Ficolins Promote Fungal Clearance in vivo and Modulate the Inflammatory Cytokine Response in Host Defense against *Aspergillus fumigatus*

Ninette Genster^a Elisabeth Præstekjær Cramer^b Anne Rosbjerg^a
Katrine Pilely^a Jack Bernard Cowland^b Peter Garred^a

^aLaboratory of Molecular Medicine, Department of Clinical Immunology, Rigshospitalet, Faculty of Health and Medical Sciences, University of Copenhagen, and ^bThe Granulocyte Research Laboratory, Department of Hematology, Copenhagen University Hospital, Copenhagen, Denmark

Key Words

Ficolins · Lectin pathway · Opsonization · Knockout model

Abstract

Aspergillus fumigatus is an opportunistic fungal pathogen that causes severe invasive infections in immunocompromised patients. Innate immunity plays a major role in protection against *A. fumigatus*. The ficolins are a family of soluble pattern recognition receptors that are capable of activating the lectin pathway of complement. Previous in vitro studies reported that ficolins bind to *A. fumigatus*, but their part in host defense against fungal infections in vivo is unknown. In this study, we used ficolin-deficient mice to investigate the role of ficolins during lung infection with *A. fumigatus*. Ficolin knockout mice showed significantly higher fungal loads in the lungs 24 h postinfection compared to wild-type mice. The delayed clearance of *A. fumigatus* in ficolin knockout mice could not be attributed to a compromised recruitment of inflammatory cells. However, it was revealed that ficolin knockout mice exhibited a decreased production of proinflammatory cytokines in the lungs compared to wild-type mice following *A. fumigatus* infection. The impaired clearance and cytokine production in ficolin knockout mice was

independent of complement, as shown by equivalent levels of *A. fumigatus*-mediated complement activation in ficolin knockout mice and wild-type mice. In conclusion, this study demonstrates that ficolins are important in initial innate host defense against *A. fumigatus* infections in vivo.

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Introduction

The mold *Aspergillus fumigatus* is an opportunistic fungal pathogen and one of the leading causes of invasive fungal infections [1]. Invasive pulmonary aspergillosis, a disease of high morbidity and mortality which affects immunocompromised individuals, is of growing concern due to the increased use of immunosuppressive drugs (e.g. for organ transplantation) and chemotherapy treatments [2]. Despite the availability of newer antifungal agents, the prevention and treatment of invasive pulmonary aspergillosis remain problematic [1, 2]. It is essential to elucidate the contribution of innate and adaptive immune responses to *A. fumigatus* to understand the development of invasive disease and to design targeted immunotherapy and vaccination against the fungus.

A. fumigatus is a saprotroph widespread in nature that produces numerous airborne spores known as conidia. Innate immunity plays a major role in protection against *A. fumigatus*. In immunocompetent hosts, anatomical barriers and professional phagocytes prevent disease when conidia are inhaled on a daily basis [3]. In immunocompromised individuals, however, inhaled conidia germinate and form invasive hyphae which can penetrate pulmonary tissues to cause disease. Upon inhalation of *A. fumigatus* conidia into the lungs, initial recognition is managed by alveolar macrophages which lead to intracellular degradation of the conidia and secretion of proinflammatory mediators which further recruit neutrophils to assist in fungal clearance [3]. The immune response to *A. fumigatus* is highly dependent on the recognition of fungal-specific structures, such as cell wall components, by pattern-recognition receptors. Several classes of cell-associated or soluble pattern-recognition receptors are involved in the initial recognition of conidia, including Toll-like receptors [4–7], C-type lectin receptors [8], collectins [9, 10], and pentraxins [11], which facilitate the interaction of phagocytes with fungi.

