

# Transcriptional Response of *Candida albicans* upon Inter

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Citation Report

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1	Recent advances in the genomic analysis of <i>Candida albicans</i> . <i>Revista Iberoamericana De Micologia</i> , 2005, 22, 187-193.	0.4	8
2	Use of genome information for the study of the pathogenesis of fungal infections and the development of diagnostic tools. <i>Revista Iberoamericana De Micologia</i> , 2005, 22, 238-241.	0.4	13
4	Comparative Gene Expression Analysis by a Differential Clustering Approach: Application to the <i>Candida albicans</i> Transcription Program. <i>PLoS Genetics</i> , 2005, 1, e39.	1.5	124
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6	The <i>Candida albicans</i> Vacuole Is Required for Differentiation and Efficient Macrophage Killing. <i>Eukaryotic Cell</i> , 2005, 4, 1677-1686.	3.4	56
7	Release from Quorum-Sensing Molecules Triggers Hyphal Formation during <i>Candida albicans</i> Resumption of Growth. <i>Eukaryotic Cell</i> , 2005, 4, 1203-1210.	3.4	133
8	<i>Candida albicans</i> -Conditioned Medium Protects Yeast Cells from Oxidative Stress: a Possible Link between Quorum Sensing and Oxidative Stress Resistance. <i>Eukaryotic Cell</i> , 2005, 4, 1654-1661.	3.4	116
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12	<i>Cryptococcus neoformans</i> Gene Expression during Murine Macrophage Infection. <i>Eukaryotic Cell</i> , 2005, 4, 1420-1433.	3.4	184
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15	Global analysis of altered gene expression during morphogenesis of <i>Candida albicans</i> in vitro. <i>Biochemical and Biophysical Research Communications</i> , 2005, 334, 1149-1158.	1.0	22
16	<i>Candida albicans</i> protein analysis during hyphal differentiation using an integrative HA-tagging method. <i>Biochemical and Biophysical Research Communications</i> , 2005, 337, 784-790.	1.0	10
17	<i>Candida albicans</i> biofilm development, modeling a host-pathogen interaction. <i>Current Opinion in Microbiology</i> , 2006, 9, 340-345.	2.3	190
18	Transmembrane domain prediction and consensus sequence identification of the oligopeptide transport family. <i>Research in Microbiology</i> , 2006, 157, 395-406.	1.0	17
20	Microarrays for Studying Pathogenicity in <i>Candida Albicans</i> . , 0, , 181-209.		18

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649	<i>Escherichia coli</i> , but Not <i>Staphylococcus aureus</i> , Functions as a Chelating Agent That Exhibits Antifungal Activity against the Pathogenic Yeast <i>Candida albicans</i> . <i>Journal of Fungi (Basel, Switzerland)</i> , 2023, 9, 11.	1.5	9
650	“Under Pressure” How fungi evade, exploit, and modulate cells of the innate immune system. <i>Seminars in Immunology</i> , 2023, 66, 101738.	2.7	2
651	The F <sub>1</sub> F <sub>o</sub> -ATP synthase $\epsilon$ subunit of <i>Candida albicans</i> induces inflammatory responses by controlling amino acid catabolism. <i>Virulence</i> , 2023, 14, .	1.8	2
652	RAD51-dependent genetic pathways are essential for DNA-protein crosslink repair and pathogenesis in <i>Candida albicans</i> . <i>Journal of Biological Chemistry</i> , 2023, 299, 104728.	1.6	3
653	<i>Candida albicans</i> Hyphal Morphogenesis within Macrophages Does Not Require Carbon Dioxide or pH-Sensing Pathways. <i>Infection and Immunity</i> , 2023, 91, .	1.0	4