Abbreviated identification of *Candida albicans* by the presence of a pseudohyphal fringe (‘spiking’ appearance) – some caveats

*Candida albicans* is the most commonly encountered yeast pathogen in our clinical laboratory. Workers such as Barnes & Vale (2005) and Nagaishi & Baron (1997) have described the rapid identification of *C. albicans* solely based on the presence of a pseudohyphal fringe (‘spiking’ colony morphology) with excellent sensitivity (even surpassing that of the well-established germ-tube test) and specificity. This method is also supported by the Clinical and Laboratory Standards Institute M35-A2 document ‘Abbreviated Identification of Bacteria and Yeast’ (CLSI, 2008), which states that colonies growing for less than 48 h on blood-containing medium that exhibit mycelial ‘star-like’ or ‘feet’ projections into the agar may be identified as *C. albicans*. The use of an obvious morphological feature to immediately identify *C. albicans* is of great interest to us as it would save considerable time, effort and cost, and potentially translate to more timely information being conveyed to the clinician.

We therefore evaluated the performance of this method using our standard trypticase soy agar with 5% sheep blood (TSA–BAP), following overnight (18–24 h) incubation in a CO₂ incubator, using our standard trypticase soy agar with 5% sheep blood (TSA–BAP), and note to identify *C. albicans/C. dubliniensis*. Of these 42 germ-tube positive isolates, only 33 showed the presence of a pseudohyphal fringe (‘spiking’), yielding a sensitivity of 78.6% for the observation of ‘spiking’ in *C. albicans/C. dubliniensis*. Of the 70 germ-tube negative yeast isolates, 67 did not show ‘spiking’, yielding a specificity of 95.7%. The overall inter-test agreement was good (Cohen’s κ=0.76). These results are summarized in Table 1. Interestingly, the three germ-tube negative isolates that demonstrated pseudohyphal ‘spiking’ showed the presence of arthroconidia (in addition to blastoconidia) on microscopic (Gram-stain) examination, and further phenotypic workup confirmed these isolates to be *Trichosporon* sp. rather than *C. albicans*.

Even though we found this rapid method of morphological identification of *C. albicans/C. dubliniensis* to be less sensitive than previously described, it appears to be feasible to use it for the presumptive identification of yeasts for semi-critical and non-sterile sites. It is, however, crucial to note that non-*Candida* yeasts (such as *Trichosporon* spp.) may give false-positive results for ‘spiking’, as in our study. Although the presence of a pseudohyphal fringe in colonies of *Trichosporon* spp. has been recognized previously (www.mycology.adelaide.edu.au – Mycology Online, University of Adelaide, Australia), the possibility of mistaking such colonies for ‘spiking’ colonies of *C. albicans*, to the best of our knowledge, has not been described. As the susceptibility profiles of these two genera of yeasts may differ significantly (Pfaller & Diekema, 2004), it would be crucial for the clinical laboratory to differentiate between the two, especially when the isolate is derived from a sterile site or suspected to be causing a more serious illness. We therefore suggest that the Gram-stain morphology of any ‘spiking’ isolate be carefully examined for the presence of arthroconidia (which are indicative of *Trichosporon* spp.). Additional analysis, such as urease testing, may be indicated in suspicious cases.

### Table 1. Comparison of the performance of pseudohyphal fringe (‘spiking’) appearance at 18–24 h incubation on 5% TSA–BAP in CO₂ to the germ-tube test using horse serum in identifying *C. albicans/C. dubliniensis*

<table>
<thead>
<tr>
<th>Pseudohyphal fringe (‘spiking’)</th>
<th>Germ tube</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Positive (+)</td>
<td>Negative (−)</td>
</tr>
<tr>
<td>Present (+)</td>
<td>33</td>
<td>3</td>
</tr>
<tr>
<td>Absent (−)</td>
<td>9</td>
<td>67</td>
</tr>
<tr>
<td>Total</td>
<td>42</td>
<td>70</td>
</tr>
</tbody>
</table>

Sensitivity of ‘spiking’ in comparison to the germ-tube test = 33/42 = 78.6%; specificity of ‘spiking’ in comparison to the germ-tube test = 67/70 = 95.7%; positive predictive value of ‘spiking’ in comparison to germ-tube test = 33/36 = 91.7%; negative predictive value of ‘spiking’ in comparison to germ-tube test = 67/76 = 88.2%; overall inter-test agreement (Cohen’s κ) = 0.76 (0.64–0.89).
There are some limitations to our study. The lower sensitivity observed in our study may possibly be due to the use of an abridged incubation time of 18–24 h, as compared to the 24–48 h described in other studies (Barnes & Vale, 2005; Nagaishi & Baron, 1997). However, a shorter incubation period may be more realistic in the clinical laboratory setting, where clinicians may expect quicker delivery of preliminary information to guide patient management. Furthermore, adoption of this method by our laboratory would result in a significant reduction in our germ-tube testing volume, translating into savings in terms of labour and cost of reagents.

Neither the pseudohyphal fringe nor the germ-tube test allows confident separation between C. albicans and C. dubliniensis. For instance, Barnes & Vale (2005) noted that 20% of C. dubliniensis isolates produced ‘spiking’ colonies and 40% showed positive germ-tube test results. We cannot determine what proportion of our study isolates were C. dubliniensis; however, recent susceptibility data from Pfaller et al. (2010) seem to suggest that the antifungal susceptibility profile C. dubliniensis is comparable with that of C. albicans. Further testing to separate these species may be indicated for critical infections.

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