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A Murine Model for Disseminated Candidiasis in Neonates

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Abstract

Candida albicans is the leading fungal pathogen causing invasive disease in immunocompromised patients including the neonate. A reliable animal model for disseminated candidiasis in the neonate is needed to study the unique aspects of this host-pathogen interaction. To establish such a model, two day old BALB/c mouse pups were given intraperitoneal injections with varied inocula of *C. albicans* or saline control. Pups were examined every 3-8 hours for death. Surviving pups were sacrificed at 72 hours. Kidney, lung, spleen, liver and brain were homogenized and plated for colony counts and/or fixed for histological staining. Intraperitoneal injection of *C. albicans* led to mortality in a dose-dependent fashion. Disseminated infection was confirmed by colony counts of homogenized kidney, lung, and brain, as well as by histological examination. Infection with a *C. albicans* mutant lacking the cell surface adhesin, Als3p, led to significant reduction in mortality relative to wild-type ($P = 0.03$). This model will be useful to study the unique aspects of antifungal defense in a neonatal host and will provide a means to test novel therapeutic strategies.

Introduction

Candida albicans is the leading fungal pathogen in immunocompromised patients (1), and the third most common pathogen overall causing late-onset sepsis in premature infants (2). Well described risk factors for disseminated disease in this population include gastrointestinal (GI) colonization, prolonged hospitalization, broad spectrum antibiotic use, central venous catheters and parenteral nutrition (3-5). Colonization of preterm infants has been documented to occur through both vertical (from mother to infant) and horizontal routes (6). Even with antifungal therapy, candidiasis is often fatal among premature infants and is associated with neurodevelopmental impairment among survivors. Follow-up examinations at 18-22 months corrected age show significant increases in rates of cerebral

palsy, blindness, deafness, and mental retardation (7). The severity of these infections has led to development of prophylactic strategies to reduce colonization and limit invasive disease (8-11). Although effective, these strategies require prolonged exposure to antifungal agents with their associated risks (12-13). Novel therapeutic strategies are needed to improve these outcomes. However, the mechanisms leading to immune compromise in the neonate are likely different from other patient populations at risk (14). An animal model of disseminated candidiasis in a neonatal host is therefore needed to recapitulate unique aspects of this host-pathogen interaction.

Murine models have been a feasible and reliable method to study the pathogenesis of candidiasis, but have their limitations. Unlike humans, the mouse is not naturally colonized in the GI tract with *C. albicans*. In order to achieve persistent colonization, the animals must be treated with antibiotics and/or immunosuppressive agents, and dissemination is uncommon (15-16). Two strategies have been employed to circumvent these issues. The most widely used model involves intravenous injection of adult animals via the lateral tail vein. This model has been used extensively to study virulence properties of the organism and immunological adaptations of the animal in response to hematogenous infection (15,17). A second strategy is gastric inoculation of neonatal mice, which leads to persistent, albeit decreasing colonization over time with some dissemination and mortality. However, mortality was strain dependent and amounted to approximately 50% or less in these studies (18-19).

In the present study, we sought to develop a model that would result in more reliable disease burden while still maintaining some clinical relevance. The goal was to avoid pharmacological immunosuppression so as to allow inquiry into inherent immune status in infected neonates, as well as to obviate the variability and technical challenges inherent to GI colonization. GI pathology including abdominal surgery is an independent risk factor for disseminated disease (3,5). When the integrity of the bowel mucosa is compromised, translocation of the organism is likely facilitated with spread via the enteric or lymphatic circulation. Direct inoculation of the peritoneum can also occur in the setting of bowel perforation with similar routes to dissemination. This study was structured to test the hypothesis that disseminated disease and subsequent mortality could be induced in a reliable and reproducible fashion by the intraperitoneal (i.p.) route in neonatal mice without additional immunosuppression. This model provides the framework to study the unique host-pathogen interface in neonatal candidiasis as well as the development of novel therapeutic strategies.

