



Published in final edited form as:

*J Med Microbiol.* 2006 October ; 55(Pt 10): 1323–1327. doi:10.1099/jmm.0.46737-0.

## ***Candida albicans* HWP1 gene expression and host antibody responses in colonization and disease**

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### **Abstract**

*In vivo* expression of the developmentally regulated *Candida albicans* hyphal wall protein 1 (*HWP1*) gene was analysed in human subjects who were culture positive for *C. albicans* and had oral symptoms ( $n=40$ ) or were asymptomatic ( $n=29$ ), or had vaginal symptoms ( $n=40$ ) or were asymptomatic ( $n=29$ ). *HWP1* mRNA was present regardless of symptoms, implicating hyphal and possibly pseudohyphal forms in mucosal carriage as well as disease. As expected, in control subjects without oral symptoms ( $n=10$ ) and without vaginal symptoms ( $n=10$ ) who were culture negative in oral and vaginal samples, *HWP1* mRNA was not detected. However, exposure to Hwp1 in healthy culture-negative controls, as well as in oral candidiasis and asymptomatic mucosal infections, was shown by the existence of local salivary and systemic adaptive antibody responses to Hwp1. The results are consistent with a role for Hwp1 in gastrointestinal colonization as well as in mucosal symptomatic and asymptomatic infections. Overall, Hwp1 and hyphal growth forms appear to be important factors in benign and invasive interactions of *C. albicans* with human hosts.

### **INTRODUCTION**

*Candida albicans* is a member of the normal flora of the gastrointestinal tract that frequently causes serious oral and vaginal mucosal invasion and systemic disease in hosts with impaired immune defences. For pathogens that persist in the host permanently or for extended periods, understanding the mechanisms that lead to progression from commensalism to virulence is an emerging area of medical research. It has recently become recognized that for pathogens whose ecological niche is the host, molecular factors believed to be important for virulence may also be considered adaptive factors that play an essential role in allowing the pathogen to persist in the host (Falkow, 2006). In these organisms, mechanisms of persistence and virulence may overlap via common determinants that function in both states. For *C. albicans*, the factors contributing to low-level asymptomatic persistence in the gastrointestinal tract, i.e. colonization, among inhibitory microbial flora, to frequent occurrence of asymptomatic oral and vaginal carriage, and to virulence in immunocompetent and immunodeficient hosts, are poorly understood. In previous work, we

have found that expression of specific secreted aspartyl proteinase (*SAP*) genes is correlated with active disease and anatomical location, whereas expression of other *SAP* genes reflects the presence of the organism but not disease (Naglik *et al.*, 2003).

Hyphal wall protein (Hwp1), a transglutaminase substrate which functions as an adhesin, is important for the pathogenesis of candidiasis (Staab *et al.*, 1996, 1999; Sundstrom *et al.*, 2002). *HWP1* mRNA and protein are abundant in hyphae *in vitro*, and greatly reduced or undetectable in yeast forms (Staab *et al.*, 1996, 1999; Sharkey *et al.*, 1999; Nantel *et al.*, 2002). Hypha-specific expression of Hwp1 extends to fungal growth in the host, as shown by the presence and absence of Hwp1 on hyphae and yeast, respectively, in candidiasis of the murine stomach (Sundstrom *et al.*, 2002). To gain insight into the relationship between *HWP1* gene expression and commensalism or symptomatic tissue invasion, we analysed the same carrier and candidiasis specimens that had been used previously to determine *SAP* and *PLB* gene expression in samples of whole unstimulated saliva and vaginal swabs (Naglik *et al.*, 2003), for *HWP1* gene expression by RT-PCR. Host antibody responses to Hwp1 were also measured. The results supported a potentially important role for hyphal forms in asymptomatic infections with *C. albicans*.