Ficolins are a family of soluble pattern-recognition receptors that activate the lectin pathway of complement similar to mannose-binding lectin (MBL) [12–14]. They comprise a collagen-like domain and a fibrinogen-like domain, of which the latter is involved in ligand recognition. Binding of ficolin to target structures on the pathogen surface activates associated serine proteases which results in the cleavage of downstream complement components C4 and C2 and formation of the C3 and C5 convertases. This cascade of events leads to the formation of cleavage products that function in opsonization, cell lysis, and the generation of an inflammatory response [15]. In humans, three ficolins have been identified (ficolin-1/M-ficolin, ficolin-2/L-ficolin and ficolin-3/H-ficolin), whereas in mice two ficolins have been identified (ficolin-A and ficolin-B) [13]. Human and murine ficolins are believed to be closely related based on their phylogenetic and biochemical properties [16]. Ficolin-A is regarded as the murine orthologue of ficolin-2, while ficolin-B is the orthologue of ficolin-1. The third ficolin gene exists as a pseudogene in mice.

Research over the past decade has revealed that ficolins recognize a wide range of pathogens [17–20], and that deficiency or low levels of human ficolins is associated with specific infectious diseases [21–23]. In addition, in vivo studies have demonstrated a protective effect of ficolins in murine models of bacterial and viral infections [20, 24–26]. While in vitro studies suggest that ficolins par-

ticipate in the elimination of *A. fumigatus* [27–31], the in vivo role of ficolins during fungal infection has not yet been addressed. Therefore, we used ficolin-deficient mice to elucidate the role of ficolins in host defense against *A. fumigatus* infection in the present study.

Materials and Methods

Mice

Ficolin-A deficient (*Fcna*^{-/-}) and ficolin-B-deficient (*Fcnb*^{-/-}) mice raised on a C57BL/6J background were kindly provided by Dr. Endo of Fukushima University, Japan [24], and further backcrossed to C57BL/6J for five generations in our laboratory. From these two mouse lineages, a ficolin double knockout (KO) lineage was generated. Because *Fcna* and *Fcnb* are located close together on the same chromosome, we initially crossed *Fcna*^{-/-}*b*^{+/+} (F5) and *Fcna*^{+/+}*b*^{-/-} (N5) mice to obtain double heterozygous offspring that were crossed with *Fcna*^{-/-}*b*^{+/+} mice to generate a double heterozygous mouse carrying the haplotype *Fcna*⁻*b*⁻. The obtained mouse carrying genotype *Fcna*⁻*b*^{+/+}*a*⁻*b*⁻ was subsequently crossed with a C57BL/6J mouse to generate double heterozygous offspring carrying the haplotype *Fcna*⁻*b*⁻, and eventually these were interbred. The resulting offspring (*Fcna*^{-/-}*b*^{-/-} mice and their wild-type, WT, littermates) were set up for homozygous breeding; the mice used throughout this study belong to the F2–F4 generations. Female mice were used at 10–12 weeks of age. The mice were housed in pathogen-free, temperature-controlled and air-conditioned facilities with a 12-hour light/dark cycle. Breeding as well as all animal experiments were conducted at the Department of Experimental Medicine, University of Copenhagen, and National University Hospital, Rigshospitalet, Copenhagen, Denmark. All animal experiments were approved by the Danish Animal Experiments Inspectorate (permission No. 2014-15-2934-996) and conducted according to the national Animal Experimentation Act (LBK No. 474 from May 15, 2014). The review board at the Faculty of Health and Medical Sciences, University of Copenhagen, approved this study (P14-122).

Infection

The *A. fumigatus* strain was obtained from a clinical isolate 6871 (kindly provided by Prof. Luigina Romani, University of Perugia, Italy) and grown on Sabouraud dextrose agar (SDA; Oxoid) for 4 days at 37°C. Conidia were harvested by washing plates with sterile phosphate-buffered saline (PBS; pH 7.4) supplemented with 0.025% Tween 80 (PBS-T). The suspension was filtered through a 40-μm cell strainer (BD Biosciences) to separate conidia from the contaminating mycelium and washed three times in PBS-T before resuspension in sterile PBS. The absence of mycelium in filtrate was verified microscopically when conidia were counted in a hemocytometer and the concentration was adjusted in sterile PBS. The conidial preparations were stored at 4°C for up to 4 months. The conidia viability of the challenge inoculum was confirmed by plating serial dilutions on SDA plates.

