Screening Protocol for Torulopsis (Candida) glabrata

GEOFFREY LAND,^{1*} JACKIE BURKE,¹ CAROL SHELBY,¹ JUDITH RHODES,² JOYCE COLLETT,² ISABELLE BENNETT,³ AND JETTIE JOHNSON³

Clinical Microbiology and Mycology Laboratories, Methodist Medical Center, Dallas, Texas 75203¹; University Hospitals, The University of Cincinnati School of Medicine, Cincinnati, Ohio 45267²; and Baylor University Medical Center, Dallas, Texas 75246³

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A screening test has been developed for the presumptive identification of *Torulopsis* (*Candida*) glabrata from other common clinical isolates of yeast-like fungi. An interlaboratory comparison of a protocol consisting of morphology on cornneal Tween 80 agar and trehalose fermentation at 42°C was successful in differentiating *T. glabrata* from other taxa that are frequent or possible clinical isolates. The screening results for 517 clinical yeast isolates, 241 of which were *T. glabrata*, were compared with their final identification via commercial systems (API 20C Yeast Identification System [bioMERIEUX, Hazelwood, Mo.] and Rapid Yeast Identification Panel [Dade Microscan, Sacramento, Calif.]). The trehalose screening test has a sensitivity and a specificity of 97.8 and 95.8%, respectively, and a positive predictive value of 97.4% and a negative predictive value of 96.5%. Overall, the trehalose screen had an efficiency rating of 93.9% for ruling in or out *T. glabrata*. Since *T. glabrata* represents a substantial part of the workload in a clinical laboratory, a significant reduction in direct and indirect costs should be realized.

Torulopsis glabrata, a member of the family Cryptococcaceae, is an anamorphic, nonfilamentous yeast inhabiting the gastrointestinal and genitourinary tracts of humans and animals (20, 23). Although its possible association with bronchopneumonia was reported in 1937, T. glabrata has long been considered a saprophyte because it is most often isolated from stools, skin, mouth, vagina, and improperly collected urine specimens (16, 20, 21). Recently, T. glabrata has emerged as an opportunistic pathogen in patients with severe underlying primary diseases and/or undergoing complex therapeutic regimens. It has been the suspected etiologic agent of a variety of infections including meningitis (1), endocarditis (4, 8), cholecystitis (20), osteomyelitis (5, 15), spondylitis (3), hyperplasia and infection of the oral pharynx (6, 18), and fungemia (8, 16, 21). Amphotericin B and/or flucytosine have been the standard therapy for these infections, augmented by surgery when necessary (10, 21). However, a potential therapeutic problem is that some clinical strains of T. glabrata readily show in vitro resistance to the synthetic imidazoles, suggesting that imidazole resistance could be an in vivo problem as well (7, 10, 11, 24).

As a result of its frequent association with humans, *T. glabrata* is usually one of the three most common yeasts isolated in the clinical laboratory. Because of its ubiquity, possible clinical relevance, and potential therapeutic problems, it has become necessary to identify *T. glabrata* in a timely and costeffective fashion. Stockman and Roberts developed a screening test based upon *T. glabrata*'s ability to assimilate trehalose in the presence of cycloheximide within 1 h, when incubated at $37^{\circ}C$ (22). The test was in a microtiter format, and by equating bromcresol green reduction with trehalose metabolism and observing the plate coordinates, large numbers of isolates could be surveyed at one time. Although this was an excellent approach, the medium has not been made commercially available, nor have published studies been forthcoming as to its efficacy.

Our goal was to develop a screening test for *T. glabrata* that used commercially available reagents, was cost-effective, and

Preliminary studies (data not shown) revealed that we could combine growth at elevated temperatures with trehalose fermentation and that either a heating block or an incubator could be used for incubation. Additionally, we found that we could usually use yeasts from primary cultures on chocolate agar, blood agar, or Columbia CNA. Other modifications which may be used include substituting a deep mineral oil overlay for Vaspar and bromthymol blue dye instead of bromcresol purple as an indicator (yeast fermentation broth with bromthymol blue with trehalose with Durham tube and yeast fermentation broth with bromcresol purple with trehalose with Durham tube, respectively [Remel Laboratories, Lenexa, Kans.]).

From this departure point, we developed the following protocol based upon standard procedures (20, 23). Upon isolation of a yeast from a primary clinical specimen, a germ tube test was performed, morphology medium (cornmeal Tween 80 or its equivalent) was inoculated, and when necessary, a plate was streaked for isolation of pure colonies. Growth from pure

offered a relatively rapid turnaround time. In reviewing two yeast compendia and comparing a number of tests, it became apparent that no one test would work with the effectiveness of the germ tube test (2, 12). However, we felt we could capitalize on a combination of tests consisting of morphology on cornmeal agar and two unique characteristics, namely, fermentation of the nonreducing disaccharide trehalose and growth at 42°C (Table 1) (2, 12). Of 33 yeast taxa considered frequent or possible clinical isolates (23), 5 grow well at 42°C and 10 exhibit strain-dependent or variable growth (Table 1). The concomitant ability of these same yeasts to ferment trehalose in a standard Wickerham broth is variable among seven species (Candida albicans, Candida lusitaniae, Candida parapsilosis, Candida tropicalis, Candida viswanathii, Hansenula polymorpha, and Saccharomyces cerevisiae), whereas only T. glabrata consistently does both. Since cornmeal morphology or its equivalent was required by the commercial systems already in place in our respective laboratories (API 20C and Dade Microscan's Rapid Yeast Identification Panel), only media for the occasional growth plate, media for trehalose fermentation, and a means to attain 42°C had to be supplied.

^{*} Corresponding author.

