5 and IL-13 results in upregulation of low-affinity IgE receptors on B cells, which subsequently leads to an increase in total IgE and specific anti-*Aspergillus* IgE and IgG, causing type I and type III hypersensitivity reactions [99–101]. Without early diagnosis and sufficient treatment, ABPA becomes chronic and persistent inflammation results in bronchiectasis and severe lung fibrosis [40, 102, 103]. The role of ABPA in lung function decline, and unstable disease was demonstrated in a number of studies [92, 97, 104–107]. Kraemer et al. [97] demonstrated a significant negative effect of ABPA on FEV₁ and forced expiratory flow at 50% (FEF₅₀) in a retrospective study with an observational period of 28 years in 122 mostly pediatric CF patients. In a prospective study in 50 CF adults, Chotirmall et al. [105] described a mean transient decline of 6.7% in predicted FEV₁ in patients with ABPA followed by a normalization after treatment. Another study in 56 adult CF patients analyzed the factors associated with the time until the next pulmonary exacerbation (TUNE) and found ABPA aside of CF-related diabetes to be correlated with shorter TUNE [107]. A study from Germany found a significant association of pet ownership with ABPA and increased numbers of pulmonary exacerbations in 109 CF patients [108]. In the aforementioned study by Baxter et al. [92], CF patients with ABPA had a more pronounced FEV₁ decline over a 2-year period compared to the non-diseased individuals.

However, the optimal method to distinguish between AS and ABPA in CF is still a matter of debate. For example, different variable cutoff values for specific anti-*Aspergillus* IgE and IgG have been proposed [92, 98, 101, 109]. Furthermore, novel markers such as thymus- and activation-regulated chemokine (TARC)/CCL17 [100, 110],

galactomannan in sputum or *Aspergillus* recovery by PCR have been introduced as diagnostic tools for *Aspergillus*-related disease entities in CF [92]. Other studies have suggested using the basophil activation test (BAT) by measurement of CD203c expression on basophils upon stimulation with *Aspergillus* antigen as a diagnostic tool for ABPA [111] or discriminating AS from ABPA [112, 113]. Recommendations from the Cystic Fibrosis Foundation Consensus Conference in 2001 were published in 2003 for diagnostic and screening criteria as well as treatment [114]. Nevertheless, a large part of the studies on ABPA in CF published after issue of this consensus paper used different diagnostic criteria [93]. This might be due to a need for improved diagnostic and monitoring techniques for ABPA in CF [92, 115].

Clinical Impact of Other Fungal Pathogens in the CF Airways

Colonization by Non-*Aspergillus* Filamentous Fungi

Molds belonging to the *Scedosporium* genus and *Lomentospora prolificans* (formerly *Scedosporium prolificans*) are the second most common filamentous fungi detected in the airways of CF patients [65]. Normally, *Scedosporium*/*Lomentospora* infections include sinusitis, pneumonia, lung fungus ball, chest abscesses or disseminated infections in immunocompromised patients [116]. While there are case studies reporting detrimental effects of *Scedosporium*/*Lomentospora* infections in CF patients without previous lung transplantation, the clinical impact of these pathogens is not fully understood. One case study reported on the development of a lung mycetoma, respiratory distress and suggestive cerebral involvement due to neurological symptoms within 18 months after initial isolation of *S. apiospermum* in a young male CF patient. After neurological symptoms occurred, the clinical state deteriorated dramatically, resulting in coma and death within a few days [117]. Another case report showed a rapid clinical impairment within 2 years in a young female CF patient who had excellent exercise tolerance before the acquisition of *S. apiospermum*. After numerous therapeutic approaches including systemic and inhaled antimycotics, intravenous antibiotics, corticosteroids and

immunoglobulins, in the end, lung transplantation was required [118]. In a third study, a 11-year-old CF male presented with respiratory distress and reduced physical tolerance. After isolation of *S. apiospermum* for the first time and insufficient response to intravenous voriconazole and corticosteroid therapy, respiratory distress worsened with a significant decline in FEV₁, necessitating therapeutic bronchoscopy. Bronchoalveolar lavage cultures were positive for *S. apiospermum* only, which showed resistance to voriconazole. The authors suspected a plastic bronchitis due to *S. apiospermum* [119]. Despite these case reports, systematic studies have not found any significant impact of *Scedosporium/Lomentospora* species on lung function. Blyth et al. [67] did not find any association between FEV₁ and colonization by these fungi in their prospective study with 69 adult CF patients. This was confirmed in another recent prospective study with 161 adolescent and adult CF patients showing no difference in FEV₁ between colonized and non-colonized patients. In the latter study, colonization by *Scedosporium/Lomentospora* species was associated with younger age, less colonization with *Haemophilus influenzae* and mucoid *P. aeruginosa* [120]. Interestingly, in both prospective studies, CF patients colonized by these fungi tended to have higher FEV₁ values compared to the non-colonized individuals, although not statistically significant. Nevertheless, colonization by *Scedosporium/Lomentospora* species was associated with higher risk of ABPA [33, 120] or was even linked to symptoms of allergic bronchopulmonary disease similar to ABPA but independently from *Aspergillus* species [65].