Age- and weight-matched mice were anesthetized with isoflurane and a sublethal inoculum dose of 2×10^7 of *A. fumigatus* conidia in a 50-μl PBS suspension was placed on the nostrils for inhalation. Control mice received 50 μl of sterile PBS. After inocula-

tion, all animals fully recovered from the procedure within 30 min. The clinical appearance and body weight of the animals were monitored daily. A pilot study was carried out to determine the optimal dose (i.e. nonlethal with mild clinical signs of disease and weight loss) for challenge.

Quantification of the Fungal Burden

The fungal burden in the lungs was determined by quantitative colony-forming unit (CFU) assay. At selected time points postinfection (p.i.), mice were sacrificed by cervical dislocation and the lungs and spleens were aseptically removed. The wet mass of the lungs was determined using a precision balance before the tissue specimens were homogenized in 2 ml of sterile PBS for 1 min using a mechanical homogenizer (TissueRuptor; Qiagen). Primary homogenates were cultured by serial dilution, plated in triplicate on SDA plates, incubated at 37°C for 24 h, and the number of *A. fumigatus* colonies was counted. The results are expressed as CFU per lung.

Lung Myeloperoxidase Assay

Mice were sacrificed by cervical dislocation 24 h p.i. and the lungs were removed and snap-frozen on dry ice. Lysis buffer (200 mM NaCl, 5 mM EDTA, 10 mM Tris, 10% glycine, 1 mM PMSF, 1 µg/ml leupeptin and 28 µg/ml aprotinin, pH 7.4) was added to the lung tissue before homogenization on ice using the TissueRuptor (Qiagen) and the supernatant was immediately frozen and stored at -80°C until analysis. Samples were diluted 500× and assayed in duplicates by a mouse myeloperoxidase (MPO) ELISA (HK210; Hycult Biotech) following the manufacturer's recommendations.

Cytokine Gene Expression

Mice were sacrificed by cervical dislocation 24 h p.i. and the lungs were removed and stored in RNAlater (Applied Biosystems). Lung tissue was homogenized with a mechanical homogenizer (TissueRuptor; Qiagen) and RNA was extracted using the Maxwell 16 tissue LEV Total RNA Purification Kit (Promega) following the manufacturer's recommendations. Total RNA was reverse transcribed with the MuLV reverse transcriptase (Applied Biosystems No. N808-0018) and cDNA was quantified using a Qubit Fluorometer (Invitrogen). Gene expression was assessed by quantitative real-time PCR with TaqMan probes specific for mouse TBP (mM00446971_m1), B2M (Mm00437762_m1), IL-1β (Mm0434228_m1), IL-6 (Mm00446190_m1), and TNF-α (Mm00443258_m1) obtained from Applied Biosystems. Quantitative real-time PCR was performed in triplicate loading 20 ng/µl (TBP and IL-6) or 10 ng/µl (B2M, IL-1β and TNF-α) of cDNA per reaction and carried out on the Stratagene Mx3005P real-time PCR system (Agilent Technologies). Reaction conditions included incubation at 50°C for 2 min and 95°C for 10 min followed by 40 cycles at 95°C for 15 s and 60°C for 1 min. PCR runs included no-template controls and no-reverse transcriptase controls. The target genes were normalized to a housekeeping gene, TBP or B2M, and results were expressed as fold changes from PBS mock-infected WT mice using the delta-delta Ct method.