	1 8 8	1 ())			
Organism	Morphology on CMA	42°C growth	Trehalose fermentation		
Torulopsis glabrata	MLB, NF	+	+		
Blastoschizomyces capitatus	FSN, TH, Ar	+	_		
Candida albicans	MLB, S/E PSH, TH	+	V		
Candida catenulata	MLB, S/E PSH	_	_		
Candida guilliermondii	MLB, F, N to S/E PSH	V	_		
Candida kefyr	MLB, F, N to S/E PSH	V	_		
Candida krusei	MLB, S/E PSH	V	_		
Candida lambica	MLB, S/E PSH	_	_		
Candida lipolytica	MLB, F, N to E PSH	_	_		
Candida lusitaniae	MLB, F, N to E PSH	V	V		
Candida parapsilosis	MLB, S/E PSH	V	V		
Candida rugosa	MLB, S/E PSH	_	_		
Candida tropicalis	MLB, S/E PSH, TH	+	V		
Candida viswanathii	MLB, E PSH, TH	V	V		
Candida zeylanoides	MLB, F, N to S/E PSH	_	D		
Cryptococcus neoformans	MLB, NF	_	_		
Cryptococcus albidus	MLB, NF	_	_		
Cryptococcus laurentii	MLB, NF	_	_		
Cryptococcus luteolus	MLB, NF	_	_		
Cryptococcus terreus	MLB, NF	_	_		
Cryptococcus uniguttulatus	MLB, NF	_	_		
Debaryomyces spp.	PB, F	_	_		
Geotrichum candidum	FSN, TH, Ar	_	_		
Hansenula anomala	MLB, F, N to S/E PSH	_	_		
Hansenula polymorpha	MLB, NF	+	V		
Pichia ohmeri	MLB, E PSH	V	_		
Prototheca wickerhamii	EC	_	_		
Rhodotorula glutinis	MLB, NF	_	_		
Rhodotorula minuta	MLB, NF	_	_		
Rhodotorula pilimanae	PB, NF	_	_		
Rhodotorula rubra	MLB, NF	_	_		
Saccharomyces cerevisiae	MLB, F, N to S PSH	V	V		
Torulopsis candida	MLB, F, N or S PSH	_	D		
Torulopsis pintolopesii	MLB, F, N or S PSH	V	_		
Trichosporon beigelii	FSN, PB, TH, S/E PSH, Ar	V	_		
Trichosporon pullulans	FSN, PB, TH, S/E PSH, Ar	—	—		

TABLE 1. The ability	to grow at 42°C an	d ferment trehalos	e for those yeast	ts considered to b	e possible	clinical isolates
an	d their morphology	on cornmeal Twee	en 80 agar or its	equivalent (2, 12,	$(23)^{a}$	

^{*a*} CMA, cornneal agar; MLB, multiple lateral budding; N, no/none; F, filaments; S, simple; E, elaborate; PSH, pseudohyphae; TH, true or septate hyphae; PB, polar budding; Ar, arthroconidia; EC, endocytoplasmic cleavage; FSN, vegetative reproduction by splitting. Example: MLB, F, N to S/E PSH = multiple lateral budding; filaments, none to simple or elaborate pseudohyphae. V, strain variable; D, delayed fermentation (\geq 36 h).

cultures of small (3 to 5 μ m in diameter), nonfilamentous yeasts were transferred to 7.0-ml Durham yeast fermentation tubes containing bromcresol purple (0.0016%) and 1% trehalose to the density of a 3 or 4 McFarland standard (Remel), overlaid with melted paraffin or Vaspar, and incubated at 42°C for 24 h. For those isolates demonstrating sparse primary growth, the trehalose tube was inoculated from a growth plate on the following day. A positive result was the production of both acid and gas in trehalose broth at the elevated temperature. All yeasts were completely identified via the API 20C or Dade Microscan Yeast Identification Panel commercial system, according to their respective manufacturer's directions, and microscopic morphology was determined at 24 to 48 h.

In a survey of 517 clinical yeast isolates, isolates of *T. glabrata* and *C. tropicalis* were the only taxa that fermented trehalose at 42°C (Table 2). *C. tropicalis*, however, was an inconsistent fermenter as only 5% (4 of 87) strains produced both acid and gas in trehalose at 42°C, while 96% (231 of 241) of *T. glabrata* isolates were positive (Table 2, P = <0.0001, chi square with Yates' correction). *T. glabrata* was readily differentiated from *C. tropicalis* by the rapid fermentation of trehalose (8 to 24 h) and the lack of pseudohyphae or filaments on morphology medium. None of the other frequently encountered yeasts with the potential to ferment trehalose appeared

capable of doing so at the restrictive temperature. Several taxa demonstrated delayed acid production in trehalose broth at $42^{\circ}C (\geq 36 \text{ h})$ but did not produce both acid and gas (Table 2). Some isolates of *S. cerevisiae* bear a slight morphological resemblance to *T. glabrata* and are said to survive elevated temperatures and ferment trehalose (12, 21) and thus have the potential to be confused with *T. glabrata*. However, the 12 isolates that we tested did not appear capable of fermenting trehalose above $37^{\circ}C$. Should a rare strain of *S. cerevisiae* be found capable of fermenting trehalose at the restrictive temperature, it can be differentiated from *T. glabrata* on the basis of size and the inability of the latter to form rudimentary pseudohyphae composed of short chains of elongated budding cells.

The overall sensitivity and specificity for the trehalose screen were 97.8 and 95.8%, respectively. This translated to a positive predictive test value of 97.4% and a negative predictive value of 96.5%, indicating that 93.9% of the time the screen would effectively rule in or rule out *T. glabrata*. These values are well within the range considered acceptable for the *C. albicans* germ tube test (14). In using the trehalose screen for 6 months in a routine clinical microbiology laboratory (Methodist Medical Center), there were three instances (333 total isolates, 0.9%) in which a yeast failing to ferment trehalose at 42°C

