

Published in final edited form as:

*Microbes Infect.* 2009 ; 11(8-9): 753–761. doi:10.1016/j.micinf.2009.04.018.

## Biofilm formation in clinical *Candida* isolates and its association with virulence

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### Abstract

Biofilm formation, an important virulence trait of *Candida* species was measured in 107 *Candida* isolates from 32 candidemic patients by XTT [2,3-bis (2-methoxy-4-nitro-5-sulfo-phenyl)-2H-tetrazolium-5-carboxanilide] activity and compared to biofilm formation of *Candida* isolates from oropharyngeal lesions of 19 AIDS patients. Biofilm formation by XTT varied among species and *C. albicans*; *C. lusitaniae* and *C. krusei* produced more biofilm than the other *Candida* species. *C. tropicalis* was the most dominant species isolated from blood followed by *C. albicans*, and other non-*albicans* species whereas only *C. albicans* was recovered from oral lesions. Importantly, though Biofilm formation was variable within a species it was stable in sequential isolates during chronic infection. Sequential isolates exhibited identical Karyotype pattern or RAPD patterns unless patients were co-infected with more than one strain. High biofilm formation was associated with slow growth rate but not with adherence. Murine infection studies demonstrated that in mice, degree of biofilm formation was associated with enhanced virulence as mice infected both with no and low biofilm formers survived longer than mice infected with high biofilm formers ( $p \leq 0.001$ ). We conclude that biofilm formation is a stable but strain specific characteristic that can greatly vary among *C. albicans* and non-*albicans* strains, and plays an important role in persistence of infection.

### Keywords

*Candida*; Biofilm; PFGE; XTT

## 1. Introduction

Several *Candida* species are commensals and colonize skin and mucosal surfaces of humans. Critically ill patients and otherwise immunocompromised patients are more prone to develop both superficial and life threatening *Candida* infections [1]. *Candida* infections also constitute the most common fungal infection in AIDS patients (HIV) [2]. These patients predominantly develop oropharyngeal candidiasis (OPC), which can lead to malnutrition and interfere with absorption of medication. In countries like India, where HAART is not universally available, OPC is still common in HIV infected individuals. Although *C. albicans* remains the most commonly isolated yeast in the US, non-*albicans* species are emerging and other countries report higher numbers among their patient specimens [3]. In that respect a recent report from

*Candida* blood stream isolates at All India Institute of Medical Science (AIIMS), one of the major tertiary care centers in India indicated that the proportion of non-*albicans Candida* was higher than reported in the US and Europe [4].

*Candida* infections are commonly associated with biofilm formation (BF) that can occur both on mucosal surfaces and on plastic surfaces of indwelling devices. These biofilm consist of matrix-enclosed micro-colonies of yeast, hyphae and pseudohyphae, arranged in a complex structure [5,6]. Biofilm are inherently resistant to antifungal agents including amphotericin B (AMB) and fluconazole (FLU) [6-9], and thus affected devices generally need to be removed [10]. Among clinical *Candida* strains BF can vary and depends on the *Candida* species [11]. However, the question if variability in BF is an independent virulence factor that affects outcome of infection is still not fully answered. The objective of this study was to investigate BF in clinical *Candida* isolates from blood and oral lesions from patients at AIIMS. In addition, BF of individual strains during chronic disease was studied. Last, the contribution of BF to virulence was examined in murine infection models.

## 2. Materials and Methods

### 2.1. *Candida* Isolates and study population

A total of 126 *Candida* isolates were obtained at various days post diagnosis from blood of 32 candidemic patients, and mucosal swabs of 19 HIV patients with OPC at AIIMS (Table 1). The identification of *Candida* species was done by germ tube test, chlamyospore formation on cornmeal agar and sugar assimilation test, API20C (bioMerieux, Marcy Etoile, France). Most of the candidemic patients were admitted to the intensive care unit (ICU). All HIV patients were clinic patients. Data on patients were collected according to the AIIMS internal review board regulations. MIC to FLU and AMB were determined for planktonic cells according to CLSI M27A guidelines and their biofilm grown counterparts as previously published protocol [12].

### 2.2. Determination of Biofilm production

A semi- quantitative measure of BF and viability was detected by XTT [2,3-bis (2-methoxy-4nitro-5-sulfo-phenyl)-2H-tetra-zolium-5-carboxanilide] reduction assay in 107 isolates. In addition biofilm mass was directly measured by the crystal violet assay in 72 isolates as described [13,14]. *C. albicans* strain SC5314 was used as a control strain in each experiment. *C. albicans* BF was compared on three different surfaces, polystyrene, polypropylene (Greiner Bio One) and silicon elastomers (Cardiovascular Instrument Corp) as described above [13]. Briefly, 96 well plates were used for biofilm formation on polystyrene and polypropylene. To observe biofilm on silicon, silicon elastomers were cut of equal size and used to grow biofilm in 96 well plates. Growth conditions for BF on all three surfaces were the same. "High" and "low" biofilm formers (HBF and LBF) were defined as isolates that exhibited XTT activity greater than or less than the geometrical mean (OD of 0.50). No biofilm formers (NBF) are with optical density of < 0.1.

