

Melanie Legrand

List of Publications by Year in descending order

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18
papers

702
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759233

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839539

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21
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21
docs citations

21
times ranked

820
citing authors

#	ARTICLE	IF	CITATIONS
1	Overexpression approaches to advance understanding of <i>Candida albicans</i> . <i>Molecular Microbiology</i> , 2022, 117, 589-599.	2.5	12
2	Multiple Stochastic Parameters Influence Genome Dynamics in a Heterozygous Diploid Eukaryotic Model. <i>Journal of Fungi (Basel, Switzerland)</i> , 2022, 8, 650.	3.5	1
3	Factors that influence bidirectional long-tract homozygosis due to double-strand break repair in <i>Candida albicans</i> . <i>Genetics</i> , 2021, 218, .	2.9	1
4	Use of CRISPR-Cas9 To Target Homologous Recombination Limits Transformation-Induced Genomic Changes in <i>Candida albicans</i> . <i>MSphere</i> , 2020, 5, .	2.9	10
5	Identification of Recessive Lethal Alleles in the Diploid Genome of a <i>Candida albicans</i> Laboratory Strain Unveils a Potential Role of Repetitive Sequences in Buffering Their Deleterious Impact. <i>MSphere</i> , 2019, 4, .	2.9	5
6	<i>Candida albicans</i> : An Emerging Yeast Model to Study Eukaryotic Genome Plasticity. <i>Trends in Genetics</i> , 2019, 35, 292-307.	6.7	35
7	Generating genomic platforms to study <i>Candida albicans</i> pathogenesis. <i>Nucleic Acids Research</i> , 2018, 46, 6935-6949.	14.5	30
8	Genome Diversity and Dynamics in <i>Candida albicans</i> . , 2017, , 205-232.		4
9	Analysis of Repair Mechanisms following an Induced Double-Strand Break Uncovers Recessive Deleterious Alleles in the <i>Candida albicans</i> Diploid Genome. <i>MBio</i> , 2016, 7, .	4.1	31
10	A FACS-Optimized Screen Identifies Regulators of Genome Stability in <i>Candida albicans</i> . <i>Eukaryotic Cell</i> , 2015, 14, 311-322.	3.4	19
11	A study of the <i>DNA</i> damage checkpoint in <i>Candida albicans</i> : uncoupling of the functions of <i>Rad53</i> in <i>DNA</i> repair, cell cycle regulation and genotoxic stress-induced polarized growth. <i>Molecular Microbiology</i> , 2014, 91, 452-471.	2.5	39
12	A Versatile Overexpression Strategy in the Pathogenic Yeast <i>Candida albicans</i> : Identification of Regulators of Morphogenesis and Fitness. <i>PLoS ONE</i> , 2012, 7, e45912.	2.5	103
13	Modular Gene Over-expression Strategies for <i>Candida albicans</i> . <i>Methods in Molecular Biology</i> , 2012, 845, 227-244.	0.9	18
14	The contribution of the S-phase checkpoint genes <i>MEC1</i> and <i>SGS1</i> to genome stability maintenance in <i>Candida albicans</i> . <i>Fungal Genetics and Biology</i> , 2011, 48, 823-830.	2.1	28
15	Analysis of base excision and nucleotide excision repair in <i>Candida albicans</i> . <i>Microbiology (United Kingdom)</i> 157, 187-195. doi:10.1099/mic/0/000000.0	1.8	35
16	Haplotype Mapping of a Diploid Non-Meiotic Organism Using Existing and Induced Aneuploidies. <i>PLoS Genetics</i> , 2008, 4, e1.	3.5	129
17	Role of DNA Mismatch Repair and Double-Strand Break Repair in Genome Stability and Antifungal Drug Resistance in <i>Candida albicans</i> . <i>Eukaryotic Cell</i> , 2007, 6, 2194-2205.	3.4	95
18	Homozygosity at the MTL locus in clinical strains of <i>Candida albicans</i> : karyotypic rearrangements and tetraploid formation. <i>Molecular Microbiology</i> , 2004, 52, 1451-1462.	2.5	104