

# High resolution melting analysis for rapid identification of twenty-five *Candida* species from culture



eP380

E. Němcová<sup>1, 2</sup>, M. Černochová<sup>1</sup>, F. Růžička<sup>3</sup>, B. Mališová<sup>1</sup>, M. Vaněrková<sup>1</sup>,  
H. Šuranská<sup>1</sup>, P. Němec<sup>1, 2</sup>, T. Freiburger<sup>1</sup>



<sup>1</sup> Molecular Genetics Laboratory, Centre for Cardiovascular Surgery and Transplantation, Brno, Czech Republic

<sup>2</sup> St. Anne's University Hospital - International Clinical Research Center, Brno, Czech Republic

<sup>3</sup> Microbiological Institute, Faculty of Medicine, Masaryk University and St. Anne's University Hospital, Brno, Czech Republic

## Introduction

Although *Candida albicans* remains the most common fungal isolate from blood, many studies have described an increasing trend in non-*albicans* infections (Krcmery and Barnes 2002, Leroy *et al.* 2009). Correct identification of *Candida* species is important for targeted antifungal therapy and for epidemiological purposes. The aim of this study was to identify 25 *Candida* sp. from culture using high resolution melting analysis (HRMA).

## Materials and Methods

We have developed real-time PCR followed by HRMA for differentiation of 25 *Candida* species. The assay was performed with the reference collections strains confirmed by ITS2 sequencing (Table 1).

## Results

Real-time PCR with HRMA was performed with *Candida* reference strains in duplicates. Species could be identified by comparing the shapes of the melting curves and values of melting temperatures (Tm) with reference list (Figure 1, Table 1). Mean Tm  $\pm$  3SD values together with the shape of the melting curves were conclusive in 16 *Candida* species, while normalized temperature - shifted difference plot against *C. krusei* had to be used to distinguish other 5 species. Only Tm values of two pairs of *Candida* sp.: *C. fabianii*/*C. guilliermondii* and *C. orthopsilosis*/*C. metapsilosis* occurred repeatedly in the narrow ranges, therefore these species within 2 groups could not be reliably distinguished.

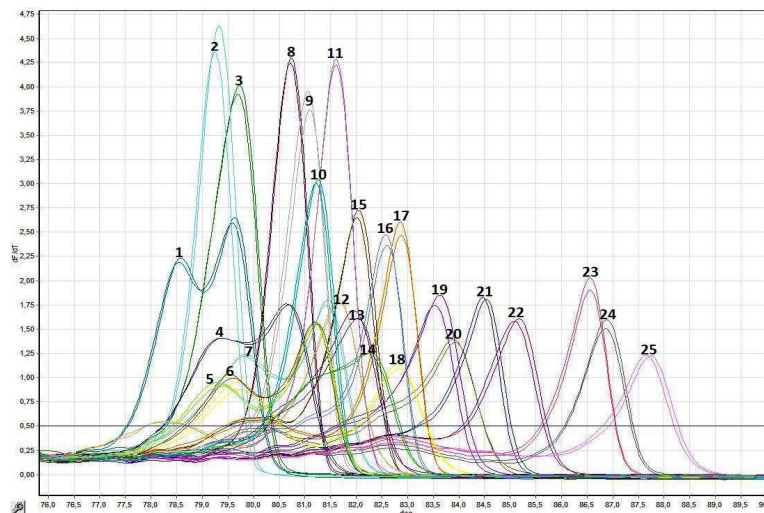


Figure 1: Melting curves of reference *Candida* species:

1) *C. pelliculosa*, 2) *C. zeylanoides*, 3) *C. saitoana*, 4) *C. tropicalis*, 5) *C. metapsilosis*, 6) *C. orthopsilosis*, 7) *C. parapsilosis*, 8) *C. utilis*, 9) *C. fabianii*, 10) *C. guilliermondii*, 11) *C. glabrata*, 12) *C. lipolytica*, 13) *C. intermedia*, 14) *C. pulcherrima*, 15) *C. dubliniensis*, 16) *C. kefyr*, 17) *C. albicans*, 18) *C. catenulata*, 19) *C. lusitanae*, 20) *C. rugosa*, 21) *C. inconspicua*, 22) *C. californica*, 23) *C. krusei*, 24) *C. norvegensis*, 25) *C. lambica*

## Conclusions

Real-time PCR followed by HRMA is a simple, rapid and inexpensive tool to identify *Candida* species. It seems to be a suitable complement to current clinical diagnostic approach based on usage of commercially available biochemical kits. This method should be further validated using clinical isolates.

**References:** •Krcmery V., Barnes A.J. Non-albicans *Candida* spp. causing fungaemia: pathogenicity and antifungal resistance. J Hosp Infect. 2002; 50(4):243-60.  
•Leroy O., Gangneux J.P., Montravers P., *et al.* Epidemiology, management, and risk factors for death of invasive *Candida* infections in critical care: A multicentre, prospective, observational study in France (2005-2006). Crit Care Med. 2009; 37(5):1612-8.

species	reference strain No.	mean Tm (SD) °C	
		peak 1	peak 2
<i>C. albicans</i>	CCM 8320	80,098 (0,048)	82,938 (0,085)
<i>C. glabrata</i>	CCM 8270	81,705 (0,070)	
<i>C. parapsilosis</i>	CCM 8260	79,863 (0,071)	81,508 (0,061)
<i>C. tropicalis</i>	CCM 8264	79,433 (0,061)	80,738 (0,096)
<i>C. krusei</i> *	CCY 29-9-17	86,605 (0,043)	
<i>C. guilliermondii</i>	CCY 39-23-6	81,255 (0,039)	
<i>C. fabianii</i>	CCY 38-20-1	81,083 (0,051)	
<i>C. utilis</i> *	CCY 29-38-74	80,720 (0,073)	
<i>C. inconspicua</i> *	CCY 26-26-11	84,603 (0,112)	
<i>C. dubliniensis</i>	CCY 29-177-1	80,200 (0,064)	82,090 (0,089)
<i>C. pelliculosa</i>	CCY 29-6-7	78,625 (0,060)	79,683 (0,065)
<i>C. saitoana</i>	CCY 26-9-9	79,695 (0,052)	
<i>C. lusitanae</i>	CCY 29-59-1	83,663 (0,079)	
<i>C. lipolytica</i>	CCY 29-26-42	78,325 (0,139)	81,710 (0,052)
<i>C. rugosa</i>	CCY 29-15-1	83,980 (0,079)	
<i>C. norvegensis</i> *	CCY 29-47-2	86,930 (0,087)	
<i>C. catenulata</i>	CCY 29-17-3	79,730 (0,067)	82,850 (0,028)
<i>C. intermedia</i>	CCY 29-12-10	81,968 (0,032)	
<i>C. lambica</i>	CCY 29-97-12	87,678 (0,039)	
<i>C. pulcherrima</i>	CCY 29-2-128	82,178 (0,062)	
<i>C. orthopsilosis</i>	MUCL 49939	79,563 (0,037)	81,208 (0,034)
<i>C. metapsilosis</i>	MUCL 46179	79,333 (0,093)	81,213 (0,083)
<i>C. zeylanoides</i>	MUCL 27735	79,328 (0,063)	
<i>C. kefyr</i>	MUCL 29857	82,658 (0,084)	
<i>C. californica</i> *	CCY 29-93-4	85,170 (0,083)	

Table 1: Melting temperatures (Tm) of reference *Candida* species

\* normalized temperature - shifted difference plot against *C. krusei* had to be used to distinguish them