Cytokine ELISA Assays

Mice were euthanized 24 h p.i. by intraperitoneal injection of pentobarbital before the trachea was exposed through a midline incision. The trachea was cannulated with a sterile 22-gauge catheter (BD Biosciences) and bronchoalveolar lavage (BAL) was per-

formed by instilling and withdrawing 1 ml of cold sterile PBS five times. The BAL fluid (BALF) was centrifuged and the supernatant was removed, immediately frozen and stored at -80°C. In separate experiments, serum was obtained from submandibular blood collection from live animals 24 h p.i. The protein levels of IL-1β, IL-6 and TNF-α in BALF and serum were measured in duplicates by mouse cytokine-specific Quantikine ELISA kits (R&D Systems) according to the manufacturer's recommendations.

Histology and Immunohistochemistry

Mice were euthanized 24 h p.i. by intraperitoneal injection of pentobarbital before the trachea was exposed through a midline incision. The trachea was cannulated with a sterile 22-gauge catheter (BD Biosciences) and the lungs were expanded with 1 ml of 10% neutral-buffered formalin. Lungs were fixed for 24 h and stored in 70% ethanol prior to paraffin embedding. For histology, tissue sections of 5 µm were stained with hematoxylin and eosin (HE). For Immunohistochemistry, tissue sections of 5 µm were deparaffinized and subjected to antigen retrieval by Proteinase K digestion (Roche) for 15 min. Sections were stained for C3d with a polyclonal goat-anti-mouse antibody (6.7 µg/ml, catalog No. AF2655; R&D Systems) or a normal goat IgG control antibody (6.7 µg/ml, catalog No. AB-108-C; R&D Systems) followed by BrightVision poly-HRP anti-goat (Immunologic) and hematoxylin counterstain. Bright field photographs of slides were obtained and analyzed using a NanoZoomer RS slide scanner (NanoZoomer Digital Pathology System and NDP view software; Hamamatsu). No morphometric quantification was performed.

Complement Component C3a ELISA

The BALF was collected as described above for the cytokine ELISA measurements. The concentration of the complement activation product C3a in BALF was measured in duplicates by a mouse-specific ELISA kit (E-EL-MO0337; Elabscience) following the manufacturer's recommendations.

Statistical Analyses

Data were analyzed using GraphPad Prism software, version 6. Comparisons between groups of WT and KO mice were made using the unpaired Student t test and $p < 0.05$ was considered statistically significant.

Results

*Impaired Fungal Clearance in Ficolin KO Mice following *A. fumigatus* Infection*

To determine whether ficolins participate in host defense against *A. fumigatus*, groups of 10- to 12-week-old WT and ficolin KO mice were intranasally inoculated with a sublethal dose of 2×10^7 *A. fumigatus* conidia. Mice were sacrificed 12–72 h p.i. for the quantification of fungal burden (CFU/lung) in whole lung homogenates (fig. 1a). The results showed that ficolin KO mice displayed statistically significant higher fungal loads than their WT counterparts at 24 h p.i., which demonstrates

Fig. 1. Increased pulmonary fungal loads in ficolin-deficient mice 24 h after *A. fumigatus* infection. WT and ficolin KO mice were intranasally inoculated with 2×10^7 *A. fumigatus* conidia. **a** Mice were sacrificed at the indicated time points for quantification of the fungal load (CFU/lung) in whole-lung homogenates. **b** Weight changes in mice during *A. fumigatus* infection. Results (mean $\pm$ SD) were obtained from two independent experiments ($n = 5$ – 7 /group in each experiment). * $p < 0.05$ for ficolin KO to WT comparisons at each time point (unpaired two-tailed Student's *t* test).

that a lack of ficolins results in increased sensitivity to *A. fumigatus* infection. We found no fungal growth in the spleens of *A. fumigatus*-challenged mice 24 or 48 h p.i., which established that the infection did not disseminate from the lungs. This finding was expected because we utilized a nonlethal model of *A. fumigatus* infection. Accordingly, conidia were almost entirely cleared from the

lungs (fig. 1a) and the body weight was regained (fig. 1b) in both ficolin KO and WT mice 72 h p.i. To elucidate whether both of the murine ficolins contribute to *A. fumigatus* clearance, the fungal burden in ficolin-A KO mice, ficolin-B KO mice and WT mice was determined in subsequent experiments (online suppl. fig. 1; see www.karger.com/doi/10.1159/000447714 for all online suppl. material). Neither ficolin-A nor ficolin-B KO mice showed increased fungal loads over WT mice, which suggests a redundant role of the two murine forms of ficolins in the immune response to *A. fumigatus*.