## METHODS

### Clinical samples

The clinical signs of oral and vaginal candidiasis have previously been described in detail (Naglik *et al.*, 2003). Oral candidiasis included pseudomembranous candidiasis (PC) (thrush), erythematous candidiasis (EC), chronic atrophic candidiasis (CAC), and xerostomia with chronic candidiasis (CC). For vaginal candidiasis, the signs included oedema (swelling), erythema, pseudomembranous plaques, and discharge; and symptoms included pruritus (itch), pain, and soreness. Briefly, samples from individuals who were culture positive for *C. albicans* (for methods, see Naglik *et al.*, 2005) and symptomatic for oral ( $n=40$ ) or vaginal ( $n=40$ ) candidiasis, or culture positive and asymptomatic for oral ( $n=29$ ) and vaginal ( $n=29$ ) candidiasis were compared to asymptomatic oral and vaginal culture-negative (from 10 oral and 10 vaginal samples) non-carriers for the presence of *HWP1* mRNA. Antifungal therapy had not been administered. Colony counts were  $>2 \times 10^3$  c.f.u. ml<sup>-1</sup> or between 2 and  $>10^4$  c.f.u. per swab in oral and vaginal candidiasis, respectively, whereas oral carriers and vaginal carriers had  $<800$  c.f.u. ml<sup>-1</sup> and 4–550 c.f.u. per swab, respectively. The collection of clinical samples was conducted according to the rules of the Guy's and St Thomas' Hospital Trust ethical review board. Informed consent was obtained from all patients regarding the nature of the study.

To determine whether secretory and humoral antibody responses existed in the sample groups, it was necessary to obtain fresh saliva samples from additional subjects, who were classified as patients, carriers and controls according to the above criteria, from individuals attending the Oral Medicine clinic at Guy's Hospital. Serum samples originally collected from some of the individuals in the oral RT-PCR study were supplemented with new samples from patients from the Oral Medicine clinic to obtain adequate numbers for comparison among groups. Sample numbers were based on power calculations from our previous studies (Naglik *et al.*, 1999, 2003).

### RT-PCR analysis of *C. albicans* *HWP1* mRNA expression

*HWP1* gene expression was examined using qualitative and quantitative RT-PCR. A radioactive qualitative RT-PCR was developed to enable detection of the low levels of *C. albicans* mRNA in the oral or vaginal carrier state, a requirement for studies aimed at comparing the carrier state with candidiasis (Naglik *et al.*, 1999, 2003). Four RT-PCR

reactions were performed for each RNA sample prepared from each clinical sample. One reaction to detect *HWP1* was performed, accompanied by three different control reactions: an *ACT1* control to demonstrate the presence or absence of *Candida* species, a negative (water) control, and a positive control using genomic DNA isolated from *C. albicans* NCPF 3156 cells. Each RNA sample was analysed in duplicate, and in many cases in triplicate, to verify *HWP1* gene expression. Complete congruence in gene expression was required in two separate analyses using the same RNA sample. RT-PCR experiments using the *ACT1* and *HWP1* primers were performed using the Access RT-PCR system (Promega). Template RNA was added to RT-PCR mix containing 1× AMV/Tfl buffer (Promega), 1 mM MgSO<sub>4</sub>, 0.1 mM dNTPs, 0.6 mM primers, 3.75 U AMV reverse transcriptase, and 1 μCi (37 kBq) [<sup>32</sup>P]-dCTP (ICN). Radioactive labelling was used to maximize sensitivity. After reverse transcription (48 °C for 45 min), the sample was denatured at 94 °C for 3 min, and 2.5 U Tfl DNA polymerase was added to the reaction (hot start). Cycling times were as follows: denaturation at 94 °C, annealing at 60 °C, and extension at 72 °C, each for 30 s. A final extension at 72 °C for 10 min followed cycling. All radiolabelled RT-PCR products were electrophoresed through a 7 % denaturing 7 M urea polyacrylamide gel, exposed to autoradiography film at -70 °C, and developed. PCR reactions were performed according to the directions of the manufacturer (Promega), incorporating <sup>32</sup>P radiolabel as previously described (Naglik *et al.*, 2003). The primers were: *HWP1* forward, 5'-CCATGTGATGATTACCCACA-3'; and reverse, 5'-GCTGGAACAGAAGATTCAGG-3' (572 bp). Actin (*ACT1*) [primers: forward, 5'-GATTTTGTCTGAACGTGGTAACAG-3'; and reverse, 5'-GGAGTTGAAAGTGGTTTGGTCAATAC-3' (271 bp)] was utilized as a control gene to detect the presence of *C. albicans*. The authenticity of the amplicons was confirmed by DNA sequencing. Real-time, quantitative RT-PCR was performed with the Qiagen SYBR green kit and the ABI 5700 thermal cycler, following the directions of the manufacturers.