### 2.3. Genotype analysis by RAPD and PFGE

For RAPD DNA was isolated as described previously [15], and RAPD was done using M13 primer (5' GAGGGTGGCGTTCT 3'). For PFGE, chromosomal DNA plugs were prepared using  $5 \times 10^8$  *Candida* cells as described [16]. Chromosomal separation was done on 1% pulse-field-certified agarose gel (SeaKem) in a CHEF DRIII pulse-field electrophoresis system (Bio-Rad) in 0.5 X Tris-boric acid-EDTA at 14 °C. The electrophoresis conditions were a switch-time of (i) 120 s for 24 h (ii) 240s for 36 hr, at 4.5V and an angle of 112° each. *Saccharomyces cerevisiae* chromosomal DNA was used as size marker. DNA gel was stained with ethidium bromide and photographed after destaining for 15 min.

## 2.4. Phenotypic characterization of yeast isolates

Growth was measured by absorbance at 490 nm at different time interval after growth in RPMI (pH 7) at 150 rpm, and 37 °C. For adhesion assays  $10^3$  *Candida* cells were added into 96-well polystyrene plates (corning) and incubated in RPMI at 37 °C for 2 hours. Adhesion was determined by cell counts using a Photo Zoom inverted light microscope (Cambridge Instrument, MA). The number of *Candida* cells attached to the bottom of each well was averaged per 40-power field. The assay was done in triplicates.

## 2.5. Mice study

In vivo pathogenesis of HBF, LBF and NBF *C. albicans* isolates was studied in female BALB/c mice (6-12 weeks, body weight 18-20 gm) obtained from NCI (Bethesda, MD). Ten mice per group were injected intra-venous with HBF, LBF and NBF *C. albicans* (grown for 12h in SDA at 30°C) strains with similar doubling time with  $3 \times 10^6$  cells in 100 µl sterile PBS via tail vein [17]. Dilutions of the inoculums (all blastophores) were back plated onto SDA plates to assure that comparable numbers of yeast cells were injected. Fungal burden was obtained d-3 in lung, liver and kidney. For survival 5 mice were observed daily for signs of disease till death. Moribund mice were sacrificed according to standard procedures. Each experiment was repeated once. At death organs were immediately excised and processed for organ fungal burden and histopathology. Formalin fixed organs were stained with periodic acid Schiff stain.

## 2.6. Statistical analysis

The statistical tests used were  $X^2$  and student's t test. Survival differences were analyzed using Kaplan-Meier survival curve, SPSS version 8.0 (SPSS Inc., Chicago IL). P value < 0.05 was considered significant.

# 3. Results

## 3.1. Patient characteristics

BF was determined in 126 *Candida* isolates from 51 patients. Nineteen isolates were cultured from the mucosal lesion of 19 HIV patients with OPC and 107 isolates were grown from the blood of 32 in-patients. Of the 32 patients, 28 were admitted to an ICU, 26 were on mechanical ventilator and exhibited risk factors for BSI including 53 % had indwelling central venous catheters. The mortality was 68 % (Table 1 and 2). Associated risk factors and outcome of the patients did not correlate with extent of BF. Eleven (34%) candidemic patients were on antifungal prophylaxis. In 14 (44%) patients therapy was started after infection. In 3 of these patients infection was cleared and in 11, infection persisted. Thus the majority of patients exhibited no or partial response to anti-fungal therapy (AFT). AFT varied and included both single and combined regimens. The HIV patients with OPC were treated as out patients and thus less sick. The majority of HIV patients exhibited low CD4+ counts but were not on HAART.

## 3.2. Species distribution and antifungal susceptibility and molecular typing

Various *Candida* species were identified with *C. tropicalis* (Ct, 38%) being the most common species followed by *C. albicans* (Ca, 21%), *C. parapsilosis* (Cp, 13%), *C. glabrata* (Cg, 3%), *C. krusei* (Ck, 3%), *C. pelliculosa* (Cpe, 3%), *C. lusitaniae* (Clu, 3%), mixed infection (MI, 13%), unidentified (UI, 3%). All isolates from oral lesions of HIV, OPC patients were *C. albicans* except in two patients co-infection of *C. albicans* with *C. tropicalis* and *C. krusei* were documented. For HIV patients all the *C. albicans* except one were sensitive (MIC 0.125-4 µgml<sup>-1</sup>) and one *C. krusei* isolate was resistant (MIC >64 µgml<sup>-1</sup>) to FLU. All 107 blood isolates were susceptible to AMB (MIC <1 µgml<sup>-1</sup>), and most to FLU (MIC 0.25-8 µgml<sup>-1</sup>). Eleven isolates exhibited dose dependence (MIC, 16-32 µgml<sup>-1</sup>) and included 4 *C.*

*pelliculosa*, 1 *C. parapsilosis* and 3 each of *C. tropicalis* and *C. glabrata*. As expected all *C. krusei* isolates were resistant to FLU. The majority of *Candida* isolates grown as biofilm were resistant to antifungals. Only 17% exhibited dose dependent sensitivity to FLU and 33% were sensitive to AMB.