Exophiala dermatitidis grows as a black yeast at 37 °C and as a filamentous fungus at room temperature. In CF airways, *E. dermatitidis* may be present transiently or chronically, but usually without any clinical signs. A single case study from Germany reported on a young CF girl who presented with respiratory distress symptoms due to pneumonia with parahilar mottles in the X-ray, which is atypical for CF [121, 122]. In two prospective studies from a Swedish cohort with 98 CF patients aged more than 12 years, *E. dermatitidis* or increased serum levels of IgG against *E. dermatitidis* were associated with pancreatic insufficiency or more frequent colonization by non-tuberculous mycobacteria, increased blood inflammation markers, more frequent intravenous antibiotic treatment and lower FEV₁, respectively [70, 123].

Although the clinical impact of this pathogen is an ongoing matter of research, its presence in the airways should be considered as a CF biomarker and isolation in context with bronchiectasis should dispose the physician to search for an atypical manifestation of CF (see *Exophiala* colonization and infection as a surrogate marker for cystic fibrosis in the elderly in this special issue).

Other filamentous fungi such as *Rasamsonia argillacea* (formerly *Geosmithia argillacea*, but first reported in CF as *Penicillium emersonii*) or *Acrophialophora fusispora* are almost exclusively isolated from CF patients [124]; however, there have been very limited studies on the clinical relevance of these fungi in CF [125, 126].

Colonization by Yeasts

Chronic colonization by *Candida* species was similarly associated with aggravated CF lung disease. In the aforementioned cross-sectional study from the ERCF, colonization by *Candida* species was associated with slightly impaired lung function (5–10% FEV₁) in CF patients [83]. In a prospective longitudinal study over 11 years from Ireland in 89 adult CF patients, intermittent and chronic colonization by *C. albicans* was a significant predictor for hospitalization due to pulmonary exacerbation and accelerated the decline in FEV₁ and body mass index (BMI) [62]. Another prospective longitudinal study over a 6-year period in 91 pediatric and adult CF patients reported that baseline FEV₁ was significantly lower and annual decline was significantly higher in patients chronically colonized by *C. albicans* compared to non-colonized patients, while intermittently colonized individuals had intermediate values. However, in this study, no difference was observed in BMI between the groups, although BMI <20 kg/m² was an independent predictor for chronic *C. albicans* colonization. Furthermore, colonization by *A. fumigatus* was significantly associated with chronic *C. albicans* colonization and was also an independent risk factor for chronic *C. albicans* colonization [64]. In our previously mentioned longitudinal study, we did not find any negative association between *C. albicans* colonization and lung function or BMI [85].

Other common members of the *Candida* species include *C. dubliniensis*, *C. glabrata*, *C. krusei*, *C. parapsilosis*, *C. lusitaniae* and *C. tropicalis*. Very little

is known about the impact of these pathogens in CF lung disease. For *C. glabrata*, the above-mentioned study by us showed a more rapid FEV₁ decline over time in patients colonized by this pathogen compared to those patients who have never had positive culture [85]. In two prospective studies with 54 or 20 pediatric and adult CF patients, both conducted to investigate the prevalence of *C. dubliniensis* in CF airways, all patients colonized by *C. dubliniensis* were clinically stable over the study period of 7 months or 2 years, respectively [127, 128].