## ELISA

Immulon 2 96-well microtitre plates (Thermo Labsystems) were coated with 100 μl of 1 μg ml<sup>-1</sup> purified recombinant Hwp1N13 (rHwp1) consisting of the transglutaminase substrate domain of Hwp1 produced as previously described (Staab *et al.*, 2004) antigen in sodium carbonate buffer, pH 9.6, for 1 h at 37 °C. Unused antigen sites were blocked overnight with 200 μl PBS/BSA/Tween 20. For control experiments, the optimum dilution of the rabbit antisera raised against Hwp1 was 1 : 8000. For sample experiments, the optimum dilutions (with minimal background) for each of the stages to carry out the ELISA on individual patient saliva samples were as follows: saliva at 1 : 2, mouse anti-human IgA at 1 : 100, and rabbit anti-mouse-alkaline phosphatase (AP) conjugate at 1 : 100. For patient serum samples, the dilutions were: serum at 1 : 15, mouse anti-human IgG at 1 : 500, and rabbit anti-mouse-AP at 1 : 500. All incubations took place for 1 h at 37 °C, and after each stage the wells were washed three times with PBS. AP enzyme activity was measured by the addition of 100 μl K-blue substrate (Neogen). The reaction was stopped by the addition of 100 μl 1 M hydrochloric acid after sufficient blue colour had developed, and was measured in an ELISA reader (Anthos) at 405 nm. Positive controls were rabbit antiserum (R1) generated against rHwp1, and purified serum IgG from a pool of 10 patients with oral candidiasis [using protein G HiTrap columns (Pharmacia) (Naglik *et al.*, 2005)]. For each assay, the negative controls were as follows: no rHwp1 coated on the plate; no sample (i.e. saliva or serum); no primary detection antibody. Samples were tested in duplicate and experiments were repeated at least once. IgG and IgA antibody titres to rHwp1 were defined as the inverse of the dilution at which the optical density was twofold greater than background.

## RESULTS AND DISCUSSION

For *C. albicans*, the relationships between factors that contribute to the three states of existence in the host, (1) low-level asymptomatic persistence, i.e. colonization, in the gastrointestinal tract among inhibitory microbial flora, (2) frequent occurrence of asymptomatic oral and vaginal carriage, and (3) virulence in immunocompetent and immunodeficient hosts, are poorly understood.

In this study, the presence of *HWPI* mRNA was found to be correlated with the presence of *C. albicans* in both asymptomatic carriers and in cases of candidiasis at oral and vaginal sites. In the oral cavity, 37/40 individuals with candidiasis and 28/29 carriers were positive for *HWPI* mRNA by the radioactive RT-PCR, whereas 59/59 vaginal samples from carriers and candidiasis cases were positive. RT-PCR results for *ACT1* and *HWPI* were uniformly negative for the culture-negative oral and vaginal control samples.