### 3.3. Degree of BF in patients was dependent on the *Candida* species

BF in blood *Candida* isolates differed considerably by XTT method and was dependent on species (Fig. 1a). BF was highest in *C. albicans* and comparable to *C. lusitaniae* and *C. krusei* followed by *C. tropicalis*, *C. parapsilosis*, one unidentified strain, *C. glabrata* and *C. pelliculosa*. *C. albicans* was the highest biofilm producer, also by crystal violet analysis and results for the two methods were significantly correlated (correlation coefficient,  $r=0.72$ ,  $p=0.02$ ), *C. glabrata* was a low or no biofilm producer by both methods. BF was highly variable among *C. albicans* strains isolated both from blood and mucosa (Fig. 1b) but did not differ significantly ( $P=0.11$ ). BF was also variable in *C. parapsilosis* but comparable among *C. tropicalis* (Fig. 1c). However for *C. tropicalis* strains crystal violet analysis exhibited a more variable BF than the XTT method. Four patients were infected with more than one *Candida* species. In two patients two different *Candida* species were grown from a single blood sample and in both scenarios *C. albicans* was the species with higher BF (data not shown).

### 3.4. BF in sequential *Candida* isolates from same patients

A total of 107 sequential blood isolates were available from 32 patients. Duration between the first and last isolation ranged from 0 to 85 days (Table 1). BF remained stable between most sequential isolates with mean % variability of 11 (range 0.2 to 30%), even in isolates that were recovered 85 days apart (Fig. 2a). In contrast, BF in sequential isolates was more variable (mean variability 56%) if the patient was co-infected with 2 species (P15, P16, P22, P23) or infection with different strains (P20). No significant change in BF in sequential isolates after the initiation of AFT was documented. Hence we conclude BF is stable during chronic infection and both HBF and LBF can co-exist in the same host.

### 3.5 Molecular characterization

Genotype characterization of biofilm forming *Candida* strains by PFGE and/or RAPD demonstrated that the same *Candida* strain persists during chronic infection in most cases. Only sequential *C. tropicalis* strains gave variable RAPD patterns and by PFGE did not exhibit enough chromosome bands. Sequential isolates with stable biofilm exhibited identical genotype unless co-infection with two different strain of same species (*C. parapsilosis*, P20 I & II). Sequential *C. parapsilosis* isolates with genotype changes exhibited significant differences in their BF, RAPD and PFGE typing, which confirmed that they were distinct strains (Fig. 2b and d(iii)). For *C. glabrata*, genotypic variability was documented in sequential isolates, all of which produced minimal or no biofilm (Fig. 2b). *C. albicans* isolates from different patients with identical genotype (RAPD and PFGE) exhibited variable BF. In HIV patients with OPC, 7 different karyotypes were identified in 17 *C. albicans* isolates (Fig. 2c). The majority of OPC isolates exhibited karyotype III ( $n=7$ , 41%), which was also exhibited by all *C. albicans* isolates from candidemic patients, however RAPD analysis differentiated these strains but showed identical RAPD in sequential isolates. Hence, for most *Candida* species stability in BF among serial isolates was associated with stability in their Karyotype pattern.

### 3.6. Association of BF with phenotypic characteristics of *Candida*

We compared adherence of HBF with LBF *Candida* strains as well as adherence among different *Candida* species. No statistical difference in adherence of blastospore after 2 hr was found between HBF vs LBF or between different *Candida* species. Most *Candida* strains

exhibited comparable growth rates irrespective of species except *C. krusei*, which all grew slower and some *C. albicans* strains which were also slow growers (doubling time > 5 hr). Among *C. albicans* a trend of higher BF was observed in slow growers ( $0.83 \pm 0.24$  vs  $0.69 \pm 0.10$ ,  $p=0.08$ ) when compared to fast growers. No LBF *Candida* isolate exhibited a slow growth rate (Fig. 3). Hyphae formation after prolonged culture in RPMI was more extensive in HBF in comparison to LBF. NBF have no hyphae formation and have the least adherence to the surface ( $OD < 0.1$ ). In addition, we observed that “slow growers” and HBF strains after 24hrs at 37°C produced more filaments in comparison to fast growers, but there was no association with adherence of their blastospores, which was more pronounced in LBF strains. We also compared BF on three different plastic surfaces and determined that BF was the most pronounced on polypropylene followed by polystyrene and silicon (data not shown).

### 3.7. Correlation of BF with virulence

A retrospective chart review determined that in this patient cohort no patient characteristic or potential risk factor was associated with BF or mortality however mortality was too high to draw conclusion. Hence, we examined the virulence of *C. albicans* strains that differ in BF in a murine infection model. For these studies we selected two *C. albicans* strains each for HBF, LBF and NBF with similar growth rate and compared their virulence in an intra-venous murine infection model. We selected strains which were identified as HBF, NBF and LBF on all three plastic surfaces. In mice infected i.v. the median survival time differed significantly ( $p < 0.001$  by log rank) and correlated with BF (median time of survival was 1 and 4 days vs 10 and 10.8 days vs 20 and > 40 days,  $p < 0.001$ ) for 2 HBF, 2 LBF and 2 NBF strains respectively (Fig. 3a). Fungal burden in kidney, liver and lung at the time of death of HBF infected mice was detected in all three organs ( $\log 5.84 \pm 0.10$ ,  $\log 3.87 \pm 0.16$  and  $\log 2.78 \pm 0.36$ , respectively), whereas LBF infected mice at the time of death had  $\log 6 \pm 0.66$  CFU in the kidney but no yeast was detectable in liver and lung. NBF infected mice had no detectable CFU at day 7. Histopathological data demonstrated massive accumulation of yeast and hyphal elements in cortex, medulla and the papilla associated with degraded glomerulus and renal tubules in HBF infected mice. In contrast, kidney of LBF infected mice were showing infiltration of fungal element and leukocytes mainly in the collecting ducts of the papilla, and only few in cortex and medullary region with small inflammatory foci. Renal clearance could be observed in NBF infected mice and the kidney appeared healthy (Fig. 3b). This result strongly supports the conclusion that variability in *C. albicans* BF plays an important role in host pathogen interaction and their outcome.