Colonization by *Pneumocystis jirovecii*

Pneumocystis jirovecii, initially described as a protozoan in the early twentieth century, was classified officially as a fungus in 1999. While *P. jirovecii* pneumonia (PJP) plays a major role in morbidity and mortality in hematologic malignancies and in immunodeficient and immunosuppressed patients [129, 130], the role of this pathogen in CF lung disease is unknown. A recent case report described a pulmonary *P. jirovecii* infection in a female CF toddler that was successfully treated with cotrimoxazole [131]. A recent prospective longitudinal study over 1 year in 111 adult CF patients found that *P. jirovecii* was more likely isolated when patients experienced an acute pulmonary exacerbation. However, pulmonary exacerbation with positive *P. jirovecii* detection was milder as suggested by lower pulmonary exacerbation score and lower serum C-reactive protein levels [132]. In a prospective multicenter study from France in 104 adult CF patients, colonization with *P. jirovecii* was associated with the absence of *P. aeruginosa* and with higher FEV₁ values [75]. Furthermore, no PJP was observed and patients were stable in two prospective studies with 95 or 88 CF patients colonized by *P. jirovecii* over a 2-month or a 1-year follow-up period, respectively [72, 73]. Although it has been shown that *P. jirovecii* might induce inflammatory changes and airway obstruction in chronic obstructive pulmonary disease [133, 134], there is no evidence until now that it might play a significant role in CF lung disease. A possible explanation might be that CF airways are more frequently colonized by some less pathogenic *P. jirovecii* genotypes compared to the genotypes found in immunosuppressed patients suffering from PJP [135].

Treatment of Fungal Colonization and Infection in CF Lung Disease

Although a number of case studies on treatment of CF patients with fungus-related lung diseases have been published, almost no randomized, controlled trials have been conducted to date. *Aspergillus* colonization and infection are treated with antimycotics, while ABPA is mostly treated with corticosteroids and sometimes with adjunctive itraconazole or voriconazole. Furthermore, the successful use of the anti-IgE drug omalizumab in CF-ABPA has been reported in several case reports [136–147]. Only one double-blind, randomized, placebo-controlled study was initiated for omalizumab, but has been prematurely terminated due to the inability to recruit participants into the study [148]. For antifungal therapy, the only completed randomized, controlled trial to date was conducted by Aaron et al. [21] in a total of 35 adolescent and adult CF patients with *A. fumigatus* colonization (non-ABPA), 18 receiving itraconazole 5 mg/kg/d and 17 receiving placebo over 24 weeks. No beneficial effect was observed after treatment with itraconazole. On the contrary, the time to first pulmonary exacerbation was shorter in the treatment group (77 days) compared to the placebo group (134 days), without reaching statistical significance. However, this study had two major limitations—firstly, the number of patients were too small, and secondly, itraconazole blood levels were subtherapeutic in 43% of the treated patients [21]. A French multicenter, open-label trial on the impact of itraconazole and voriconazole in *Aspergillus*-infected CF patients has been completed, recently, but the results have not been published, until now (clinicaltrials.gov). In a single-center open-label study, 13 CF patients with *A. fumigatus* colonization (non-ABPA, non-AS with total serum IgE < 190 IU/ml, negative cutaneous test to *Aspergillus* antigens) were treated with itraconazole over 6 weeks, resulting in a significant decrease in airway fungal burden, improved respiratory symptoms and mosaic pattern changes in computed tomography as well as preserved lung function and a decrease in episodes of pulmonary exacerbations in the follow-up period of up to 12 months. The authors concluded that *A. fumigatus* eradication by itraconazole resulted in reduced levels of the fungal metabolite gliotoxin that was responsible for down-regulation of the nuclear vitamin D receptor.

Decreased vitamin D receptor expression led to increased production of IL-5 and IL-13 and hence to an enhanced Th2-associated immune response. Therefore, the beneficial effect of high-dose vitamin D₃ supplementation (4,000 IU/day) in AS patients with history of ABPA as shown by Nguyen et al. [149] might be hampered by *A. fumigatus* gliotoxin and might necessitate concurrent elimination of *A. fumigatus* in order to permit vitamin D receptor expression [34]. Hilliard et al. [150] reported on a retrospective study in 21 pediatric CF patients receiving voriconazole over various durations (1–50 weeks), with diverse clinical indications and either as a monotherapy or in combination with an oral corticosteroid. Thirteen patients had ABPA, and 7 of the remaining 8 patients had a history of *A. fumigatus* colonization or previous ABPA. In the group with ABPA, voriconazole (monotherapy and in combination with corticosteroids) resulted in significant clinical and serological improvement, while it had no effect in the group without ABPA [150]. In another retrospective study encompassing 21 CF patients with ABPA over a 5-year period, treatment with itraconazole (5–10 mg/kg/day for 6–54 months in 14 patients or continuously in 7 patients) as monotherapy had no significant effect on lung function, while in combination with corticosteroids, FEV₁ and FVC significantly increased. Total IgE decreased in 42% of the patients receiving monotherapy and in 56% of the patients on combination therapy [151]. In the aforementioned study by Shoseyov et al. [91], six CF patients with *Aspergillus* bronchitis received itraconazole or voriconazole after antibacterial therapy had been ineffective, and azole treatment improved lung function, nutritional status and serological markers. The study by Kanthan et al. [98] comparing two 5-year periods (1996–2000 and 2001–2005) in a pediatric CF cohort with AS noted that the patients received significantly more antifungal treatment in the 2001–2006 period compared to the first 5-year period, which was associated with better lung function. Furthermore, several case studies on antimycotics reported the safe use of nebulized amphotericin B for ABPA [152, 153] or nebulized voriconazole for *S. apiospermum* [154] in CF. However, there are insufficient data on safety, necessity and effectivity of antifungal treatment in CF lung disease to date, both for ABPA [155] and for other fungus-related disease entities. More randomized, controlled studies are urgently needed to answer the

