

EDITORIAL

Identifying host immune effectors critical for protection against *Candida albicans* infections

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Candida albicans is one of the most important human fungal pathogens and can manifest as a wide variety of infections, ranging from benign superficial disease to life-threatening invasive and disseminated infection. Humans with intact immune systems are highly resistant to *C. albicans*, and in fact most fungal infections.¹ Thus, many fungal pathogens are labeled as “opportunistic.” But this label unfairly anthropomorphizes the organism and imbues it with an ulterior motive when in fact the fungi is merely responding to a change in the host’s condition or physiology.² Neither the microbe nor the host is static, and it is clearly the dynamics that are critical for determining whether an infection develops. While many researchers define virulence only in terms of the pathogen, virulence may be better defined as “a reflection of the outcome of host-microbe interaction in a susceptible host, rather than a stable or predictable microbial trait.”³

Therefore dissecting and elucidating the host immune effectors critical for maintaining protection against *C. albicans* infections is critical for truly understanding *Candida* infection pathogenesis (or virulence). For example, phagocytes are generally thought to be the most effective cellular immune effector for controlling and clearing *Candida* infections. Neutrophils play a critical role in innate host defense against both *Candida* mucosal and invasive infections.^{4,5} Haematopoietic growth factors (specifically granulocyte colony-stimulating factors, G-CSF, and granulocyte-macrophage colony stimulating factor, GM-CSF) are potent drivers of phagocyte differentiation, recruitment and activation. In preclinical models, G-CSF therapy protects against lethal fungal infections⁶ and augments antifungal drug activity.⁷ In clinical studies, prophylactic use of GM-CSF resulted in reduced incidence of fatal infections in acute myeloid leukemia patients⁸ and improved survival from fungal

infections in stem cell transplant patients.⁹ It is also well-established that cytokines play a key role in epithelial immunity against *Candida* infection. For instance, patients with chronic mucocutaneous candidiasis have severely defective Th17 responses.¹⁰ When further exploring the Th17 response (specifically the dectin-1/CARD9/Th17 pathway), it was noted that patients with defective c-type lectin pattern recognition receptor dectin-1¹¹ and/or downstream adaptor CARD9¹² suffer from mucocutaneous candidiasis. Interestingly, one study showed that treatment with G-CSF in patients with isolated chronic mucocutaneous candidiasis resulted in a complete clinical remission and complete restoration of IL-17 levels¹³ — although these results could not be reproduced by another group.¹⁴ Finally, there is growing appreciation for the role that mucosal epithelial cells play in the first line defense against *Candida* infections. In addition to its function as a physical barrier to fungal invasion, recent evidence has revealed the critical role that epithelial cells play in triggering innate immune responses. For example, when human gingival epithelial cells are exposed to *Candida spp.*, toll-like receptors are upregulated; antimicrobial peptides (specifically β -defensins) are induced; and fungal burden is decreased.^{15,16} In the gastrointestinal tract, commensal gut bacteria can induce antimicrobial peptides (namely LL-37/CRAMP via HIF-1a) to maintain *C. albicans* colonization resistance in mice, and pharmacologic activation of these gut immune effectors not only reduces *C. albicans* colonization but significantly decreases dissemination.¹⁷ Thus, elucidating host immune effectors critical for protecting against *C. albicans* colonization or invasion not only provides critical insight into infectious disease pathogenesis but may provide the foundation for novel therapeutic strategies as well.

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In the study by Break et al.¹⁸ in this issue of *Virulence*, the role that dendritic cells play in antifungal (specifically anti *Candida albicans*) host defense is further explored. Dendritic cells (DCs) are immune cells found to be important against numerous pathogens and critical for both innate and adaptive immune responses. In terms of protection against *Candida spp.*, DCs can phagocytose and kill ingested *Candida* and produce cytokines in response to *Candida*. In this study, the authors explore whether a specific subclass of DCs (CD103⁺ DCs) are important for protection against systemic or mucosal *Candida* infection. The rationale for pursuing these studies is that CD103⁺ DCs have been found to be critical for parasitic and viral infections and furthermore CD103⁺ DC dependent IL-12 production has also been shown to be important for helminth infections. Prior to this study, the *in vivo* role of CD103⁺ DCs in innate immunity against *Candida* infections has been largely unexplored. The authors leverage the fact that the transcription factor *Batf3* is critical for the development of CD103⁺ CD11b⁻ DCs in order study mice (*Batf3*^{-/-} mice) that lack CD103⁺ DCs. Both a systemic (the commonly used *Candida albicans* tail vein injection model that emulates candidemia) and a mucosal (the oropharyngeal candidiasis model) *C. albicans* infection models are utilized. Given that *Candida* strain-specific differences in antifungal immune responses have been reported *in vivo*,¹⁹ the authors test both laboratory and disease model-appropriate clinical strains.

Interestingly, *Batf3* appears to be critical for accumulation of CD103⁺ DCs in the target organs of their respective disease models (kidney for systemic, tongue for mucosal) and for expansion during systemic and mucosal infections. Yet, deficiency of CD103⁺ DCs does not reduce survival or fungal clearance during either systemic or mucosal infections. Of note, a recent study showed that *Batf3* deficient mice did modulate the host response to a secondary *Candida* infection.²⁰ Thus, the role of CD103⁺ DCs in antifungal host defense during secondary or chronic *Candida* infections may be important and necessitates future study. This also elucidates the importance of using various infectious models to dissect the role of specific immune effectors.

The scientific focus of a large proportion of microbiology and infectious disease research centers almost solely on the microbial pathogen, yet most infections can only accurately be understood in the context of host-pathogen interactions and dynamics. Many “pathogens,” including *Candida spp.*, typically colonize the skin or mucosal surfaces of healthy hosts without any deleterious sequelae. If the pathogen invades into the adjacent tissues or spreads into the systemic bloodstream, the host’s primary (innate) defense systems have failed, usually as a result of

iatrogenic causes or in some cases genetic predisposition. In any case, understanding and unraveling the dynamics of host-pathogen interactions will be critical for devising novel therapies in the future. Thus studies like Break et al.¹⁸ that elegantly dissect the interaction between the microbial and host immune factors that lead to development of infection are essential for a more holistic appreciation and approach to correcting the often very dysfunctional host-pathogen interactions that have arisen as a consequence of great advances in medical therapies and technology.

Disclosure of potential conflicts of interest

No potential conflicts of interest were disclosed.

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