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Laboratory Isolation and Identification of Candida Species Page 5/26

cerning C. albicans and related species. Of course, these studies also emphasized the use of novel methods V that proved indispensable to the particular investigations. More important, many of the protocols are of general interest for use by researchers studying Candida species. Thus, upon being approached again to edit a new volume

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The yeast Candida being the main cause of candidiasis is a Page 10/26

commonly isolated pathogen from immunocompromised patients. Successfully identifying the species of Candida is important in the treatment and management of the disease. The trend in the resistance acquired by some species of Candida leads to the importance of identification to the species level.

Identification of

Candida Species from Clinical Samples in ... DNA fingerprinting Candida species Ogy Claude Pujol and David R. Soll Genomics and Proteomics. ... In "Candida Albicans: Methods and Protocols", expert researchers explore these exciting new insights, focusing on the study of medically important fungi and Candida spp in

particular. Chapters examine critical aspects of molecular methods, providing ...

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candidiasis caused by Candida albicans are described in this unit. Detailed materials and methods for C. albicans growth and detection are also described.

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Here, we are
describing a ChIP
protocol that is

optimized for Candida albicans. The protocol

requires 4–5 days for completion of the assay and has been used to produce robust ChIP results for...

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Molecular identification of Candida species was performed using a widely used method based on PCR-RFLP [21, 23, 24]. The amplification reaction

was directed to the ribosomal region comprising the internal transcribed spacers ITS1 and ITS2 and the 5.8S gene.

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15 Typing Candida Species using Microsatellite Length Polymorphism and Page 19/26

Multilocus Sequence Typing Dea Garcia-Hermoso, Marie Desnos-Ollivier, and Stephane Bretagne; 16 Biology Diagnostic Methods for Detection of Bloodborne Candidiasis Cornelius J. Clancy, and M. Hong Nguyen. (source: Nielsen Book Data) Summary

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discussion of a variety
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poorly known fungal species occasionally involved in human infections. A molecular analysis of the lology sequences of the D1/D2 domains and the internal transcribed spacer (ITS) region of the ribosomal genes of 24 clinical isolates phenotypically identified as C. rugosa demonstrated that only 10 (41.6%) isolates belonged to that species. Page 22/26

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