questions on when (in colonized, bronchitis or sensitized patients?), which compound, route of administration (inhaled or systemically?) and how long (monitoring the fungal burden or clinical improvement?). Importantly, the increasing azole resistance (around 5–8% azole-resistant strains have been reported [156–158]) and drug–drug interactions due to inhibition of cytochrome P450 enzymes (increased serum concentration of corticosteroids and other cytochrome P450-dependent drugs such as Ivacaftor) [159] have to be taken into account when considering eradication therapy of fungal pathogens.

Treatment of fungal infections in CF and azole resistance in *A. fumigatus* will be discussed in detail in separate reviews of this thematic special issue (see “Progress in treatment and prevention of fungal infections in CF” and “Azole resistance in *Aspergillus fumigatus* in patients with cystic fibrosis: a matter of concern?”).

Pathophysiology of Fungi in CF

Figure 1 gives an overview of the pathogen recognition receptors on innate immune cells involved in fungal detection.

Interaction of Fungi with Other Microbes in CF

Yeasts

CF lungs consist of a complex microbial community, and resulting biofilms contain different microbes [79, 160]. Initially, it was believed that biofilms are traditionally monomicrobial in nature. However, there is growing evidence that these biofilms in CF lung provide a niche to many microbes to thrive [161, 162]. *Candida albicans* is one of the major players involved in polymicrobial communities in CF lungs. This yeast can often be found to grow with bacteria in these biofilms, and these pathogen–pathogen interactions lead to unprecedented events in terms of virulence of the pathogens. *P. aeruginosa* and *S. aureus* are the most prevalent opportunistic pathogens in individuals with CF lung disease followed by *H. influenzae* and other species [6, 7, 9, 14, 15]. Several studies have demonstrated that *C. albicans* and *P. aeruginosa* interact with each other in the lung and also cause synergistic infections [163, 164]. Studies show that *P. aeruginosa*

