

عنوان مقاله:

Development a hydrolysis probe-based quantitative PCR assay for the specific detection and quantification of *Candida auris*

محل انتشار:

سرطان معده، دوره 6، شماره 3 (سال: 1399)

تعداد صفحات اصل مقاله: 7

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خلاصه مقاله:

Background and Purpose: *Candida auris* is an emerging multidrug-resistant pathogen. The identification of this species with the conventional phenotypic or biochemical mycological methods may lead to misidentification. Molecular-based species-specific identification methods such as quantitative real-time polymerase chain reaction (qPCR) facilitate a more reliable identification of *C. auris* than mycological methods. Regarding this, the present study aimed to develop a hydrolysis probe-based qPCR assay for the rapid, accurate identification of *C. auris*. **Materials and Methods:** The internal transcribed spacer 2 regions in the nuclear ribosomal DNA of *C. auris* and other related yeasts were assayed to find a specific PCR target for *C. auris*. A 123-base-pair target was selected, and primers and a probe were designed for hydrolysis probe-based real-time PCR with TaqMan chemistry. Ten-fold serial dilutions of *C. auris* ranging from 10⁶ to 10⁰ CFU/mL were prepared to establish a standard curve to quantify the yeast. **Results:** The qPCR assay was able to identify and quantify *C. auris* with a detection limit of 1 *C. auris* CFU per reaction. Specificity was confirmed by the non-amplification of the sequences belonging to other *Candida* species, yeasts, molds, bacteria, or human DNAs. The standard curve of the assay showed a highly significant linearity between threshold values and dilution rates ($R^2=0.99$; slope=-3.42). **Conclusion:** The applied qPCR assay facilitated the rapid and accurate identification and quantification of emerging opportunistic *C. auris*. Therefore, considering the promising test validation results, we succeeded to develop a rapid and accurate hydrolysis probe-based qPCR assay for the screening and identification of *C. auris*.

کلمات کلیدی:

Candida auris, Quantification, Real-Time PCR

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