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OrganismNo. of strainsMorphology on CMA 42° Crehalose"Time positive (b)Torulopsis (Candida) glabrata241MLB, NF(231) A/G8–18Blastoschizovyces capitatus5FSN, TH, ArCandida albicans31°MLB, S/E PSH, THCandida quillermondii5MLB, F, N to S/E PSHCandida furbia6MLB, S/E PSHCandida furbia6MLB, S/E PSHCandida furbia6MLB, S/E PSHCandida furbia2MLB, F, N to S/E PSHCandida turbica6MLB, S/E PSHCandida positionia2MLB, S/E PSHCandida nogii2MLB, S/E PSHCandida philyla2MLB, S/E PSHCandida rugosa2MLB, S/E PSHCandida rugosa2MLB, S/E PSHCandida valvanutii4MLB, S/E PSHCandida svalvanotifi4MLB, S/E PSHCandida tropicalis1MLB, S/E <th></th> <th></th> <th></th> <th>in the b</th> <th colspan="2"></th>				in the b		
$ \begin{array}{cccccccc} Torology is (Candida) glabrata 241 MLB, NF (231) A/G 8-18 \\ Blastoschizomyces capitatus 5 FSN, TH, Ar \\ Candida albicans 31c MLB, S/E PSH, TH \\ Candida catenulata 1 MLB, S/E PSH \\ Candida catenulata 1 MLB, F, N to S/E PSH \\ Candida kefyr 1 MLB, F, N to S/E PSH \\ Candida kefyr 1 MLB, F, N to S/E PSH \\ Candida kefyr 2 MLB, S/E PSH \\ Candida tambica 6 MLB, S/E PSH \\ Candida tambica 2 MLB, F, N to E PSH \\ Candida parapison 2 MLB, F, N to E PSH \\ Candida parapison 2 MLB, F, N to E PSH \\ Candida parapison 2 MLB, F, N to E PSH \\ Candida parapison 2 MLB, F, N to E PSH \\ Candida parapison 2 MLB, F, N to E PSH \\ Candida parapison 2 MLB, F, N to E PSH \\ Candida parapison 2 MLB, F, N to FPSH \\ Candida ragosa 2 MLB, S/E PSH \\ Candida ragosa 2 MLB, S/E PSH \\ Candida ragosa 2 MLB, F, N to F PSH \\ Candida ragosa 3 MLB, F, N to F PSH \\ Candida ragosa 3 MLB, F, N to F PSH \\ Candida ragosa 4 MLB, F, N to F PSH \\ Candida spinondes 3 MLB, F, N to S/E PSH \\ Candida spinondes 3 MLB, F, N to S/E PSH \\ Candida spinontes 4 MLB, NF \\ Candida spinontes 1 MLB, NF \\ Candida spinontes remus 1 MLB, NF \\ Candida spinontes 1 MLB, NF \\ Cand$	Organism	No. of strains	Morphology on CMA	42°C trehalose ^b	Time positive (h)	
Blassochizomyces capitatus5FSN, TH, ArCandida divisoras31°MLB, SE PSH, THCandida catenulata1MLB, SE PSH, THCandida guillermondii5MLB, F, N to S/E PSHCandida kgly1MLB, F, N to S/E PSHCandida kfly1MLB, SF, N to S/E PSHCandida knosci8MLB, SE PSHCandida lusitaniae2MLB, F, N to E PSHCandida parapsilosis24MLB, SE PSHCandida pringia2MLB, SF, PSHCandida pringia2MLB, SF, PSHCandida pringia2MLB, SF, PSHCandida pringias2MLB, SF, PSHCandida pringias2MLB, SF, PSHCandida viswanathi8PSH, THCandida viswanathi4MLB, SF, PSHCandida viswanathi1MLB, NFCandida zeylanoides3MLB, F, N to S/E PSHCandida viswanathi1MLB, NFCandida zeylanoides1MLB, NFCandida zeylanoides1MLB, NFCondida viswanathi1MLB, NFCondida viswanathi1MLB, NFCondida viswanathi1 <td>Torulopsis (Candida) glabrata</td> <td>241</td> <td>MLB, NF</td> <td>(231) A/G</td> <td>8-18</td>	Torulopsis (Candida) glabrata	241	MLB, NF	(231) A/G	8-18	
$ \begin{array}{ccc} Candida abicans & 31^c & MLB, SE PSH, TH & - & - & - \\ Candida catenulata & 1 & MLB, SE PSH & - & - & - \\ Candida catenulata & 1 & MLB, F, N to S/E PSH & - & - & - \\ Candida keyr & 1 & MLB, F, N to S/E PSH & - & - & - \\ Candida kusci & 8 & MLB, SE PSH & - & - & - \\ Candida lambica & 6 & MLB, SE PSH & - & - & - \\ Candida nogü & 2 & MLB, F, N to E PSH & - & - & - \\ Candida parapsilosis & 24 & MLB, SF PSH & - & - & - \\ Candida parapsilosis & 24 & MLB, SF PSH & - & - & - \\ Candida parapsilosis & 24 & MLB, SF PSH & - & - & - \\ Candida parapsilosis & 24 & MLB, SF, NPSH & - & - & - \\ Candida rugosa & 2 & MLB, SF, NPSH & - & - & - \\ Candida rugosa & 2 & MLB, SF, NPSH & - & - & - \\ Candida tropicalis & 87 & MLB, SF, PSH & - & - & - \\ Candida tropicalis & 87 & MLB, SF, NPSH & - & - & - \\ Candida viswanathii & 4 & MLB, E PSH, TH & (4) A/G \geq 236 \\ Candida viswanathii & 4 & MLB, F, N to SFE PSH & - & - \\ Candida synonides & 3 & MLB, F, N to SFE PSH & - & - \\ Candida spinonides & 1 & MLB, NF & - & - \\ Candida spinonides & 1 & MLB, NF & - & - \\ Candida spinonides & 1 & MLB, NF & - & - \\ Captococcus albidus & 1 & MLB, NF & - & - \\ Captococcus albidus & 1 & MLB, NF & - & - \\ Captococcus albidus & 1 & MLB, NF & - & - \\ Captococcus subiduts & 1 & MLB, NF & - & - \\ Captococcus subiduts & 1 & MLB, NF & - & - \\ Captococcus subiduts & 1 & MLB, NF & - & - \\ Captococcus subiduts & 1 & MLB, NF & - & - \\ Captococcus subiduts & 1 & MLB, NF & - & - \\ Captococcus subiduts & 1 & MLB, NF & - & - \\ Captococcus subiduts & 1 & MLB, NF & - & - \\ Captococcus subiduts & 1 & MLB, NF & - & - \\ Captococcus subiduts & 1 & MLB, NF & - & - \\ Captococcus subiduts & 1 & MLB, NF & - & - \\ Captococcus subiduts & 1 & MLB, NF & - & - \\ Captococcus subidutis & 1 & MLB, NF & - & - \\ Captococcus subidutis & 1 & MLB, NF & - & - \\ Captococcus subidutis & 1 & MLB, NF & - & - \\ Captococcus subidutits & 1 & MLB, NF & - & - \\ Captococcus subidutits & 1 & MLB, NF & - & - \\ Captococcus subidutita & 3 & MLB, F, N to S/E PSH & - & - \\ Captococcus subidutita & 3 $	Blastoschizomyces capitatus	5	FSN, TH, Ar	_ /	-	