To quantitate the levels of *HWPI* mRNA during oral candidiasis, real-time PCR was performed on samples (five per group) from individuals with EC or PC. EC is characterized by a reddening of tissue, but not by the raised white aphthae, consisting of mixtures of fungi and epithelial cells, that are found in PC. An *HWPI* message was detected in all samples, with levels that were two- to 20-fold lower than those of the *ACT1* message. Equivalent levels of *HWPI* mRNA were found in individuals with PC or EC [PC, median transcript level of 106 (interquartile ranges 106–188); EC, median 128 (interquartile ranges 128–138), relative to an arbitrary transcript level of 1000 for *ACT1*]. Quantification of *HWPI* mRNA from carriers by real-time PCR was attempted, but was inconsistent and non-reproducible, probably because the amount of message was below the necessary threshold for reliable detection by this method (results not shown).

The RT-PCR results showed that carriers and candidiasis cases were equivalent in terms of the expression of *HWPI*. Furthermore, the five oral specimens in which *HWPI* mRNA was not detected were derived from both groups, indicating that cases were not more likely to have *HWPI* mRNA than candidiasis carriers. In this respect, *HWPI* gene expression results are similar to frequencies seen for *SAP2* and *SAP5*, which are the *SAP* genes most commonly expressed during candidiasis and carriage (Naglik *et al.*, 1999, 2003). However, in the case of *HWPI*, the results suggested an increased level of *HWPI* mRNA in candidiasis cases. *HWPI* mRNA in candidiasis probably exceeded that in asymptomatic conditions, based on the uniformly positive results of RT-PCR in samples from symptomatic, but not asymptomatic, infections. Thus the presence of *HWPI* gene expression appears to parallel the tissue burden, as reflected in the higher levels of c.f.u. in the symptomatic group.

Superimposed upon questions regarding mechanisms of persistence in health are the relative contributions of yeasts, hyphae and pseudohyphal growth forms of *C. albicans* to disease, to carrier states on mucosal surfaces, and to colonization in the lower gastrointestinal tract. The enhanced adherence and invasive properties of hyphal forms relative to yeast forms (Sundstrom, 2006), and the attenuated virulence of mutants unable to form hyphae (Lo *et al.*, 1997), support the belief that equates hyphal forms with invasiveness and yeast forms with commensalism; however, little experimental support exists for this dogma today. Hyphae, pseudohyphae and yeast are found in the same tissue specimens in candidiasis (Sundstrom, 2006), and hyperfilamentous mutants are also attenuated in virulence (Laprade *et al.*, 2002; Bahn *et al.*, 2003), leading to the view that reversible bud–hypha transitions are required for candidiasis. Demonstration of growth morphology during gastrointestinal tract colonization and the carrier state in health is not possible, because of the small numbers of

fungi amongst large numbers of normal bacteria in sites that are not easily amenable to specimen collection.

Because *HWP1* mRNA is abundant in hyphae compared to yeast, an implication of the presence of *HWP1* mRNA in carriers is that hyphal forms are present in the host in the absence of symptoms. This agrees with our previous report, which concluded that hyphae are probably present in most samples, based on widespread expression of *SAP4-6* (Naglik *et al.*, 2003). However, *HWP1* mRNA may also arise from pseudohyphal growth forms (Snide & Sundstrom, 2006), or possibly from yeast forms, although experimental data to support the presence of *HWP1* mRNA in yeast forms has not been reported.

The high frequency of *HWP1* gene expression in oral and vaginal specimens from carriers strongly implicates Hwp1 and hyphae in the ability of *C. albicans* to establish and maintain its presence on mucosal surfaces of human hosts. The results do not support a predominance of yeast forms with absent hyphal forms in the asymptomatic carrier state or in seropositive individuals who lack *C. albicans* in oral or vaginal specimens. It is also unlikely that PC and EC differ with respect to the morphological composition of *C. albicans* in the tissue, since the measurement of *HWP1* gene expression by real-time PCR in EC versus PC showed equivalent levels in both groups.