Methods

Strains and Media

C. albicans strains used in this study include wild type strain SC5314 (20) and strain 1843 containing a homozygous deletion in *ALS3* (*iro1-ura3Δ :: λimm⁴³⁴/iro1-ura3Δ :: λimm⁴³⁴als3laΔ/als3saΔ-URA3*) (21) generously provided by Lois Hoyer. Starter cultures for injection were grown 16 h at 37°C with vigorous agitation in YEPD medium comprised of 1% yeast extract, 2% peptone and 2% dextrose (Difco Laboratories; Becton, Dickinson and Company; Franklin Lakes, NJ). Cultures were predominantly (>99%) yeast forms following this incubation. Prior to inoculation, overnight cultures of *C. albicans* were washed, enumerated on a hemacytometer, and resuspended in pyrogen-free saline (Hospira, Inc.; Lake Forest, IL). The concentration was adjusted such that the desired unit dose per gram could be delivered in a volume of 10 μl.

Injection of Neonatal Mice

Timed pregnant BALB/c mice were obtained from Charles River Laboratories (Wilmington, MA). Pregnant dams were maintained in individual cages with unlimited access to food and water. Mice were monitored to determine the date of parturition. Pups were delivered in litters ranging from 3-9 pups and were randomized prior to inoculation to either sterile saline (control) or 10^x colony forming units (CFU)/g of *C. albicans* yeast. Randomization was performed within cages rather than by litter to account for maternal and litter variations. Although cross-contamination of pups assigned to different experimental groups was theoretically possible, this risk was minimized by the short duration of the experiment. Further, in experiments using an endpoint such as mortality that is influenced by many variables, the risk of cross-contamination was outweighed by the risk of confounding effects of maternal and litter variability inherent to a litter-based randomization scheme. Pups were injected on post-partum day 2. Just prior to injection, each pup was weighed to the nearest tenth of a gram. Weight of pups was closely clustered around 2 g, so each pup received a standard dose of 20 μ l yeast or sterile saline injected i.p. in the lower half of the abdomen. After injection, pups were examined every 3-8 hours for death or signs of illness. The pups were dissected at the time of natural death or sacrificed for dissection when found moribund. All surviving animals were sacrificed at 72 hours after injection. A single kidney and lung were harvested and immediately homogenized for colony counts. If the time of natural death could not be accurately determined within 2 hours, the organ colony counts were excluded from analysis. The remaining kidney, lung, spleen, liver and brain were fixed in 10% buffered formalin (Fisher Scientific, Kalamazoo, MI) for subsequent histology. In selected studies, brain tissue was also collected for homogenization and colony counts.

Selected tissues underwent histological preparation and silver staining at the institutional core facility. Kidney, lung, and brain were homogenized with a FastPrep-24 Instrument (MP Biomedicals, Inc., Solon, OH) using Lysing Matrix D (Qbiogene, MP Biomedicals, Inc., Solon, OH) in 1 ml sterile saline and appropriate dilutions were plated on YEPD. Colony counts were performed following an overnight incubation at 37°C. All animal studies were reviewed and approved by the Lifespan Institutional Animal Care and Use Committee, which oversees the animal care facility where animals were housed for this study.

Results

C. albicans infection in neonatal mice

Mouse pups were injected i.p. on post-partum day 2 with concentrations of *C. albicans* strain SC5314 ranging from 10^4 to 10^8 CFU/g or with saline. Mice were followed closely for signs of illness and sacrificed at 72 hours following injection. Figure 1 shows the Kaplan-Meier survival curve summarizing these experiments. Doses of 10^6 CFU/g and below were not associated with mortality, and caused no apparent clinical symptoms. Doses above 10^6 CFU/g caused increased mortality in a dose dependent fashion. Because 10^7 CFU/g led to near complete mortality by study end point with a range of time of death throughout the observation period, this dose was selected for subsequent experiments.