	Candida albicans	31^{c}	MLB, S/E PSH, TH	_	-	
	Candida catenulata	1	MLB, S/E PSH	_	-	
$ \begin{array}{llllllllllllllllllllllllllllllllllll$	Candida guilliermondii	5	MLB, F, N to S/E PSH	_	-	
Candida krinsei8MLB, S/E PSHCandida lambica6MLB, S/E PSHCandida lusitaniae2MLB, F, N to E PSHCandida pogi2MLB, PSH/clusters of BCandida propsilosis24MLB, S/E PSHCandida propsilosis24MLB, S/E PSHCandida propsilosis24MLB, S/E PSHCandida rugosa2MLB, F, N to E PSHCandida vigosa2MLB, S/E PSH, TH(4) A/G \geq 36Candida vigosa87MLB, S/E PSH, THCandida sellatoidea2MLB, F, N to S/E PSHCandida sellatoides3MLB, F, N to S/E PSHCandida sequencies3MLB, SFCandida sequencies3MLB, NFCandida sequencies1MLB, NFCondida sequencies2MLB, NFCondida sequencies1MLB, NFCoptococcus laurenti1MLB, NFCoptococcus uniguitulatus1MLB, NFCoptococcus uniguitulatus1MLB, NFCoptococcus uniguitulatus1MLB, NFCoptococcus uniguitulatus1MLB, NF <t< td=""><td>Candida kefyr</td><td>1</td><td>MLB, F, N to S/E PSH</td><td>_</td><td>-</td></t<>	Candida kefyr	1	MLB, F, N to S/E PSH	_	-	
Candida lambica6MLB, SE PSHCandida lusitaniae2MLB, F, N to E PSHCandida progi2MLB, F, N to E PSHCandida progi2MLB, SE PSHCandida prilyla2MLB, SF, NPSHCandida rugosa2MLB, SF, PSHCandida stellatoidea2MLB, F, N to PSHCandida visvanathi4MLB, SE PSH, TH(4) A/G \geq 36Candida zeylanoides3MLB, F, N to SE PSHCandida zeylanoides3MLB, F, N to SE PSHCandida zeylanoides3MLB, F, N to SE PSHCandida zeylanoides3MLB, FNCandida zeylanoides1MLB, NFCoptococcus lateofus1MLB, NFCoptococcus lateofus1MLB, NFCoptococcus luteolus2MLB, NFCoptococcus unguttulatus1MLB, NFDebaryomyces sp.1PB, FCharlea anomala6MLB, NFCharlea cubifus1MLB, NFCharlea cubifus1MLB, NFCharlea cubifus1MLB, NFCoptococcus unguttulatus1MLB, NF <td>Candida krusei</td> <td>8</td> <td>MLB, S/E PSH</td> <td>_</td> <td>-</td>	Candida krusei	8	MLB, S/E PSH	_	-	
Candida lusitaniae2MLB, F, N to E PSHCandida mogi2MLB, PSH/clusters of BCandida parapsilosis24MLB, S/E PSHCandida parapsilosis24MLB, S/E PSHCandida progosa2MLB, S/E PSH(1) A ≥ 36 Candida trogosa2MLB, F, N to E PSHCandida tellatoidea2MLB, S/E PSHCandida trogosa3MLB, S/E PSH, TH(4) A/G ≥ 36 Candida viswanathii4MLB, E PSH, THCandida seylanoides3MLB, NFCandida seylanoides3MLB, NFCandida seylanoides1MLB, NFCandida seylanoides1MLB, NFCoptococcus aleoformans4MLB, NFCryptococcus luteolus2MLB, NFCryptococcus luteolus1MLB, NFCoptococcus uniguttulatus1MLB, NFCoptococcus uniguttulatus1MLB, NFChansenula polymorpha11MLB, NFHansenula polymorpha11MLB, NFPrototheca wickerhamii3MLB, NFRhodotorula giutinis15MLB, NFPrototheca wickerhamii3MLB, NFRhodotorul	Candida lambica	6	MLB, S/E PSH	_	-	
Candida mogü2MLB, PSH/Clusters of BCandida prapsilosis24MLB, S/E PSHCandida philyla2MLB, NF, NPSHCandida rugosa2MLB, S/E PSH(1) A ≥ 36 Candida tellatoidea2MLB, S/E PSH, TH(4) A/G ≥ 36 Candida viswanathii4MLB, E PSH, THCandida viswanathii4MLB, F, N to S/E PSHCandida viswanathii4MLB, PSHCandida zylanoides3MLB, F, N to S/E PSHCandida zylanoides3MLB, NFCandida zylanoides1MLB, NFCryptococcus neformans4MLB, NFCryptococcus laurentii1MLB, NFCryptococcus laurentii1MLB, NFCryptococcus laurentii1MLB, NFCryptococcus unigutudatus1MLB, NFCoptococcus unigutudatus1MLB, NFChansenula anomala6MLB, F, N to S/E PSHAnsenula anomala6MLB, FNProtoheca wickrhamii3ECRhodotorula glutinis15MLB, NFRhodotorula glutinis15MLB, NFRhodotorula glutinis15MLB, NFRhodotorul	Candida lusitaniae	2	MLB, F, N to E PSH	_	-	
Candida parapsilosis24MLB, S/E PSHCandida philyla2MLB, NF, NPSHCandida trogosa2MLB, S/E PSH(1) A \geq 36Candida trogicalis87MLB, S/E PSHCandida trogicalis87MLB, S/E PSH, TH(4) A/G \geq 36Candida trogicalis87MLB, S/E PSH, THCandida viswanathii4MLB, S/E PSH, THCandida seylanoides3MLB, F, N to S/E PSHCandida seylanoides3MLB, NFCandida seylanoides1MLB, NFCandida seylanoides2MLB, NFCryptococcus albidus1MLB, NFCryptococcus laurentii1MLB, NFCryptococcus tereus1MLB, NFCryptococcus tereus1MLB, NFCoptococcus tereus1MLB, NFDebaryomyces spp.1PB, FHansenula anomala6MLB, F, N to S/E PSHHansenula anomala3MLB, NFRhodotorula glutinis15MLB, NFRhodotorula glutinis15MLB, NFRhodotorula glutinis15MLB, NFRhodotorula glutinis15MLB, NFRhodotorula glutinis15	Candida mogii	2	MLB, PSH/clusters of B	_	-	