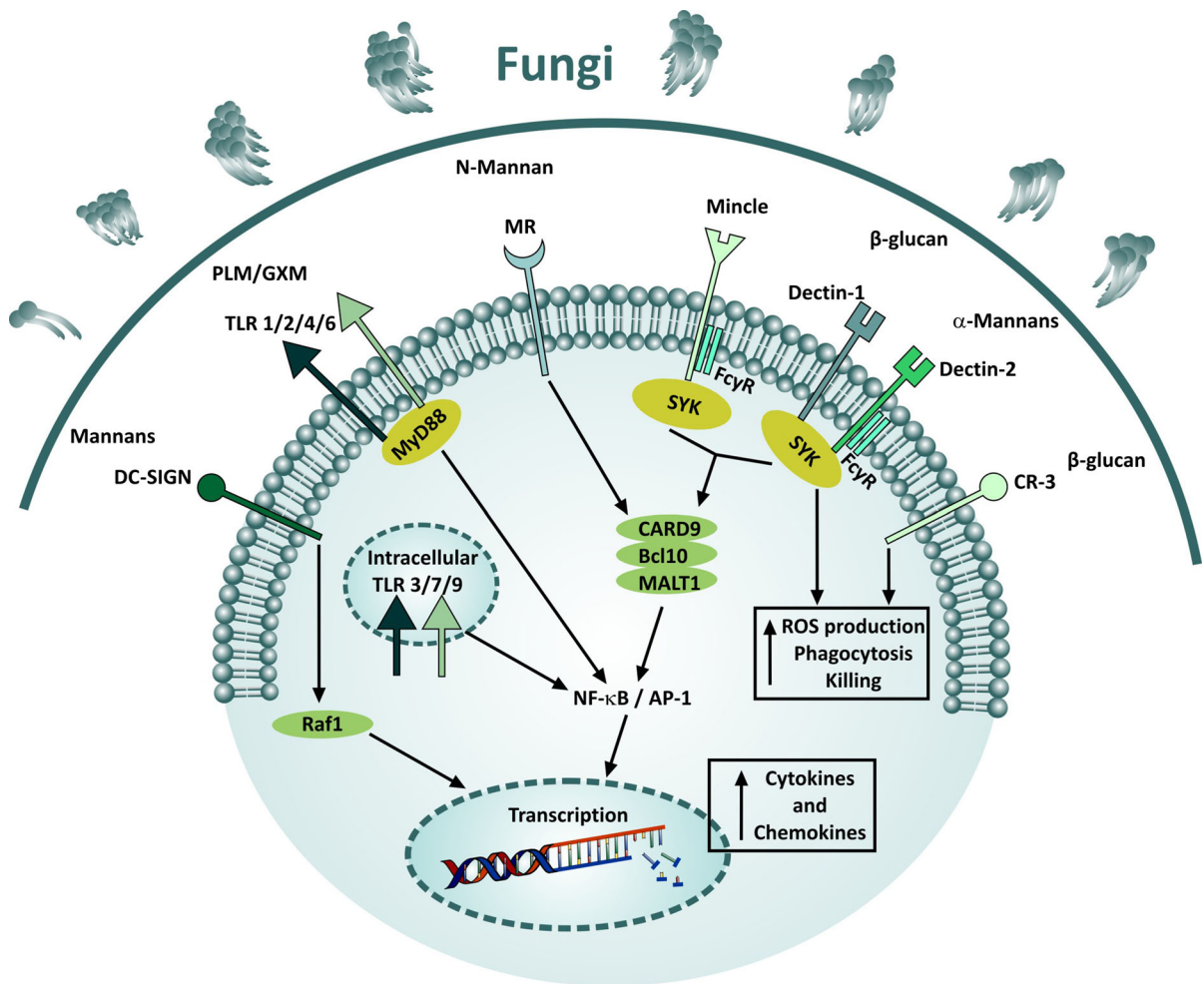


Fig. 1 Fungal recognition by innate immune cells. Fungal pathogens are detected by a variety of pattern recognition receptors (PRRs). Through receptor recognition, signaling pathways are activated leading to increased production of cytokines and chemokines and to engagement of different antifungal immune mechanisms. *AP-1* activator protein 1, *Bcl10* B cell lymphoma/leukemia 10, *CARD9* caspase recruitment domain-containing protein 9, *CR-3* complement receptor 3, *DC-SIGN* dendritic cell-specific intercellular adhesion molecule-3-

grabbing non-integrin, *GXM* glucuronoxylomannan, *MALT1* mucosa-associated lymphoid tissue lymphoma translocation protein 1, *Mincle* macrophage inducible Ca²⁺-dependent lectin, *MR* mannose receptor, *MyD88* myeloid differentiation primary response gene 88, *NF-κB* nuclear factor kappa-light-chain-enhancer of activated B cells, *PLM* phospholipomannan, *Raf1* rapidly accelerated fibrosarcoma 1, *ROS* reactive oxygen species, *SYK* spleen tyrosine kinase, *TLR* Toll-like receptor

can form a dense biofilm on *C. albicans* filaments leading to the killing of fungi. This phenomenon was dependent on morphogenetic changes as only hyphal form of *Candida* was affected by killing and there was no effect on the yeast form [163, 165]. Secreted mediators-based signaling between both the organisms also occurred, validating an important role of *P. aeruginosa* pigments in interaction with *Candida* quorum-sensing molecules. Hyphal formation by *Candida* also helped in acquiring the nutrition for *P. aeruginosa*

[165]. As an example, Chen et al. [166] reported that *Candida* ethanol stimulated the biofilm formation and adhesion of *P. aeruginosa* through inhibition of locomotion. This effect was accompanied by enhanced production of antifungal phenazine derivatives by *P. aeruginosa*, molecules that, in turn, were shown to promote ethanol production within fungi [166]. McAlester et al. [167] reported *C. albicans* farnesol inhibiting the swarming capability of *P. aeruginosa* in CF isolates. Using a combination of clinical and genomic methods,

Kim et al. [168] showed how pathogens and their interactions adapt to the CF environment. The study reported that *C. albicans* isolates from CF patients could go through adaptive mutation in *NRG1* (*Pro-neuregulin-1*), which led to filamentation without any environmental stimulus. Interestingly, these filamentous forms of *C. albicans* were resistant to *P. aeruginosa*-mediated factors, which in general are known to suppress the filamentous growth of the fungus [168].