Pulmonary Cellular Infiltration of *A. fumigatus*-Infected Mice

Next, we tested whether the impaired fungal clearance observed in ficolin KO mice was associated with a reduced recruitment of inflammatory cells to the lungs. Mice were intranasally inoculated with *A. fumigatus* conidia or PBS as a control. After 24 h the lung tissues were collected for histology and stained with HE to assess the cellular infiltration and inflammation (fig. 2). Evaluation of the lung sections revealed the presence of inflammatory infiltrates in both WT and ficolin KO mice challenged with *A. fumigatus* (fig. 2b), and a comparable morphology of the two strains was observed. Normal lung histology was evident in sections from control mice mock infected with PBS (fig. 2a). In another experiment MPO levels were quantified in lung homogenates as a measure of neutrophil influx (fig. 3). The level of MPO in *A. fumigatus*-infected mice was increased 24 h p.i. compared to PBS mock-infected mice, confirming that neutrophils were recruited to the lungs following *A. fumigatus* challenge. There was no statistically significant difference in MPO levels between WT and ficolin KO mice.

Impaired Production of Proinflammatory Cytokines in Lungs of Ficolin KO Mice following *A. fumigatus* Infection

To further evaluate the inflammatory response, we analyzed the lung proinflammatory cytokine profiles following *A. fumigatus* infection. WT and ficolin KO mice were intranasally inoculated with *A. fumigatus* conidia or PBS as a control. Mice were sacrificed 24 h p.i. to measure the mRNA expression of IL-1 β , IL-6 and TNF- α in lung homogenates (fig. 4a). In addition, the expression of IL-10 was measured; however, expression of this cytokine was not detected in any case. The results revealed that expression of IL-1 β and IL-6 was significantly decreased in ficolin KO mice compared to WT mice following

Fig. 2. Lung histology of mice following *A. fumigatus* infection. WT and ficolin KO mice were intranasally inoculated with PBS as controls (a) or 2×10^7 *A. fumigatus* conidia (b) and sacrificed 24 h p.i. Lung tissues were harvested, fixed, and HE stained for

histological examination. Representative lung sections from WT or ficolin KO mice are shown. Images are representative of 2 mice in PBS mock-infected groups and 6 mice in the *A. fumigatus*-infected groups obtained from one experiment.

A. fumigatus infection. The difference in expression of TNF- α did not reach statistical significance. To support these findings, we measured the levels of the three cytokines in BALF 24 h p.i. in another set of experiments (fig. 4b). These results confirmed the mRNA findings; the protein levels of all three proinflammatory cytokines were decreased in ficolin KO mice compared to those in WT mice following *A. fumigatus* challenge, although only differences in IL-1 β levels were statistically significant. The levels of all three cytokines in BALF from PBS mock-infected mice were below the detection limit in both strains. In addition, serum levels of the cytokines were measured; IL-1 β and TNF- α were undetectable in the serum of both ficolin KO and WT mice following *A. fumigatus* challenge. In accordance with the lung cytokine levels, serum IL-6 was decreased in ficolin KO mice over WT mice, although the difference was not statistically significant (online suppl. fig. 2). Altogether, these results show a decreased production of proinflammatory cytokines in ficolin KO mice, which indicates that ficolins modulate the cytokine response following *A. fumigatus* infection.