Candida philyla2MLB, NF, NPSHCandida rugosa2MLB, SF, PSH(1) A ≥ 36 Candida sellatoidea2MLB, SF, N to E PSHCandida tropicalis87MLB, S/E PSH, TH(4) A/G ≥ 36 Candida viswanathii4MLB, E PSH, THCandida seylanoides3MLB, F, N to S/E PSHCandida zyelanoides3MLB, NFCandida seylanoides3MLB, NFCandida seylanoides1MLB, NFCandida seylanoides1MLB, NFCryptococcus albidus1MLB, NFCryptococcus laurentii1MLB, NFCryptococcus lurentii1MLB, NFCryptococcus lurentii1MLB, NFCryptococcus lurentii1MLB, NFCryptococcus lurentii1MLB, NFCryptococcus lungutulatus1MLB, NFCharlen anomala6MLB, F, N to S/E PSHCharlen anomala6MLB, F, N to S/E PSHPichia ohmeri4MLB, NFPichia ohmeri3ECRhodotorula glutinis15MLB, NFRhodotorula glutininae1PB, NFRhodotorula plimanae1 <td>Candida parapsilosis</td> <td>24</td> <td>MLB, S/E PSH</td> <td>_</td> <td>-</td>	Candida parapsilosis	24	MLB, S/E PSH	_	-	
Candida rugosa2MLB, S/E PSH(1) A ≥ 36 Candida rojcalis2MLB, F, N to E PSHCandida torpicalis87MLB, S/E PSH, TH(4) A/G ≥ 36 Candida viswanathii4MLB, E PSH, THCandida zeylanoides3MLB, F, N to S/E PSHCandida spp. not otherwise identified5MLB, S/ECryptococcus neoformans4MLB, NFCryptococcus albidus1MLB, NFCryptococcus lurentii1MLB, NFCryptococcus lurentis2MLB, NFCryptococcus lurentis1MLB, NFCryptococus lurentis1MLB, NFCryptococus lurentis1MLB, NFCryptococus lurentis1MLB, NFCryptococus lurentis1MLB, NFCryptococus l	Candida philyla	2	MLB, NF, NPSH	_	-	
Candida stellatoidea2MLB, F, N to E PSHCandida tropicalis87MLB, S/E PSH, TH(4) A/G ≥ 36 Candida viswanathii4MLB, E PSH, THCandida viswanathii3MLB, F, N to S/E PSHCandida seylanoides3MLB, F, N to S/E PSHCandida seylanoides3MLB, SFCandida seylanoides1MLB, NFCryptococcus neoformans4MLB, NFCryptococcus latentii1MLB, NFCryptococcus luteolus2MLB, NFCryptococcus luteolus1MLB, NFCryptococcus luteolus1MLB, NFCryptococcus uniguttulatus1MLB, NFDebaryomyces spp.1PB, FGeotrichum candidum7FSN, TH, ArHansenula anomala6MLB, F, N to S/E PSHHansenula polymorpha11MLB, NFPrototheca wickerhamii3ECRhodotorula glutinins15MLB, NFRhodotorula glutinins1PB, NFRhodotorula glutinins1PB, NFRhodotorula glutinins1PB, NFChichosopesit1MLB, NFChichosopesitie1	Candida rugosa	2	MLB, S/E PSH	(1) A	≥36	
Candida tropicalis87MLB, S/E PSH, TH(4) A/G≥36Candida viswanahii4MLB, E PSH, THCandida zeylanoides3MLB, F, N to S/E PSHCandida spp. not otherwise identified5MLB, NFCandida spp. not otherwise identified5MLB, NFCryptococcus albidus1MLB, NFCryptococcus laurentii1MLB, NFCryptococcus laurentii1MLB, NFCryptococcus laurentii1MLB, NFCryptococcus laurentii1MLB, NFCryptococcus uniguttulatus1MLB, NFDebaryomyces spp.1PB, FGeotrichum candidum7FSN, TH, ArHansenula polymorpha11MLB, NFProtoheca wickerhamii3ECProtoheca wickerhamii3BCRhodotorula glutinis15MLB, NFRhodotorula nimuta3MLB, NFRhodotorula rubra2MLB, NFProtoheca wickerhamii2MLB, NFProtoheca wickerhamii3ECRhodotorula nimuta3MLB, NFRhodotorula nimuta2MLB, NFRhodotorula nibuta2 <t< td=""><td>Candida stellatoidea</td><td>2</td><td>MLB, F, N to E PSH</td><td></td><td>-</td></t<>	Candida stellatoidea	2	MLB, F, N to E PSH		-	
Candida viswanathii4MLB, E PSH, THCandida zeylanoides3MLB, F, N to S/E PSHCandida spp. not otherwise identified5MLB, PSHCryptococcus neoformans4MLB, NFCryptococcus labidus1MLB, NFCryptococcus latentii1MLB, NFCryptococcus luteolus2MLB, NFCryptococcus luteolus1MLB, NFCryptococcus luteolus1MLB, NFCryptococcus uniguttulatus1MLB, NFCryptococcus uniguttulatus1MLB, NFChansenula anomala6MLB, F, N to S/E PSHHansenula nonmala6MLB, F, N to S/E PSHHansenula polymorpha11MLB, NFPrototheca wickerhamii3ECRhodotorula giluinis15MLB, NFRhodotorula pilumanae1PB, NFRhodotorula pilutuatu3MLB, NFRhodotorula pilunanae1PB, NFRhodotorula piluta2MLB, NFRhodotorula piluta3MLB, NFRhodotorula piluta3MLB, NFRhodotorula piluta2MLB, NFRhodotorula piluta2<	Candida tropicalis	87	MLB, S/E PSH, TH	(4) A/G	≥36	
Candida zeylanoides3MLB, F, N to S/E PSHCandida spp. not otherwise identified5MLB, PSHCryptococcus neoformans4MLB, NFCryptococcus albidus1MLB, NFCryptococcus laurentii1MLB, NFCryptococcus turious2MLB, NFCryptococcus turious1MLB, NFCryptococcus uniguitulatus1MLB, NFCryptococcus uniguitulatus1MLB, NFDebaryomyces spp.1PB, FGeotrichum candidum7FSN, TH, ArHansenula anomala6MLB, NFHansenula polymorpha11MLB, NFProtoheca wickerhamii3ECRhodotorula glutinis15MLB, NFRhodotorula plutinata3MLB, NFRhodotorula glutinis1PB, NFRhodotorula glutinis1PB, NFRhodotorula plutanae1PB, NFRhodotorula rubra2MLB, NFTorulopsis candida2MLB, F, N or S PSHTorulopsis candida2MLB, F, N or S PSHTorulopsis pintolopesii1MLB, F, N or S PSHTrichosporon pullulans <td< td=""><td>Candida viswanathii</td><td>4</td><td>MLB, E PSH, TH</td><td></td><td>-</td></td<>	Candida viswanathii	4	MLB, E PSH, TH		-	