Staphylococcus aureus is the other most prevalent bacteria in CF lung especially during early onset of the disease. Due to close proximity and niche in the CF lung, *Candida* and *Staphylococcus* also interact with each other at various physical and biochemical levels [169]. Using a mouse model, Carlson et al. [170] showed enhanced mortality of the mice after coinfection with *C. albicans* and *S. aureus*. Physical interaction between *C. albicans* and *S. aureus* led to synergistic effect on biofilm formation, followed by degradation of *S. aureus* biofilm by *C. albicans* farnesol [171, 172]. Apart from *C. albicans*, *C. dubliniensis* are also reported to show an interesting effect in CF environment. A study by Wahab et al. [128] reported that the presence of *C. dubliniensis* in the lower airways of CF patients might be due to their increased adherence in comparison with other *Candida* species. Since *C. dubliniensis* was isolated along with co-colonizing bacteria like *P. aeruginosa* and *S. aureus*, authors speculated that it might hint toward unknown survival strategies of *C. dubliniensis* and coexistence with other pathogens in CF lungs [128].

Aspergillus spp.

Some studies shed light on the interaction of *A. fumigatus* with *P. aeruginosa*. Briard et al. [173] showed that *A. fumigatus* and *P. aeruginosa* could interact at a distance through volatile compounds-mediated communication and, interestingly, *A. fumigatus* growth was enhanced in response to these compounds [173]. On the contrary, Mowat et al. [174] demonstrated that small diffusible and heat-stable molecules of *P. aeruginosa* could potentially inhibit filamentous growth of *A. fumigatus*, making it an interesting phenomenon for the CF lung. In an earlier study, Briard et al. [175] demonstrated interactions between *A. fumigatus* and *P. aeruginosa*, leading to stimulatory or antagonistic effects. They analyzed the

functions of the four *P. aeruginosa* phenazines and their mode of action on *A. fumigatus*. All four phenazines showed *A. fumigatus* growth inhibitory effects. The inhibition of *A. fumigatus* involved production of reactive oxygen species (ROS), specifically O_2^- , and the reactive nitrogen species (RNS) $ONOO^-$. They also found that subinhibitory concentrations of pyocyanin (PYO), phenazine-1-carboxylic acid (PCA) and phenazine-1-carboxamide (PCN) could enhance *A. fumigatus* [175]. In CF, Shirazi et al. [176] demonstrated that biofilm filtrates of *P. aeruginosa* strains isolated from CF patients could inhibit preformed *A. fumigatus* biofilms through apoptosis and non-CF isolates were less inhibitory. Complex and adaptive fungi–bacteria interaction was demonstrated by Moree et al. [177] who showed that reported inhibitors of *A. fumigatus*, phenazines secreted by *P. aeruginosa*, were used as siderophores by *A. fumigatus*. Another mechanism of fungi killing was described showing bacteriophage Pf4 produced by *P. aeruginosa* inhibiting *A. fumigatus* biofilm by iron sequestration [178]. The role of IL-10 in CF during infection with *A. fumigatus* and *P. aeruginosa* was addressed by Casaulta et al. [179] using an in vitro model. Using peripheral blood mononuclear cells (PBMCs) isolated from CF patients, the authors demonstrated that they secreted large amounts of IL-10 upon exposure to *A. fumigatus* antigens or heat-inactivated *P. aeruginosa* which controlled T cell response. Blockage of IL-10 led to a significant increase in T cell proliferation and interferon- γ production [179].

Other Fungi

For other fungal species co-colonizing the CF airways, evidence is still scarce in terms of their interactions with other microbes. A study by Kaur et al. [180] showed that *P. aeruginosa* inhibits the growth of *Scedosporium aurantiacum*. They also demonstrated that *P. aeruginosa* biofilm formation is important, but not crucial for inhibiting the growth of *S. aurantiacum* in a lung-mimicking model [180]. A recent study by Schwarz et al. [102] presented a novel association of ABPA and *Pseudomonas* as a new potential risk factor for *Scedosporium/Lomentospora* infections. It was also speculated that allergy might be involved in inducing immunological host reactions which eventually lead to a less effective response to *Scedosporium/Lomentospora* infections [102].