Tissue sections of kidney, spleen, liver, and brain were silver stained to detect fungal elements (figure 2). Yeast and hyphal elements were scattered in the kidney and liver parenchyma with no specific anatomic relationship (figure 2, panel A and B). There was no fungus detected in brain by histology (not shown). In the low power view of the spleen, abundant fungal elements were diffusely present in the capsular region (figure 2, panel C). A high power view of the same region demonstrated prominent hyphae with penetration into the spleen parenchyma (figure 2, panel D).

Colony counts were obtained from homogenized kidney and lung and were highly variable (table 1). In general, colony counts were higher in kidney compared to lung at a given dose, and extent of fungal burden was proportional to dose injected. Statistical analysis using a negative binomial model supported this dose response relationship with $P = 0.0005$ for kidney and $P = 0.003$ for lung. Although no fungal elements were detected in histological studies of the brain, an additional experiment was conducted to examine brain involvement by colony counts, a more sensitive measure. Among 12 pups injected, all died by study end point and 9 pups (75%) had colony counts ranging from 100-240 colonies per brain. Although a lower fungal burden was seen relative to kidney and lung, these data support some involvement of the central nervous system. No organisms were recovered from any organs taken from animals in the saline control groups, providing reassurance in regard to the possibility of cross-contamination among animals in the same litter.

Assessment of ALS3 mutant in neonatal model

To determine the utility of this model in assessing virulence determinants of *C. albicans*, a mutant (1843) carrying a homozygous deletion of the adhesin gene, *ALS3*, was evaluated in the neonatal mouse model. Prior work demonstrated that this strain had reduced adhesion to epithelial and endothelial cells *in vitro* (22). The *als3* mutant strain yielded a statistically significant reduction in mortality relative to wild-type (figure 3, $P = 0.03$). The median survival for the wild-type was 24 hours compared to 44 hours for the mutant. Tissue burden in kidney and lung was also compared in these animals (table 2). Because colony count data were again highly disperse and not normally distributed, a negative binomial model was used for analysis. Again, tissue burden was higher in kidney than in lung for both strains. Although trends toward higher colony counts in mice injected with wild-type vs. mutant could be identified, there was no significant difference in tissue fungal burden in these animals.

Discussion

Invasive candidiasis portends a poor prognosis despite available antifungal agents. A sophisticated understanding of host-pathogen interactions will be required to make additional progress in treatment and prevention of these infections. The model described here will be useful in studies to explore the unique aspects of the neonatal host in this disease. Neonatal mouse models have been successfully used to study sepsis with other microorganisms including group B *Streptococcus* (GBS) (23), *E. coli* (24), *Pseudomonas aeruginosa* (25), and *Listeria monocytogenes* (26-27). In some cases these studies have uncovered significant differences that can be defined between the neonatal and adult host and demonstrate the importance of neonatal models to study invasive disease.

Previous studies of *C. albicans* in the neonatal mouse utilized gastric inoculation as the route of infection. Using neonatal mice, Pope *et al.* provided the first report of lethal candidiasis in an animal model following GI colonization without any additional measures to compromise immunity in the animals (18). In this study, five to six day old mouse pups were inoculated via intragastric injection (5×10^8 CFU) and systemic spread of infection was seen in selected organs. Fungal invasion was seen in liver, kidney, and spleen within six hours of inoculation, suggesting timely passage across the digestive tract wall or entry into the systemic circulation, possibly through the lymphatics. However, mortality was approximately 50% or less in this model and tissue burden decreased over time. A subsequent study using a lower dose (2×10^7 CFU) in 6 day old pups by the same route also led to recovery of *C. albicans* from kidney, liver, lung, and spleen, but in relatively smaller numbers and no mortality (19). A dose of 1×10^7 CFU led to still fewer or no recovery of fungi from these organs. However, long term GI colonization was demonstrated in these animals and early colonization led to protective immune responses after later IV challenge

with *C. albicans* as adults. In another study, six day old mouse pups were orally inoculated with *C. albicans* following cortisone induced immunosuppression. Histology of the entire gastrointestinal tract showed highest frequency of invasion of mucosa by *Candida* in the jejunum. Only GI tract organs were studied in this model (28).