Candida sp. not otherwise identified5MLB, PSHCryptococcus neoformans4MLB, NF $ -$ Cryptococcus albidus1MLB, NF $ -$ Cryptococcus laurentii1MLB, NF $ -$ Cryptococcus laurentii2MLB, NF $ -$ Cryptococcus laurentii1MLB, NF $ -$ Cryptococcus uniguttulatus1MLB, NF $ -$ Cryptococcus uniguttulatus1MLB, NF $ -$ Debaryomyces spp.1PB, F $ -$ Geotrichum candidum7FSN, TH, Ar $ -$ Hansenula anomala6MLB, F, N to S/E PSH $ -$ Prototheca wickerhamii3EC $ -$ Rhodotorula glutinis15MLB, NF $ -$ Rhodotorula nitura1PB, F $ -$ Rhodotorula rubra2MLB, NF $ -$ Rhodotorula plutinis15MLB, NF $ -$ Rhodotorula nitura2MLB, NF $ -$ Saccharomyces cerevisiae12MLB, NF $ -$ Torulopsis candida2MLB, NF $ -$ Torulopsis pintolopesii1MLB, F, N or S PSH $ -$ Torulopsis pintolopesii6FSN, PB, TH, S/E PSH, Ar $ -$ Trichosporon beigelii6FSN, PB, TH, S/E PSH, Ar $ -$ Trichosporon pullulans <td>Candida zeylanoides</td> <td>3</td> <td>MLB, F, N to S/E PSH</td> <td>_</td> <td>-</td>	Candida zeylanoides	3	MLB, F, N to S/E PSH	_	-	
Cryptococcus neoformans4MLB, NFCryptococcus labidus1MLB, NFCryptococcus laurentii1MLB, NFCryptococcus lauteohus2MLB, NFCryptococcus lauteohus1MLB, NFCryptococcus lauteohus1MLB, NFCryptococcus terreus1MLB, NFCryptococcus terreus1MLB, NFDebaryomyces spp.1PB, FGeotrichum candidum7FSN, TH, ArHansenula anomala6MLB, F, N to S/E PSHHansenula polymorpha11MLB, NFPrototheca wickerhamii3ECRhodotorula glutinis15MLB, NFRhodotorula glutinis1PB, NFRhodotorula ninuta3MLB, NFSaccharomyces cervisiae12MLB, F, N to S PSHTorulopsis candida2MLB, F, N or S PSH(1) A ≥ 36 Torulopsis pintolopesii1MLB, F, N or S PSHTrichosporon pugliulans1FSN, PB, TH, S/E PSH, Ar	Candida spp. not otherwise identified	5	MLB, PSH			
Cryptococcus albidus1MLB, NFCryptococcus luteolus1MLB, NFCryptococcus luteolus2MLB, NFCryptococcus tereus1MLB, NFCryptococcus uniguttulatus1MLB, NFDebaryomyces spp.1PB, FGeotrichum candidum7FSN, TH, ArHansenula anomala6MLB, F, N to S/E PSHHansenula anomala6MLB, E PSHPichia ohmeri4MLB, NFPrototheca wickerhamii3ECRhodotorula glutinis15MLB, NFRhodotorula pilimanae1PB, NFRhodotorula pilimanae1B, NFSaccharomyces cerevisiae12MLB, NFTorulopsis candida2MLB, NFTorulopsis i untila3B, NFTorulopsis condida1MLB, F, N or S PSHTrichosporon pullulans1FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar	Cryptococcus neoformans	4	MLB, NF	_	-	
Gyptococcus laurentii1MLB, NFCryptococcus luteolus2MLB, NFCryptococcus uniguttulatus1MLB, NFCryptococcus uniguttulatus1MLB, NFDebaryomyces spp.1PB, FGeotrichum candidum7FSN, TH, ArHansenula anomala6MLB, NFHansenula anomala6MLB, F, N to S/E PSHPichia ohmeri4MLB, NFPrototheca wickerhamii3ECRhodotorula glutinis15MLB, NFRhodotorula glutinis15MLB, NFRhodotorula pilimanae1PB, NFRhodotorula pilimanae2MLB, NFSaccharomyces cerevisiae12MLB, NFSaccharomyces cerevisiae12MLB, NFTorulopsis candida2MLB, F, N or S PSHTrichosporon beigelii6FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar	Cryptococcus albidus	1	MLB, NF	_	-	
Cryptococcus luteolus2MLB, NFCryptococcus terreus1MLB, NFCryptococcus uniguttulatus1MLB, NFDebaryomyces spp.1PB, FCeotrichum candidum7FSN, TH, ArHansenula anomala6MLB, F, N to S/E PSHHansenula polymorpha11MLB, NFPrototheca wickerhamii3ECRhodotorula glutinis15MLB, NFRhodotorula glutinia3MLB, NFRhodotorula pilimanae1PB, NFRhodotorula pilimanae1PB, NFRhodotorula pilimanae1PB, NFSaccharomyces cerevisiae12MLB, F, N to S PSHTorulopsis candida2MLB, F, N or S PSHTrichosporon beigelii6FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar	Cryptococcus laurentii	1	MLB, NF	_	-	
Cryptococcus terreus1MLB, NFCryptococcus uniguttulatus1MLB, NFDebaryomyces spp.1PB, FGeotrichum candidum7FSN, TH, ArHansenula anomala6MLB, F, N to S/E PSHHansenula polymorpha11MLB, NFPichia ohmeri4MLB, E PSHPrototheca wickerhamii3ECRhodotorula glutinis15MLB, NFRhodotorula pilimanae1PB, NFRhodotorula pilimanae1PB, NFSaccharomyces cerevisiae12MLB, F, N to S PSHTorulopsis pintolopesii1MLB, F, N or S PSHTrichosporon beigelii6FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar	Cryptococcus luteolus	2	MLB, NF	_	-	
Gryptococcus uniguttulatus1MLB, NFDebaryomyces spp.1PB, FGeotrichum candidum7FSN, TH, ArHansenula anomala6MLB, F, N to S/E PSHHansenula polymorpha11MLB, NFPichia ohmeri4MLB, E PSHPrototheca wickerhamii3ECRhodotorula glutinis15MLB, NFRhodotorula glutinis15MLB, NFRhodotorula pilimanae1PB, NFRhodotorula rubra2MLB, NFSaccharomyces cerevisiae12MLB, F, N to S PSHTorulopsis candida2MLB, F, N or S PSH(1) A ≥ 36 Torulopsis pintolopesii1MLB, F, N or S PSH, ArTrichosporon beigelii6FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar	Cryptococcus terreus	1	MLB, NF	_	-	
Debaryomyces spp.1PB, FGeotrichum candidum7FSN, TH, ArHansenula anomala6MLB, F, N to S/E PSHHansenula polymorpha11MLB, NFPichia ohmeri4MLB, E PSHPrototheca wickerhamii3ECRhodotorula glutinis15MLB, NFRhodotorula glutinis1PB, NFRhodotorula pilimanae1PB, NFRhodotorula rubra2MLB, NFSaccharomyces cerevisiae12MLB, F, N to S PSHTorulopsis candida2MLB, F, N or S PSHTrichosporon beigelii6FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar	Cryptococcus uniguttulatus	1	MLB, NF	_	-	