Immune Recognition of Fungi

Yeasts

Cell wall is one of the most important factors responsible for the recognition of *Candida* by the host immune system. The outer layer of *Candida* cell wall is composed of β -(1,3)-glucan polysaccharide fibrils, which are covalently linked with chitin (a β -(1,4)-linked polymer of *N*-acetyl glucosamine) and β -(1,6)-glucans. The outer layer is made of *N*- or *O*-linked mannosylated proteins termed as mannans. These cell wall polysaccharides or pathogen-associated molecular patterns (PAMPs) are recognized by various pattern recognition receptors (PRRs) present on the host cells [181]. Major PRRs that are involved in recognition of fungal pathogens by the host are mainly classified among the Toll-like receptors (TLRs) and C-type lectin receptors (CLRs). These TLRs and CLRs are expressed on the surface of different cell types that participate in antifungal host response like airway epithelial cells, macrophages, neutrophils and monocytes [182–184]. The expression of TLR1–TLR6 and TLR9 have been reported by Greene et al. [185] on CF tracheal epithelial cell lines, and no difference was found on TLR expression between both CF and non-CF cell lines. On the other hand, Muir et al. [186] also demonstrated the presence of TLR1–TLR10 in CF bronchial epithelial cell lines and an increased TLR2 expression in the CF conditions. While membrane-bound mannoprotein-recognizing TLRs such as TLR2, TLR4 and TLR6 have been described as the most important TLRs for recognition of *Candida* species [187], there is also evidence suggesting that the intracellular TLRs that recognize cytoplasmic nucleic acids, namely TLR3 and TLR9, may also have a role in anti-*Candida* host defense [188]. Also, TLR9 has been found to be expressed on the surface of primary airway epithelial cells and in the context of CF, sinonasal epithelial cells isolated from patients with CF-associated chronic rhinosinusitis display an elevated level of TLR9 [189]. A recent study in murine knockout models demonstrated that TLR9, and also nucleotide-binding oligomerization domain-containing protein 2 (NOD2) and mannose receptor (MR), recognized fungal chitin resulting in the production of the anti-inflammatory cytokine IL-10 [190]. These findings also hint toward a broader role of TLR9 in anti-*Candida* immune response.

Candida cell wall β -(1,3)-glucan is exposed in bud scars that are revealed during the process of cell division or hyphal transition [191]. This process facilitates *Candida* recognition by the host and may provide the hint to the host of a transition from fungal colonization to infectious form [192]. The importance of dectin-1 in the control of fungal infections is highlighted by its role in fungal killing and by the susceptibility of dectin-1-deficient mice to *C. albicans* infection [193–195]. However, the role of dectin-1 in fungal host defense remains a topic of debate. While Taylor et al. [195] showed that *dectin-1*^{−/−} mice had increased susceptibility to disseminated candidiasis, another report found that *dectin-1*^{−/−} deficiency was dispensable to *C. albicans* infection [196]. Furthermore, in human, dectin-1 deficiency has been studied in a family and it was found that it can lead to susceptibility to certain fungal infections including chronic mucocutaneous and recurrent vulvovaginal candidiasis [197]. Apart from dectin-1, other CLRs such as dectin-2, mannose receptor (MR) and Mincle also play a role in recognition of *C. albicans*. Dectin-2 recognizes the α -mannan of *C. albicans* and is mainly expressed on macrophages, dendritic cells (DCs) and neutrophils and modulates T helper cell responses [198]. MR recognizes N-linked mannans present on the fungal cell wall. MR is primarily present on macrophages and was shown to be responsible for further IL-17 induction [199]. Mincle is reported to be expressed on monocytes and neutrophils. It has been shown that Mincle-deficient mice are highly susceptible to systemic candidiasis [200]. However, the ligand for Mincle is still unknown.

Aspergillus spp.

Using various in vitro studies, both TLR2 and TLR4 signaling pathways have been associated with *Aspergillus*-mediated proinflammatory cytokine production and NF- κ B translocation [201, 202]. Studies showed that blocking of TLR4 resulted in decreased TNF α production by adherent monocytes [203]. There is also evidence suggesting that morphology-based antifungal immune response takes place during *Aspergillus* pathogenesis via TLR2 and TLR4 as *Aspergillus* conidia and hyphae showed a different effect [204]. Further studies suggested that intracellular TLRs, TLR3 and TLR9 also play an important role during *Aspergillus* host defense. A TLR3-mediated

mechanism was reported for protection against *A. fumigatus* in epithelial cells [205, 206]. Similarly, TLR9 could recognize *Aspergillus* DNA and was actively recruited during phagocytosis of *A. fumigatus* conidia [207].