## 4. Discussion

BF is presumed to promote persistence of infection and thus we investigated BF in serial *Candida* isolates of patients with candidemia and of isolates of patients with OPC in a major medical center in India. Our data suggests that (i) BF depends on species and in *C. albicans* and *C. parapsilosis* species is highly variable but less variable in *C. tropicalis* (ii) Degree of BF is a specific trait of a *Candida* isolate that is not associated with differences in adherence but with slow growth. (iii) Most importantly this trait is inherited and thus remains stable in sequential isolates. (iv) No association between individual host factors and the capacity for BF in the *Candida* spp. was observed and both HBF and LBF can co-exist in the same host. (v) In animal murine models degree of BF correlates with virulence, whereas gene expression of biofilm associated genes is more variable.

*Candida* biofilms may contribute both to the pathogenesis of superficial and systemic candidiasis as they are notoriously resistant to antifungal drugs. *Candida* species are now the fourth most common cause of nosocomial BSIs [18]. Although *C. albicans* remains the most common species recovered from blood in Europe and the US [19] emergence of non-albicans



species has been documented [20]. In this and in earlier studies *C. tropicalis* was the predominant species in BSI in India [4].

We investigated BF in *Candida* isolates of patients with candidemia and OPC. Patients in this study had many predisposing factors for disseminated candidiasis including longer stay in ICU and indwelling medical devices. Patients with OPC were all HIV clinic patients that were infected with *C. albicans*. The strength of our study is that it is the first to investigate BF in blood serial isolates derived from the same patient, which yields important information on the stability of this virulence-associated trait. Our data demonstrates that BF is dependent on *Candida* species and generally highest in *C. albicans* isolates followed by *C. lusitaniae* and *C. krusei*. Most studies have found similar differences of BF among the *Candida* species [11, 21]. Of note is that BF formation in *C. glabrata* isolates in this study was found to be among the lowest similar to the result of other studies [22], whereas BF in *C. glabrata* isolates from urine was found to be much higher [13]. This could potentially suggest that certain host niches in some *Candida* species promote selection of HBFs.

BF among *C. albicans* varied significantly both by the XTT and crystal violet method and ranged from high to low both in blood and OPC isolates similar to previous studies in urine [13] and blood isolates [23]. One study reported comparable BF in 26 *Candida* strains derived from OPC [24] and another study demonstrated that *Candida* strains from BSI produce more biofilm than those from noninvasive infections [21] neither of which was observed in our study. Variability of BF was also found in *C. parapsilosis* but only by crystal violet in *C. tropicalis* isolates. The meaning of these findings is unclear and warrants further investigation. We found under direct microscopy that *C. albicans* strains exhibit differences in ratio of blastospore, hypha and pseudohyphae, which may explain BF variability, rapid death and pronounced tissue invasion. Although the growth rate was similar between isolates we noted that all slow growers were HBF even though some HBF exhibited a normal growth rate. This is consistent with other reports showing that biofilm-associated cells that are embedded in the deeper layers, show suppressed growth rates, thereby being more resistant to antifungal drugs [25].

Because of extensive co-morbidity candidemia was often persistent despite appropriate AFT and thus several serial isolates could be isolated. As previously reported most biofilms exhibited enhanced resistance to AFT. BF was stable in serial isolates from individual patients even after the initiation of AFT. Identical genotypes were demonstrated in most serial isolates except *C. glabrata*, *C. tropicalis* isolates and in one patient with *C. parapsilosis* infection, which supports conclusions of other studies that the same strain usually persists during chronic infection [26]. We demonstrate that significant differences in BF of two serial isolates are the result of co-infection with more than one species. This finding also demonstrates that both LBF and HBF can co-exist in the same niche and we hypothesize for *C. glabrata* the host may select for HBF in certain niches (urine and vagina) [13,27] and LBF in blood [28].

Candidemia ranks among the infections with highest mortality rates [1,29]. The impact of BF in *Candida* strains on mortality of candidemic patients concluded based on retrospective multivariate analysis that high BF is associated with poor outcome [11]. This conclusion has not yet been verified in a prospective study. Outcome of candidemia is affected by many factors and the mortality in our patient population was too high to draw such conclusions. However in a murine infection model we demonstrated that BF affects outcome. Infection with 6 different *C. albicans* HBF, LBF or NBF strains with comparable growth rates demonstrated that decreased survival correlated with degree of BF. Histological analysis suggested that despite comparable fungal burden in the kidney, HBF results in deeper tissue invasion as many fungal cells invade the glomerular capsule, leading to degraded glomeruli and also renal tubules in HBF infected mice. Of note is that is conceivable that the rapid mortality could be the result of more pronounced hyphae formation in HBF strains, which may promote BF and possibly

shock. In contrast, LBF infected mice presented with a less invasive infection that was predominantly limited to the papilla and did not result in tissue destruction.