Geotrichum candidum7FSN, TH, Ar $ -$ Hansenula anomala6MLB, F, N to S/E PSH $ -$ Hansenula polymorpha11MLB, NF $ -$ Pichia ohmeri4MLB, E PSH $ -$ Prototheca wickerhamii3EC $ -$ Rhodotorula glutinis15MLB, NF $ -$ Rhodotorula glutinis15MLB, NF $ -$ Rhodotrula minuta3MLB, NF $ -$ Rhodotorula rubra2MLB, NF $ -$ Rhodotrula rubra2MLB, NF $ -$ Saccharomyces cerevisiae12MLB, F, N to S PSH $ -$ Torulopsis candida2MLB, F, N or S PSH $ -$ Trichosporon beigelii6FSN, PB, TH, S/E PSH, Ar $ -$ Trichosporon pullulans1FSN, PB, TH, S/E PSH, Ar $ -$	Debaryomyces spp.	1	PB, F	_	-	
Hansenula anomala6MLB, F, N to S/E PSH $ -$ Hansenula polymorpha11MLB, NF $ -$ Pichia ohmeri4MLB, E PSH $ -$ Prototheca wickerhamii3EC $ -$ Rhodotorula glutinis15MLB, NF $ -$ Rhodotorula glutinis15MLB, NF $ -$ Rhodotrula minuta3MLB, NF $ -$ Rhodotorula rubra2MLB, NF $ -$ Rhodotrula rubra2MLB, NF $ -$ Saccharomyces cerevisiae12MLB, F, N to S PSH $ -$ Torulopsis candida2MLB, F, N or S PSH $ -$ Trichosporon beigelii6FSN, PB, TH, S/E PSH, Ar $ -$ Trichosporon pullulans1FSN, PB, TH, S/E PSH, Ar $ -$	Geotrichum candidum	7	FSN, TH, Ar	_	-	
Hansenula polymorpha11MLB, NF $ -$ Pichia ohmeri4MLB, E PSH $ -$ Prototheca wickerhamii3EC $ -$ Rhodotorula glutinis15MLB, NF $ -$ Rhodotorula minuta3MLB, NF $ -$ Rhodotorula pilimanae1PB, NF $ -$ Rhodotrula rubra2MLB, NF $ -$ Saccharomyces cerevisiae12MLB, F, N to S PSH $ -$ Torulopsis candida2MLB, F, N or S PSH $ -$ Trichosporon beigelii6FSN, PB, TH, S/E PSH, Ar $ -$ Trichosporon pullulans1FSN, PB, TH, S/E PSH, Ar $ -$	Hansenula anomala	6	MLB, F, N to S/E PSH	_	-	
Pichia ohmeri4MLB, E PSH $ -$ Prototheca wickerhamii3EC $ -$ Rhodotorula glutinis15MLB, NF $ -$ Rhodotorula minuta3MLB, NF $ -$ Rhodotorula pilimanae1PB, NF $ -$ Rhodotorula rubra2MLB, NF $ -$ Saccharomyces cerevisiae12MLB, F, N to S PSH $ -$ Torulopsis candida2MLB, F, N or S PSH $ -$ Torulopsis pintolopesii1MLB, F, N or S PSH $ -$ Trichosporon beigelii6FSN, PB, TH, S/E PSH, Ar $ -$ Trichosporon pullulans1FSN, PB, TH, S/E PSH, Ar $ -$	Hansenula polymorpha	11	MLB, NF	_	_	
Prototheca wickerhamii3EC $ -$ Rhodotorula glutinis15MLB, NF $ -$ Rhodotorula minuta3MLB, NF $ -$ Rhodotorula pilimanae1PB, NF $ -$ Rhodotorula rubra2MLB, NF $ -$ Saccharomyces cerevisiae12MLB, F, N to S PSH $ -$ Torulopsis candida2MLB, F, N or S PSH $ -$ Torulopsis pintolopesii1MLB, F, N or S PSH $ -$ Trichosporon beigelii6FSN, PB, TH, S/E PSH, Ar $ -$ Trichosporon pullulans1FSN, PB, TH, S/E PSH, Ar $ -$	Pichia ohmeri	4	MLB, E PSH	_	-	
Rhodotorula glutinis15MLB, NFRhodotorula minuta3MLB, NFRhodotorula pilimanae1PB, NFRhodotorula rubra2MLB, NFSaccharomyces cerevisiae12MLB, F, N to S PSHTorulopsis candida2MLB, F, N or S PSH(1) A≥36Torulopsis pintolopesii1MLB, F, N or S PSHTrichosporon beigelii6FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar	Prototheca wickerhamii	3	EC	_	-	
Rhodotorula minuta3MLB, NFRhodotorula pilimanae1PB, NFRhodotorula rubra2MLB, NFSaccharomyces cerevisiae12MLB, F, N to S PSHTorulopsis candida2MLB, F, N or S PSH(1) A≥36Torulopsis pintolopesii1MLB, F, N or S PSHTrichosporon beigelii6FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar	Rhodotorula glutinis	15	MLB, NF	_	-	
Rhodotorula pilimanae1PB, NFRhodotorula rubra2MLB, NFSaccharomyces cerevisiae12MLB, F, N to S PSHTorulopsis candida2MLB, F, N or S PSH(1) A≥36Torulopsis pintolopesii1MLB, F, N or S PSHTrichosporon beigelii6FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar	Rhodotorula minuta	3	MLB, NF	_	-	
Rhodotorula rubra2MLB, NFSaccharomyces cerevisiae12MLB, F, N to S PSHTorulopsis candida2MLB, F, N or S PSH(1) A≥36Torulopsis pintolopesii1MLB, F, N or S PSHTrichosporon beigelii6FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar	Rhodotorula pilimanae	1	PB, NF	_	-	
Saccharomyces cerevisiae12MLB, F, N to S PSHTorulopsis candida2MLB, F, N or S PSH(1) A≥36Torulopsis pintolopesii1MLB, F, N or S PSHTrichosporon beigelii6FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar	Rhodotorula rubra	2	MLB, NF	_	-	
Torulopsis candida2MLB, F, N or S PSH(1) A≥36Torulopsis pintolopesii1MLB, F, N or S PSHTrichosporon beigelii6FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar	Saccharomyces cerevisiae	12	MLB, F, N to S PSH	_	-	
Torulopsis pintolopesii1MLB, F, N or S PSHTrichosporon beigelii6FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar	Torulopsis candida	2	MLB, F, N or S PSH	(1) A	≥36	
Trichosporon beigelii6FSN, PB, TH, S/E PSH, ArTrichosporon pullulans1FSN, PB, TH, S/E PSH, Ar-	Torulopsis pintolopesii	1	MLB, F, N or S PSH	—	-	
Trichosporon pullulans 1 FSN, PB, TH, S/E PSH, Ar – –	Trichosporon beigelii	6	FSN, PB, TH, S/E PSH, Ar	—	-	
	Trichosporon pullulans	1	FSN, PB, TH, S/E PSH, Ar	-	-	