Fig. 3. Lung MPO concentration following *A. fumigatus* infection. WT and ficolin KO mice were intranasally inoculated with 2×10^7 *A. fumigatus* conidia and sacrificed 24 h p.i. Control mice received PBS. Lung tissues were harvested and the MPO concentration in supernatants of lung homogenates was measured by ELISA. Results represent means $\pm$ SEM from one experiment with $n = 2$ (PBS group) or $n = 6$ (*A. fumigatus* group).

Fig. 4. Impaired production of proinflammatory cytokines in the lungs of ficolin-deficient mice following *A. fumigatus* infection. WT and ficolin KO mice were intranasally inoculated with 2×10^7 *A. fumigatus* conidia and sacrificed 24 h p.i. Control mice received PBS. **a** Lungs were removed and the expression of proinflammatory cytokines in lung homogenates was measured by qPCR; data represent means $\pm$ SEM as the fold increase over the control group

(WT mock infected with PBS). **b** In separate experiments BALF was collected and cytokine concentrations in BALFs were measured by ELISA. The figures illustrate results from one experiment, representative of two independent experiments showing similar results, with $n = 2$ (PBS group) or $n = 6$ (*A. fumigatus* group) in each experiment. * $p < 0.05$ for ficolin KO to WT comparisons (unpaired two-tailed Student's *t* test).

Unimpaired Complement Activation in Ficolin KO Mice following *A. Fumigatus* Infection

Next, we sought to elucidate whether the impaired fungal clearance and cytokine induction in ficolin KO mice was due to a decreased activation of complement. Mice were intranasally inoculated with *A. fumigatus* conidia or PBS as a control; after 24 h the lung tissues were collected for immunohistochemistry and stained for the complement activation product C3d (the antibody also detects C3b and C3bi). Evaluation of lung sections revealed the deposition of C3d in mice challenged with *A. fumigatus* (fig. 5b, top panels); a similar level of C3d deposition was observed in ficolin KO and WT mice. No C3d was evident in lung sections from control mice mock infected with PBS (fig. 5a, top panels) or in sections stained with a control antibody (fig. 5a, b, bottom panels). To support these findings, we quantitatively determined the levels of another complement cleavage product, C3a,

in BALF (fig. 6). The results showed that C3a levels in the BALF of *A. fumigatus*-infected mice were increased compared to PBS mock-infected mice confirming that complement was activated following *A. fumigatus* infection. In agreement with the immunohistochemistry findings there was no difference between ficolin KO and WT mice, which implies that ficolins function independently of complement in the current infection model.

Discussion

Although ficolins have been identified as important pattern recognition molecules for bacterial and viral pathogens in vivo [20, 24–26], their role in fungal infection has not been established. Prior work from our research group has shown that human and murine ficolins bind to *A. fumigatus* in vitro [27, 28]. In the present study,

Fig. 5. Complement deposition in the lungs of mice following *A. fumigatus* infection. WT and ficolin KO mice were intranasally inoculated with PBS as controls (**a**) or 2×10^7 *A. fumigatus* conidia (**b**) and sacrificed 24 h p.i. Lungs were harvested, fixed, and immunohistochemically stained with an anti-C3d antibody (upper

panels) or a control antibody (lower panels). Sections were counterstained with hematoxylin. Representative lung sections from WT or ficolin KO mice are shown. Images are representative of 2 mice in the PBS mock-infected group and 6 mice in the *A. fumigatus*-infected group obtained from one experiment.

we identified a role for the ficolins in a mouse model of *A. fumigatus* infection. Two important points have emerged from this study. First, ficolins promote *A. fumigatus* conidia clearance in the lungs. Second, ficolins modulate the cytokine production following *A. fumigatus* infection.