Inflammatory cytokines are produced in CF airways by continuous activation of NF- κ B and TLR-independent neutrophil recruitment. Also, in CF, TLR2 and TLR5 are highly expressed on apical surfaces [208]. During infection in CF, further NF- κ B is activated due to TLR signaling. Now impaired TLR4 signaling in CF further prevents the activation of immune response resulting in failure in pathogen clearance and recurrent chronic infections [209]. The CLR dectin-1 recognizes β -(1,3)-glucans and is involved in the recognition of *A. fumigatus* [210]. Human airway epithelium and other immune cells express dectin-1 [211], making it an important candidate for *Aspergillus* recognition. Binding of dectin-1 to *Aspergillus* germ tubes leads to augmentation of TLR2-mediated proinflammatory cytokine production [212]. Apart from germ tubes, swollen *Aspergillus* conidia are also recognized by dectin-1 due to enhanced surface expression of β -(1,3)-glucan [210]. There are ample studies showing dectin-1-deficient mice being extremely susceptible to pulmonary aspergillosis [213–215]. Apart from dectin-1, there is some evidence of involvement of other CLRs, namely the dendritic cell-specific intercellular adhesion molecule-3-grabbing non-integrin (DC-SIGN), MR and dectin-2 in anti-*Aspergillus* host defense. DC-SIGN is expressed on DCs as well as macrophages and has been shown to be involved in binding and phagocytosis of *A. fumigatus* [216]. While the MR has been shown to induce production of proinflammatory cytokines by *A. fumigatus* conidia [204], dectin-2 has been found to mediate the release of inflammatory lipid mediators from murine bone marrow-derived DCs [217]. However, the function and importance of these CLRs in anti-*Aspergillus* host immune response needs to be elucidated.

Along with TLRs and CLRs, the secreted receptor pentraxin 3 (PTX3) has been also shown to play an important role in CF conditions. Polymorphisms in the *PTX3* gene were found to correlate with the risk of *P. aeruginosa* infections in CF patients [218], and PTX3 was found to be localized with *P. aeruginosa* in sputum of CF patients [219]. Paroni et al. [219] also explored the therapeutic potential of PTX3 in CF

during *P. aeruginosa* infection, by using a CF mouse model. Administration of PTX3 led to diminished inflammation in CF by a decrease in inflammatory cytokine levels and improved bacterial clearance by enhancing the phagocytosis capability of CF neutrophils [219]. Some studies shed light on the relevance of PTX3 and fungi in CF. Garlanda et al. [220] reported that conidia of *A. fumigatus* induced the PTX3 secretion in the lungs during pulmonary *Aspergillus* infection. It further binds to galactomannans of the conidial wall and facilitates the phagocytosis by macrophages [220]. Hamon et al. [50] reported decreased levels of PTX3 in the respiratory secretions of CF patients, even though PTX3 level in the CF serum was high. This decrease in PTX3 was attributed to proteolytic cleavage by elastases and *Aspergillus* protease enzymes [50]. This leads to an interesting speculation that lower PTX3 levels in CF lung might correlate with recurring fungal colonization. There is also evidence regarding involvement of the serum opsonin, H-ficolin, in modulation of the host immune response. H-ficolin binding to *A. fumigatus* led to enhanced activation of the lectin complement pathway and lung epithelial cell association with fungal colonization [221].

To summarize, there is growing evidence suggesting that patients with CF have an impaired immune response toward pathogens. Studies reporting the lack of PTX3 in the lung of CF patients, higher expression of TLR2 on apical surfaces of CF epithelial cells and defect in appropriate execution of TLR4 signaling-mediated immune response in CF hint toward some potential mechanisms responsible for recurrent fungal colonization and infection in the CF environment. In the last decade, there has been significant progress in the field of analyzing, processing and handling the CF patient samples in terms of diagnosis and identification of microorganisms colonizing the CF airways. Further studies to dissect the function of fungal PRRs and the impact of colonizing fungi on patient's health are necessary to uncover novel mechanisms and broaden our understanding on the relevance of fungal abundance in CF lung disease.

Conclusions

Novel detection methods have revealed that CF airways harbor a plethora of microbial species, including a vast

variety of fungi [76–79]. In this review, the possible role of a range of fungi, including filamentous molds such as *Aspergillus* species, *Scedosporium* species, *L. prolificans* and *E. dermatitidis*, as well as yeasts such as *Candida* species and other fungi such as *P. jirovecii*, in CF lung disease has been discussed. There is an increasing body of evidence, at least for certain fungi such as *Aspergillus* and *Candida* species, supporting a harmful role in CF lung disease rather than being mere bystanders [5, 55, 59, 62, 64, 84, 85]. However, while a few studies have proposed that fungi are associated with aggravation of CF lung disease [60, 92, 222], multicenter studies corroborating these findings are lacking. Until now, there is only one published prospective, randomized, controlled study that examined the effect of antifungal therapy on pulmonary outcome in CF patients [21]. Consequently, prospective, randomized, controlled studies on antifungal treatment in CF patients that take different fungal disease entities into account are urgently needed.

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