In summary, we conclude that BF represents a phenotypic characteristic of a *Candida* strain that is inherited and thus stable during chronic infection. BF is more variable in some species especially *C. albicans* and *C. parapsilosis*. Our data from murine infection models greatly supports the notion that enhanced BF could potentially be an independent risk factor for prolonged disease and bad outcome. Prospective patient studies may be warranted in addition to studies that investigate the regulatory pathways that control BF and hyphae transition to identify more virulent *C. albicans* strains and optimize treatment regimens of disseminated candidiasis.

## Acknowledgement

We thank Dr. Bitzer for help with the histological analysis. This work was supported by grant AI059681-05 to B.C.F., pilot funds from CFAR grant (AI 051519) and by the AIDS International Training and Research Program (program director, Vinayaka Prasad [NID D43-TW001403]) to I.X. and F.H. Furthermore, F.H. was supported by ICMR (3/1/59/MPD/2003-JRF).

## Abbreviations

BF, biofilm; HBF, high biofilm former; NBF, no biofilm former; LBF, low biofilm former.

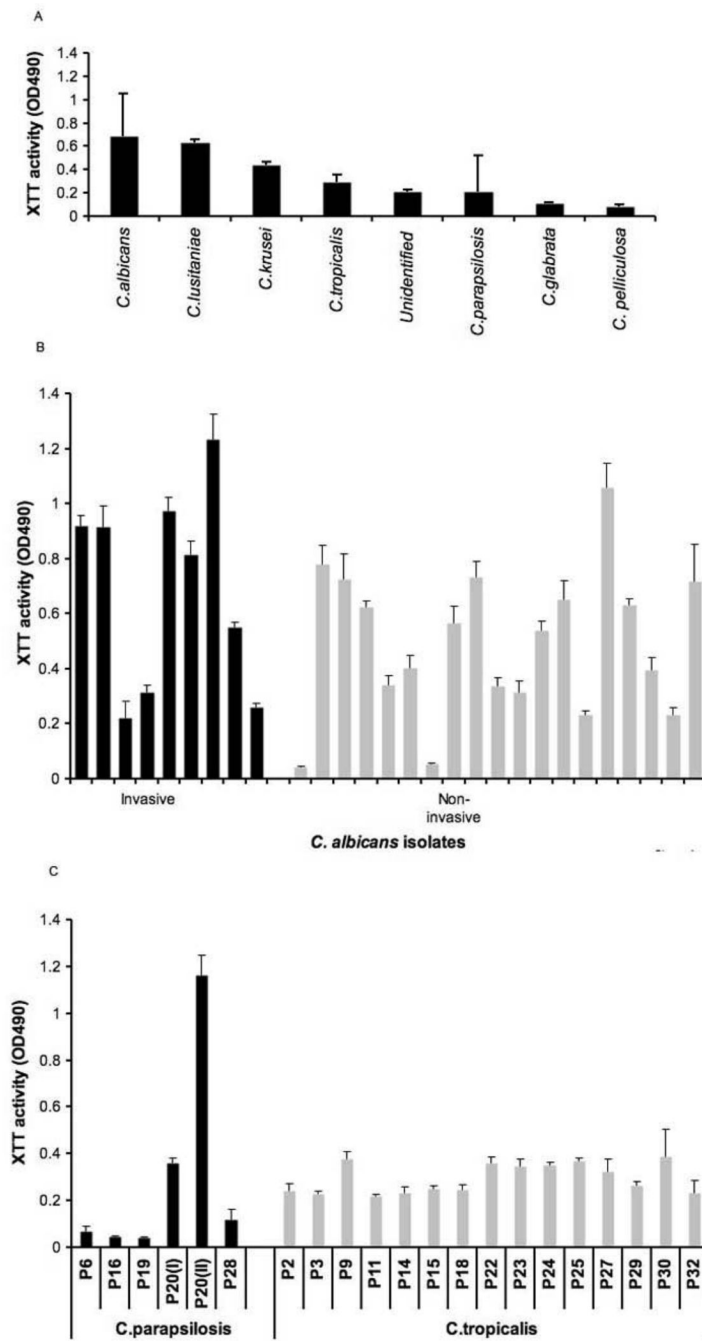
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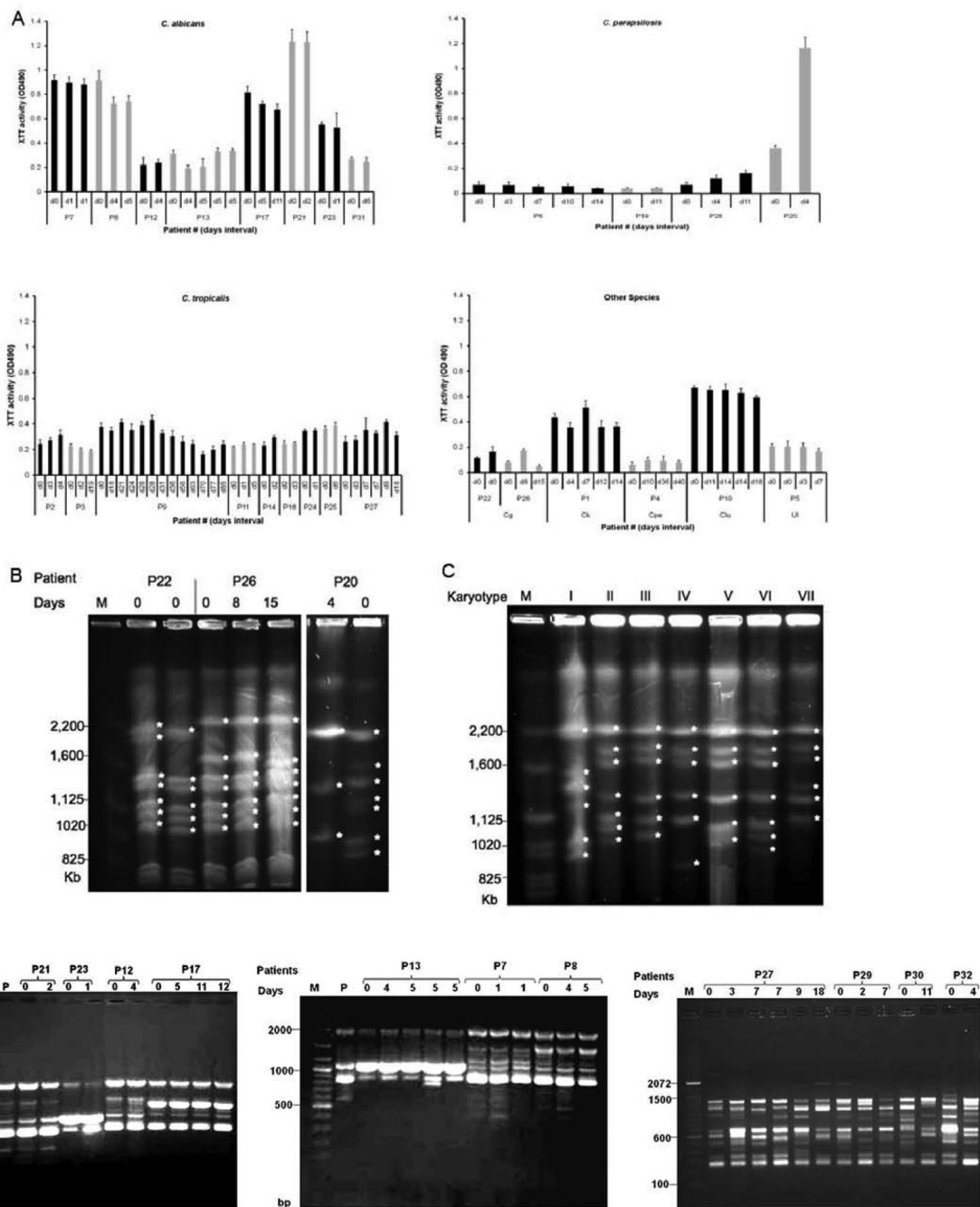
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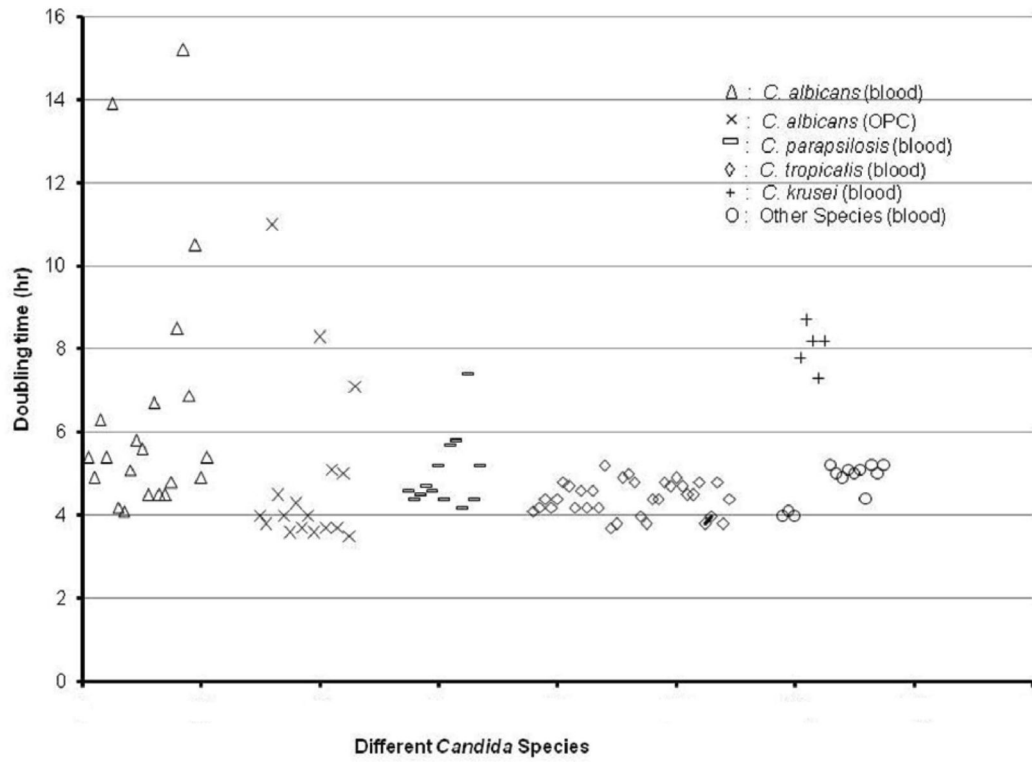
**Fig. 1.** BF is dependent on *Candida* species. (A) BF is highly variable among clinical *Candida* species (B) BF was highly variable among *C. albicans* isolates from both candidemic (invasive-black bar) and oropharyngeal lesion of HIV positive patients (non-invasive-gray bar) (C) Like *C. albicans*, BF was variable among *C. parapsilosis* but was relatively stable among *C. tropicalis* isolates from candidemic patients.