We found that fungal load was significantly higher in ficolin KO mice compared to WT mice 24 h p.i. (fig. 1a). Recent studies showed that ficolin-A opsonization of *A. fumigatus* conidia enhanced the binding of conidia to A549 airway epithelial cells [29], human monocyte-derived macrophages and neutrophils [30]. Consequently, hyphal growth was inhibited and fungal killing was increased. Earlier studies also recognized that the human homologue of ficolin-A, ficolin-2, exhibits opsonic activity [17, 31]. In the light of these data it seems likely that the delayed fungal clearance in the ficolin KO mice observed in our study could be credited to impaired opsonophagocytosis of the conidia. Interestingly, a model was proposed where pathogen-bound natural IgG:ficolin

Fig. 6. Complement activation in lungs of mice following *A. fumigatus* infection. WT or ficolin KO mice were intranasally inoculated with 2×10^7 *A. fumigatus* conidia and sacrificed 24 h p.i. Control mice received PBS. The BALF was collected and C3a concentration in BALFs was measured by ELISA. Results were obtained from two experiments (n = 4–8 mice in the *A. fumigatus* groups per experiment).

complexes enhance the uptake by phagocytes via FcγR1 independent of C3 [32]. Nevertheless, the precise mechanism by which ficolins promote fungal clearance in the lungs remains to be established. We found that 48 h after the onset of *A. fumigatus* infection the fungal loads were similar in ficolin KO and WT mice, which indicates that ficolins are of importance in the initial phase of infection. It is known that initial recognition of *A. fumigatus* conidia is managed by alveolar macrophages and pattern recognition receptors, whereas neutrophils are recruited subsequently [3]. The long pentraxin-3 (PTX3) is secreted by neutrophils and plays a pivotal role in protection against *A. fumigatus* in mice [33]. Hence, PTX3 constitutes an example of how redundant pathways might compensate in clearing conidia and modulating pulmonary inflammation over time. The objective of this study was to investigate the role of ficolins when fungal conidia are inhaled in the healthy host under normal circumstances. Therefore, we established a nonlethal model of infection in immunocompetent (ficolin KO) mice. It is conceivable that the significance of ficolins is enhanced in the immunocompromised host, e.g. during neutropenic conditions. Indeed, it is in these situations that *A. fumigatus* represents a threat to human health [2]. Therefore, it is highly relevant to further elucidate the role of ficolins during fungal infections in an immunocompromised experimental model. Interestingly, a recent study detected the serum protein ficolin-2 in BALF from the lungs of patients with a diagnosis of invasive *A. fumigatus* infection [34], which highlights a clinical perspective for future studies.

Ficolin-A is a primarily liver-derived serum protein [35], whereas ficolin-B is present in serum only in very low concentrations [24]. It was recently shown that ficolin-B is stored in and released from immature granulocytic myeloid cells, and a role in the early infection-induced cellular response of these inflammatory cells was suggested [36]. We found that mice deficient in only ficolin-A or ficolin-B exhibited a similar clearance of conidia compared to WT mice (online suppl. fig. 1). These results imply that ficolin-A and ficolin-B play redundant roles in the clearance of *A. fumigatus* conidia. In contrast, Endo et al. [24] showed that the two murine ficolins did not work competitively in protection against *S. pneumoniae*; all of the three mouse strains, including ficolin-A KO, ficolin-B KO and double-deficient mice, displayed reduced survival and increased bacterial burdens in the lungs compared to WT mice. Taken together, it appears that the individual roles of the two murine ficolins depend on the pathogen and/or the biological setting.

We found that the observed delayed clearance of *A. fumigatus* conidia from the lungs in the absence of ficolins could not be attributed to impaired recruitment of inflammatory cells (fig. 2, 3). Actually, despite reduced production of proinflammatory cytokines in the lungs of *A. fumigatus*-infected ficolin KO mice, there was no significant difference between ficolin KO and WT mice regarding cellular recruitment to the lungs. These data indicate either that compensating mechanisms for prompting cellular recruitment exist, or that recruitment is independent of these cytokines. Another possibility is that the difference in cellular recruitment between ficolin KO and WT mice was too small to detect with the employed methods.