GI colonization models using adult mice show important differences from the neonatal models. Broad spectrum antibiotic administration for 3 days prior to oral inoculation with *C. albicans* was necessary for GI colonization to develop (29). Additionally, once colonized, extraintestinal dissemination in these animals was very infrequent. When these animals were treated with dexamethasone, however, dissemination to kidney and mesenteric lymph nodes did occur, but mortality remained low (16). Features of disease associated with infection by the i.p. route in adult mice are also available. Vonk *et al.* studied the role of tumor necrosis factor- α (TNF) and lymphotoxin- α (LT) by injecting *C. albicans* i.p. in adult TNF-LT double knockout mice and their wild-type littermates (30). Unlike the present study in neonates, disseminated disease only occurred if immunosuppression was induced with cyclosporine prior to infection. Otherwise, the adult mice formed local abscesses that were cleared without disseminated disease. Taken together, these studies demonstrate that features of candidiasis in mouse models differ dramatically in the neonate compared to the adult, likely due to the relative immaturity of host defenses in the neonatal period. Such differences support the notion that features of disease unique to the neonatal host can be manifest by such an approach. However, there are additional factors placing preterm infants at risk for disseminated candidiasis that will be difficult to emulate in a murine model. Interventions such as indwelling catheters, parenteral nutrition, lack of enteral nutrition/breast milk, and many others that increase risk in the NICU are difficult to model and somewhat limit the applicability of host defense studies at this stage of mouse development to that of the preterm human.

In this study, the heavy involvement of the spleen with scattered foci in other organs supports a hematogenous route of dissemination, perhaps initiating in the spleen. Although involvement of the spleen was detected in the neonatal gastric inoculation model, colonization was similar or less than other organs (18). Presumably, the i.p. route of infection is responsible for the heavy spleen involvement in our model, either by direct contact with the organ or through lymphatic channels. Brieland *et al.* inoculated *C. albicans* (5×10^6 CFU/mouse) via lateral tail vein injections into adult, immunocompetent mice and reported growth of *C. albicans* in various organs (31). The kidney was noted to have logarithmic growth in fungal burden. However, the liver and heart fungal burden declined quickly over time. The brain, lung, and spleen were all noted to have steady fungal loads with no significant change over the duration of the infection. Consistent with these data, the kidney counts in this study were higher than in lung tissue. The Brieland study collected data over a 21-day post-infection time course. In our model, mortality occurred within 72 hours, and any surviving pups had generally cleared the infection by the 72 hour time point. The kinetics of infection were therefore quite different from the Brieland model.

The Brieland study described multiple foci of hyphal invasion in kidneys, hearts, brains and spleens of infected mice, with the largest fungal burden in the kidney. We found the largest foci of hyphae around the splenic capsule. We also did not visualize hyphal elements in brain of neonatal mice by histology, while the adult model showed brain involvement within 48 hours post-infection. Brain colony counts yielded consistent but reduced fungal burdens when compared to lung and kidney, suggesting that the fungal burden of the brain is not high enough to be detected by histology. Because tissue homogenates were the only way to assess fungal burden, involvement of vascular structures in the brain rather than the parenchyma itself is also possible. The differences in tissue distribution between these

models likely relate to the route of infection and/or the dose of inoculation, but may also be influenced by developmental stage of the animal.

We have previously described a single-chain variable fragment, scFv3, which is specific to Als3p (22). Als3p is a cell wall protein expressed on *C. albicans* hyphae, which belongs to the Als family of adhesins (32). Als3p enables adherence to both epithelial and endothelial host cells through interaction with E-cadherin and N-cadherin respectively (33). Strain 1843, carrying a homozygous deletion of *ALS3* demonstrates reduced adhesion to human epithelial and endothelial cells. Treatment of wild-type *C. albicans* with scFv3 resulted in reduced adherence, similar to the *als3* mutant (22). In our model, the *als3* deletion mutant showed somewhat attenuated mortality. Antibodies against Als3p such as scFv3 may therefore be useful to confer protection from disseminated candidiasis. This model provides fertile grounds to test this and other therapeutic strategies. Experiments are underway to evaluate scFv3 and other Als3p specific antibodies for their capacity to provide protection. Novel chemotherapeutic agents against fungi could also be tested in this model for efficacy and to assess any unique toxicity that may arise in a neonatal setting. Studies of pathogenesis with *C. albicans* frequently find differences among strains. This model can be used to extend these observations and make comparisons among isolates that are presumed to be different in pathogenic potential. Additionally, as non-*albicans* species increase in prevalence in the neonatal intensive care unit, this model will have utility to compare the pathogenic features of the different *Candida* species and potentially tailor appropriate therapies.