Specific host determinants that contribute to persistence and candidiasis are also poorly understood. It is not known whether loss of host immunity alone is enough to permit candidiasis, or if upregulation of specific virulence attributes is required. Known immune responses in healthy adults include humoral antibodies to polymannose surface proteins, and cell-mediated immunity to candidal antigens, as demonstrated by positive delayed-type hypersensitivity skin-test reactions to undefined factors in crude extracts that develop early in life (Shannon *et al.*, 1966; Munoz & Limbert, 1977; Lehmann & Reiss, 1980).

To determine if healthy adults also mount immune responses to Hwp1, antibody titres in oral-culture-negative subjects were compared to those of oral carriers and candidiasis cases by ELISA of rHwp1 (Staab *et al.*, 2004). In contrast to *HWP1* gene expression, immune responses to Hwp1 did not correlate with tissue burden in oral samples. Using the non-parametric Kruskal-Wallis analysis of variance, it was found that there were equivalent titres of serum IgG and salivary IgA anti-Hwp1 antibodies in the oral-culture-negative group as well as in the other groups, and there were equivalent titres among the groups also (Table 1). Hwp1, like polymannose, and unlike other antigens that stimulate host responses associated with candidiasis (Strockbine *et al.*, 1984), is a common target of host responses to *C. albicans* that are recognized to result from long-term colonization. The presence of Hwp1 immune responses in healthy adults does not rule out a role for Hwp1 in the pathogenesis of candidiasis.

The common occurrence of mucosal candidiasis accompanying HIV infection raises the possibility that HIV infection might have affected the antibody responses to Hwp1 seen in this study. Although some of the individuals that contributed samples to the RT-PCR results were HIV infected (Naglik *et al.*, 2003), no differences were found between HIV-positive and -negative groups with respect to the presence of Hwp1 mRNA, as detected by RT-PCR. It was not possible to address the effect of HIV infection on salivary anti-Hwp1 IgA levels, because patients providing saliva samples for IgA determinations were not asked about their HIV status. Moreover, those attending the Oral Medicine clinic at Guy's Hospital were deemed to be at low risk for HIV. Given the equivalent anti-Hwp1 antibody levels found in oral-culture-negative controls compared to those of other groups, it is unlikely that the presence of HIV affected the anti-Hwp1 antibody results in this study.

Like the virulence attributes of bacterial pathogens that establish long-term associations with human hosts (Falkow, 2006), Hwp1 appears to be important both in establishing and maintaining the commensal state, as shown by the universal presence of adaptive immune responses, and in candidiasis, as shown by the probable increase in gene expression in this study, by animal studies (Sundstrom *et al.*, 2002), and by *in vitro* adherence assays (Staab *et al.*, 1999). Developmental regulation of Hwp1 implicates hyphal and pseudohyphal forms in both commensalism and candidiasis. *C. albicans* serves as a model organism to elucidate fungal determinants that play a dual role in the establishment of long-term, permanent associations with human hosts, and in tissue invasion leading to disease.

## Acknowledgments

This work was supported by a personal Wellcome Trust Value in People (VIP) award to J. R. N., and National Institutes of Health grants R21 DE015528-01 (J. R. N.) and DE011375 (P. S.). We also thank Ron Wilson for help with statistical analysis. The authors have no conflicting financial interests.

## Abbreviations

EC	erythematous candidiasis
PC	pseudomembranous candidiasis

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**Table 1**

Salivary IgA and serum IgG median antibody titres in the three patient populations. Values shown are median antibody titres, with interquartile ranges (IQRs) in parentheses.

Antibody	Patient population			<i>P</i> value*
	Candidiasis	Carrier	Control	
Salivary IgA	13 (4.0–16.0) ( <i>n</i> =19)	13.5 (6.0–32.0) ( <i>n</i> =16)	8 (5.0–40.0) ( <i>n</i> =13)	0.672
Serum IgG	450 (405–1105) ( <i>n</i> =13)	560 (335–1820) ( <i>n</i> =13)	415 (405–1520) ( <i>n</i> =24)	0.891

\* Non-parametric Kruskal–Wallis analysis of variance.