**Fig. 2.**  
 A) BF was comparable among serial isolates isolated up to 85 days apart (*C. tropicalis*-P9). In some cases co-infection with different strains of the same species (*C. parapsilosis*-P20) resulted in variation of BF whereas others exhibited comparable BF (*C. glabrata*-P22, P26). Black and gray bar indicates serial isolates from individual patients. (B) Differences in BF of serial *C. parapsilosis* were associated with karyotype variability and infection with two strains. Serial *C. glabrata* isolates exhibited in karyotypic variability but no differences in BF unless infection with different strains of same species (P20). (C) Seven different karyotype patterns were found among *C. albicans* OPC isolates from HIV patients. (D) RAPD profile of sequential isolates by M13 primer are shown for i) *C. albicans*, P21, P23, P12, P17, P13, P7, P8 and ii)

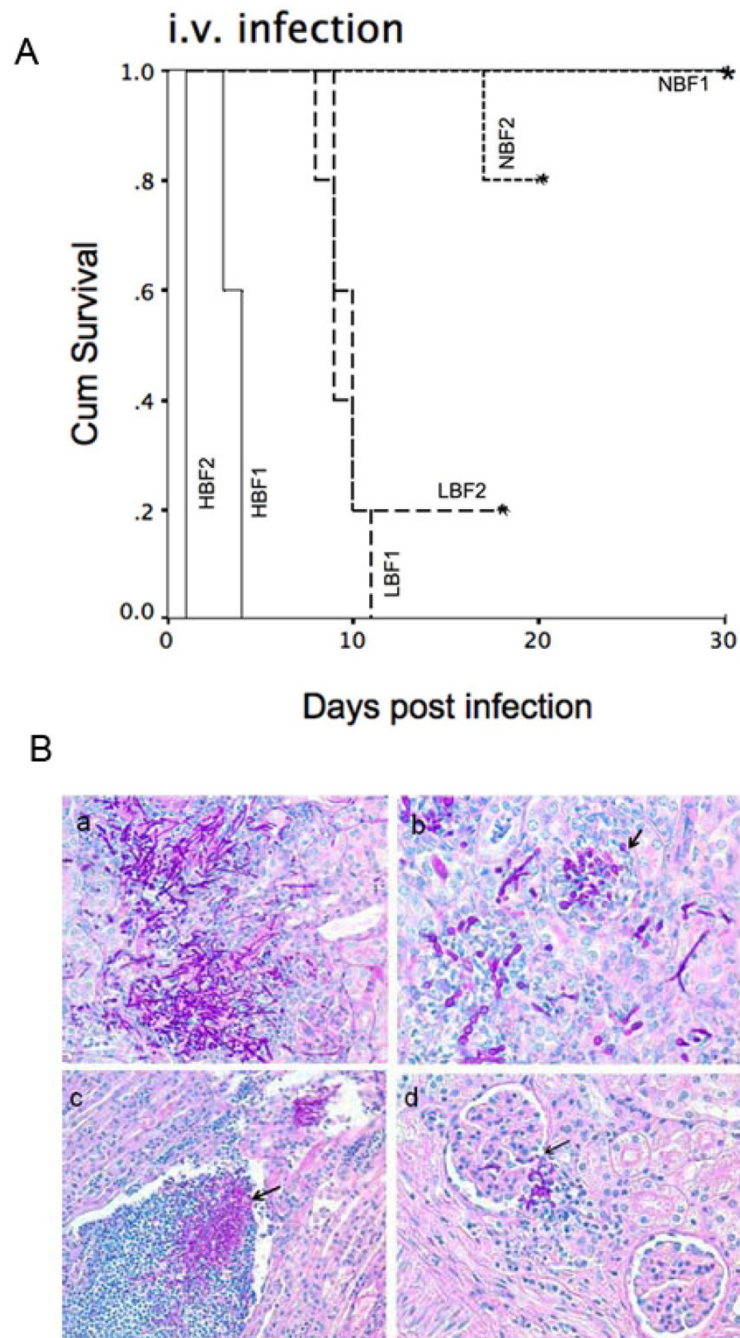
*C. tropicalis*, P27, P29, P30, P32. Note that this method demonstrated persistence of original strain for *C. albicans* isolates and more heterogeneity than karyotyping. For *C. tropicalis* RAPD patterns were very heterogeneous analogous to *C. glabrata*.

A.



**Fig. 3.** Variability in BF was associated with variability in growth. Doubling time for most *Candida* strains were comparable. Among *C. albicans* strains slow growers were found.