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Abbreviations

CFU	colony forming units
GI	gastrointestinal
scFv3	single-chain variable fragment 3

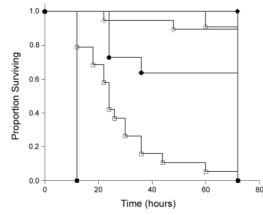
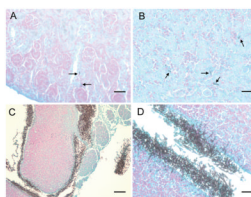


Figure 1. Kaplan-Meier survival curve by dose injected

Two day old mouse pups were injected i.p. with the following doses of wild-type *C. albicans* and survival curves were plotted: ■: 1×10^8 CFU/g (n=3); □: 1×10^7 CFU/g (n=19); ●: 5×10^6 CFU/g (n=11); ○: 1×10^6 CFU/g (n=11); ◆: 1×10^4 CFU/g (n=3); ◇: saline (n=19). Mortality occurred in a dose-dependent fashion at doses higher than 10^6 CFU/g.

**Figure 2. Tissue histology**

Silver stain of representative sections from kidney (A), liver (B), and spleen ((C) 10× magnification, (D) 40× magnification) are depicted from an animal injected with 10^8 CFU/g wild-type *C. albicans*. Arrows indicate hyphal elements visible within the organ parenchyma. Heavy involvement of the capsular and subcapsular regions of the spleen was seen. Panel A, B, D: bar = 25 microns; Panel C: bar = 100 microns.

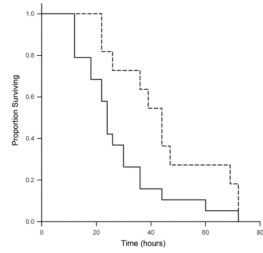


Figure 3. Kaplan-Meier survival curve comparing wild-type *C. albicans* and *als3* deletion mutant Two day old mouse pups were injected i.p. with 10^7 CFU/g wild-type *C. albicans* (solid line, n=19) or *als3* deletion mutant (dashed line, n=11) and survival curves were plotted. Median survival was 24 hours and 44 hours, respectively ($P = 0.03$ by log-rank test).

Table 1

Tissue burden by dose of wild-type *C. albicans* injected

Dose (CFU/g)	Kidney (CFU/organ)					Lung (CFU/organ)				
	Mean	Median	Minimum	Maximum	n	Mean	Median	Minimum	Maximum	n
10 ⁷	10,442	600	0	71,300	13	1,446	500	0	6,100	13
5×10 ⁶	8,734	70	0	66,000	8	1,055	30	0	7,800	8
10 ⁶	806	20	0	8,000	11	20	25	0	40	11
10 ⁴	13	20	0	20	3	0	0	0	0	3

Table 2
Tissue burden in mice infected with wild-type vs. mutant (*als3-/-*) *C. albicans*

	WT (CFU/organ) n=13				<i>als3-/-</i> (CFU/organ) n=10			
	Mean	Median	Minimum	Maximum	Mean	Median	Minimum	Maximum
Kidney*	10,442	600	0	71,300	3950	1100	0	13,500
Lung**	1446	500	0	6100	518	350	0	2200

* $P = 0.29$ for wild-type vs. mutant.

** $P = 0.14$ for wild-type vs. mutant.