In our study, *A. fumigatus*-induced production of proinflammatory cytokines was reduced by approximately 30% in ficolin KO mice compared to WT mice (fig. 4). Similarly, a proinflammatory role of MBL in a mouse model of *A. fumigatus* infection has been suggested [9]. Consistent with our data, recent studies demonstrated the ability of ficolins to modulate the inflammatory response. Specifically, it was shown that ficolin-2 stimulated murine macrophages to produce IFN-γ, IL-17A, TNF-α and IL-6 in vitro, and that ficolin-A KO mice exhibited decreased production of TNF-α and IL-17A in vivo upon stimulation with LPS [26]. Other investigators reported that ficolin-A increased the production of IL-8 from human lung epithelial cells following *A. fumigatus* challenge in vitro [29]. On the other hand, the same group showed that ficolin-A or ficolin-2 opsonization of *A. fumigatus* leads to decreased levels of IL-8, IL-1β, IL-6, and TNF-α from human monocyte-derived macrophages and neutrophils [30, 34]. However, the behavior of these cells in vitro might not represent all in vivo functions that are likely influenced by paracrine signals. Cytokines trigger a strong localized inflammatory response required for the hazard elimination, but the response must be tightly regulated to prevent excessive tissue-damaging inflammation. In the setting of overwhelming inflammation, signaling pathways might contribute to a poor outcome. In this study, we have identified a role for ficolins as cytokine modulators. In addition to the opsonizing activity, ficolins are known to activate the lectin pathway of complement via interaction with MBL-ficolin-associated serine proteases [12], a property they share with MBL. Consequently, one mechanism by which ficolins could increase the cytokine response is through activation of complement. However, our results revealed that the complement activation was unimpaired in the ficolin KO mice, which indicates that complement-independent mechanisms must be involved

(fig. 5, 6). Interestingly, it has been shown that MBL promote IL-6 and TNF- α production, not through its capacity to act as an opsonin or to activate complement, but via a cooperation with TLR2/6 [37, 38]. A similar role for ficolins could be hypothesized based on our findings.

As stated above, we found that the absence of ficolins did not reduce complement activation *in vivo* following *A. fumigatus* infection (fig. 5, 6). In addition, we found no difference between WT and ficolin KO serum regarding C3 or C4 deposition on *A. fumigatus* *in vitro* (online suppl. fig. 3). These results are consistent with another study where the contribution of ficolins to complement activation on *A. fumigatus* was investigated. The investigators used WT, ficolin-A- or MBL-deficient mouse sera and reported that ficolin-A-bound *A. fumigatus* conidia did not activate the lectin complement pathway, but that complement deposition was mediated via binding of MBL-C [29]. In addition, PTX3 has been shown to activate complement on *A. fumigatus* [39]. Therefore, it is likely that the potential contribution of ficolins to complement activation on *A. fumigatus* is negligible in the presence of other complement activators. In contrast, it was reported that ficolin KO serum showed lower complement deposition on *S. pneumoniae* compared to WT serum [24, 25]. Importantly, MBL-deficient serum showed the same level of deposition as WT serum, which

indicates that MBL did not drive lectin pathway activation on *S. pneumoniae* [25]. Hence, although MBLs are the major initiators of the lectin pathway, the data suggested that MBLs are not involved in *S. pneumoniae* infection, which might explain the difference to (lack of) ficolin-mediated complement activation on *A. fumigatus*. It should be noted that the ficolin-mediated lectin pathway appears to be more important in humans than in mice; in humans, the serum concentrations of ficolins are several times greater than MBL [13], whereas the opposite is the case for mice [40].

We conclude that ficolins promote the clearance of *A. fumigatus* and modulate the inflammatory cytokine response independent of complement activation. These results demonstrate that ficolins participate in host defense against *A. fumigatus* infections *in vivo*. The study provides supporting evidence to the role of ficolins demonstrated *in vitro* and extend our knowledge of protection against *A. fumigatus* infection.

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