TABLE 2.	Determination	of the	ability c	f clinical	yeast isol	ates to	ferment	trehalose	at 42°C	and their	concomitant
			mor	phology	on cornme	al Twe	en 80 ag	ar ^a			

^{*a*} CMA, cornneal agar; MLB, multiple lateral budding; N, no/none; F, filaments; S, simple; E, elaborate; PSH, pseudohyphae; TH, true or septate hyphae; PB, polar budding; Ar, arthroconidia; EC, endocytoplasmic cleavage; FSN, vegetative reproduction by splitting. Example: MLB, F, N to S/E PSH = multiple lateral budding; filaments, none to simple or elaborate pseudohyphae. A, acid production; G, gas production.

^b Numbers in parentheses reflect the number of isolates positive for trehalose fermentation at 42°C. *T. glabrata* was within 24 h; other yeasts were \geq 36 h.

^c Germ tube-negative C. albicans clinical isolates that were referred for further identification.

went on to be identified as *T. glabrata* by API. Each of these strains grew poorly on Sabouraud dextrose agar, the trehalose screening medium, and the API 20C basal medium supplemented with trehalose, 5 to 7 days being taken to identify them as *T. glabrata* in the last. These patients were on cytotoxic therapy for their primary diseases or had finished a course of antifungal therapy for another organism.

Although excellent commercial systems now exist for the relatively rapid identification of yeasts, the cost per identification is high for small- to modest-volume laboratories (\geq \$4.00 to \leq \$8.00). The total hands-on time involved in inoculating and reading the commercial panels is also significant. Thus, the major problem of yeast identification in this era of cost containment is not the lack of technical expertise experienced 10 years ago, but rather the cost of reagents and personnel. This challenge has been met by devising screening strategies for the more common yeasts. Precedence for the screening approach was the germ tube test developed by Reynolds and Braude (19), a mainstay for over 40 years in the clinical laboratory. It permits the laboratory to rapidly dismiss approximately 70% of

its work at less than \$2.00 per test, and its efficacy is not diluted by the fact that *Candida stellatoidea* (a sucrose-negative variant of *C. albicans*) is also germ tube positive and not all *C. albicans* strains are germ tube positive. A biochemical alternative to the germ tube test appears to be even more sensitive at about the same cost (17). Similar screening tests are available for *Cryptococcus neoformans* (birdseed agar, caffeic acid) and *C. parapsilosis* (β-glucosidase), and rapid urease and nitrate tests have been developed for screening large numbers of respiratory specimens for possible pathogenic yeasts (9, 13, 25).

In summary, *T. glabrata* and *C. tropicalis* were the only yeasts among our clinical isolates that fermented trehalose at 42°C. The nonfilamentous morphology of *T. glabrata* combined with high-temperature trehalose fermentation within 24 h readily differentiated it from *C. tropicalis* and the few other filamentous yeasts that had delayed acid production only. The trehalose screen for *T. glabrata* offers a more rapid and cost-effective alternative to commercial identification systems and takes about one-fifth of the setting-up and reading time of the more expensive panels. An experienced laboratorian can look at the colonial morphology of an unknown yeast on a Columbia CNA agar or blood agar plate, inoculate a trehalose tube and morphology medium, and have the results by the next morning (for some strains, within the same day). This is in contrast to the 3-day incubation period for the API 20C, and while the Dade Microscan Yeast Identification Panel provides an answer in 4 to 6 h, panels are more costly than the tubes (approximately \$1.50) and a 24- to 48-h growth plate is always required for the inoculum. Savings in media and time are diluted slightly if a growth plate has to be inoculated, but the overall savings compared with commercial identification systems are still significant. Another advantage of 42°C trehalose fermentation is that it is not used to discriminate between organisms having good likelihood but low selectivity identification or rare biotypes in commercial databases. These are yeast-like fungi that metabolize very few substrates with much strain variability such as Blastoschizomyces (Trichosporon) capitatus, Candida krusei, Candida lambica, Hanseniaspora valbyensis, and S. cerevisiae and are incapable of growing at 42°C, fermenting trehalose, or both (Table 1). Considering the data presented herein, we feel that a test consisting of appropriate morphology and fermentation of trehalose at 42°C effectively differentiates T. glabrata from other yeasts and has the potential not only to significantly reduce laboratory workload, but also to generate substantial savings in direct and indirect costs to the laboratory.

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