**Fig. 4.** (A) Survival of BALB/c mice (n=5 per group) after intravenous challenge with  $3 \times 10^6$  blastospore of high, low and no biofilm formers (HBF, LBF, NBF) *C. albicans* differs significantly ( $p < 0.001$ , log rank) (B) Histopathological examination of kidney. (a) HBF; massive accumulation of yeast cells, hyphal element and few leukocytes in calyx (b) HBF; kidney cortex showing infiltration of yeast and hyphal elements and degraded glomerulus. (c) LBF; papilla showing leukocytosis and fungal infiltration in collecting duct (d) LBF; little fungal infiltration in cortex and showing intact glomerulus. (Periodic acid Schiff stain was used).

**Table 1**  
Clinical summary and outcome for individual Candidemia patient

Pt #	Species	Duration of stay (AIIMS) (days)	Day of Ist (+) isolate	No. of isolates	Days recovered	Flu MIC #gml <sup>-1</sup>	AF start (day)	Diagnosis	Outcome
P1	Ck	19	1	5	0,4,7,12,14	64	8	AP & RF	D
P2	Ct	28	23	3	0,3,4	0.5-1	None	Ludwig's angina,	D
P3	Ct	180	141	3	0,2,19	32	65	ALL	S
P4	Cpe	45	4	4	0,10,36,40	16	FLU	Pancreas CA	D
P5	UI	21	10	4	0,0,3,7	4	15	Perforative peritonitis	D
P6	Cp	168	120	5	0,3,7,10,14	8	82	Cerebral atrophy	D
P7	Ca	45	41	3	0,1,1	0.5	42	Osteoarthritis	S
P8	Ca	40	31	3	0,4,5	1	31	Fungal sepsis, CHF	D
P9	Ct	203	28	13	0-85	2-8	69	ascending myelitis	S
P10	Clu	62	57	6	0-18	0.5	None	Fungal sepsis, DIC	D
P11	Ct	31	9	3	0,1,5	1	5	CHD, endocarditis	D
P12	Ca	37	6	2	0,4	0.25	6	AP, RF, sepsis	D
P13	Ca	50	16	4	0,4,5,5	0.25	8	HF, ascitis	S
P14	Ct	5	3	2	0,2	1	None	DSS	D
P15	Ca,Ct,Cp	22	5	3	0,0,12	0.5-1	VORI	ALL	S
P16	Cg,Cp	30	17	2	0,1	1	16	AP	D
P17	Ca	56	1	3	0,5,11	0.5	8	Fungal endocarditis	S
P18	Ct	47	15	2	0,3	0.5-1	13	Pelvic abcess, MTB	S
P19	Cp	60	19	2	0,11	8-16	1	RF,candidiasis,	D
P20	Cp	NA	NA	2	0,4	4-8	FLU	sepsis	S
P21	Ca	47	48	2	0,2	1	None	Gall bladder CA	D
P22	Ct,Cg	50	7	3	0,0,0	0.5-8	33	Polytoma	D
P23	Ca,Ct	35	17	3	0,1,5	0.5-2	17	Biliary peritonitis	D
P24	Ct	NA	NA	2	0,1	2	AMB	MTB	S
P25	Ct	53	22	2	0,6	0.5-1	30	Fungal endocarditis	D
P26	Cg	79	55	3	0,8,15	16	51	Cervix CA	D
P27	Ct	26	8	6	0,3,7,9,18	1	15	DSS	D
P28	Cp	70	41	3	0,4,11	0.5	26	Tetraplegia	D

Pt #	Species	Duration of stay (AIMS) (days)	Day of 1st (+) isolate	No. of isolates	Days recovered	Flu MIC $\mu\text{gml}^{-1}$	AF start (day)	Diagnosis	Outcome
P29	Ct	35	28	3	0,2,7	1	28	CHD	S
P30	Ct	NA	NA	2	0,11	0.5	NA	NA	-
P31	Ct	24	17	2	0,6	2	9	CHD, Pneumonia	D
P32	Ct	4	3	2	0,4	0.5	4	Multorgan failure, AP	D

Note: ICU -intensive care unit, AF- antifungal, AMB- amphotericin B, FLU- fluconazole, VORI- voriconazole, AP- acute pancreatitis, RF- renal failure, ALL- acute lymphoid leukemia, CA- cancer, CHF- chronic heart failure, DIC- disseminated intravascular coagulation, CHD- chronic heart disease, HF- hepatic failure, DSS- dengue shock syndrome, MTB- mycobacterium tuberculosis, NA- not available, S- survived, D- died

**Table 2**  
Underlying Candidemia and HIV patients associated variables

Variables	Candidemia Patients (N=32)	HIV patients (N=19)
Median age	± 51 (2m-85 yr)	± 34 (27-62 yr)
Male	17	15
Female	15	4
ICU admission	27 (84%)	NA
Ventilator	26 (81%)	NA
Intubated	15 (47%)	NA
Foleys catheter	27 (84%)	NA
Central venous catheter	17 (53%)	NA
Diabetes	3 (9%)	NA
Cancer patients	5 (16%)	NA
Prior surgery	13 (41%)	NA
Prophylactic antifungal	11 (34%)	1 (5%)
Organ transplant	1 (3%)	NA
Dialysis	3 (9%)	NA
Premature birth	1 (3%)	NA
Old age (>60 yr)	11(34%)	NA
Median CD4 count	NA	169 ( 14-774)
Antiretroviral therapy	NA	6 (32%)