

Judith G Berman

List of Publications by Year in descending order

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164
papers

15,623
citations

20817

60
h-index

20358

116
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226
all docs

226
docs citations

226
times ranked

10659
citing authors

#	ARTICLE	IF	CITATIONS
1	Microbial communities form rich extracellular metabolomes that foster metabolic interactions and promote drug tolerance. <i>Nature Microbiology</i> , 2022, 7, 542-555.	13.3	58
2	Tackling the emerging threat of antifungal resistance to human health. <i>Nature Reviews Microbiology</i> , 2022, 20, 557-571.	28.6	311
3	An orthologous gene coevolution network provides insight into eukaryotic cellular and genomic structure and function. <i>Science Advances</i> , 2022, 8, eabn0105.	10.3	10
4	Impact of tolerance to fluconazole on treatment response in <i>Candida albicans</i> bloodstream infection. <i>Mycoses</i> , 2021, 64, 78-85.	4.0	29
5	Combining Colistin and Fluconazole Synergistically Increases Fungal Membrane Permeability and Antifungal Cidalty. <i>ACS Infectious Diseases</i> , 2021, 7, 377-389.	3.8	17
6	The fitness costs and benefits of trisomy of each <i>Candida albicans</i> chromosome. <i>Genetics</i> , 2021, 218, .	2.9	35
7	Adaptive Resistance Mutations at Suprainhibitory Concentrations Independent of SOS Mutagenesis. <i>Molecular Biology and Evolution</i> , 2021, 38, 4095-4115.	8.9	6
8	Tunicamycin Potentiates Antifungal Drug Tolerance via Aneuploidy in <i>Candida albicans</i> . <i>MBio</i> , 2021, 12, e0227221.	4.1	22
9	Adenosine Triphosphate Released by <i>Candida albicans</i> Is Associated with Reduced Skin Infectivity. <i>Journal of Investigative Dermatology</i> , 2021, 141, 2306-2310.	0.7	2
10	Adaptation to Fluconazole via Aneuploidy Enables Cross-Adaptation to Amphotericin B and Flucytosine in <i>Cryptococcus neoformans</i> . <i>Microbiology Spectrum</i> , 2021, 9, e0072321.	3.0	19
11	Multifactorial Mechanisms of Tolerance to Ketoconazole in <i>Candida albicans</i> . <i>Microbiology Spectrum</i> , 2021, 9, e0032121.	3.0	18
12	Aneuploidy Underlies Tolerance and Cross-Tolerance to Drugs in <i>Candida parapsilosis</i> . <i>Microbiology Spectrum</i> , 2021, 9, e0050821.	3.0	14
13	Comparing the utility of in vivo transposon mutagenesis approaches in yeast species to infer gene essentiality. <i>Current Genetics</i> , 2020, 66, 1117-1134.	1.7	13
14	<i>Candida albicans</i> Genetic Background Influences Mean and Heterogeneity of Drug Responses and Genome Stability during Evolution in Fluconazole. <i>MSphere</i> , 2020, 5, .	2.9	28
15	Combination of Miconazole and Domiphen Bromide Is Fungicidal against Biofilms of Resistant <i>Candida</i> spp. <i>Antimicrobial Agents and Chemotherapy</i> , 2020, 64, .	3.2	13
16	Evaluation of Microsatellite Typing, ITS Sequencing, AFLP Fingerprinting, MALDI-TOF MS, and Fourier-Transform Infrared Spectroscopy Analysis of <i>Candida auris</i> . <i>Journal of Fungi (Basel)</i> , 2020, 6, 137-145.	1.0	10
17	Elevated Vacuolar Uptake of Fluorescently Labeled Antifungal Drug Caspofungin Predicts Echinocandin Resistance in Pathogenic Yeast. <i>ACS Central Science</i> , 2020, 6, 1698-1712.	11.3	15
18	Identification of Essential Genes and Fluconazole Susceptibility Genes in <i>Candida glabrata</i> by Profiling <i>Hermes</i> Transposon Insertions. <i>G3: Genes, Genomes, Genetics</i> , 2020, 10, 3859-3870.	1.8	27

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19	Drug resistance and tolerance in fungi. <i>Nature Reviews Microbiology</i> , 2020, 18, 319-331.	28.6	342
20	Reply to Balsa-Canto et al.: Growth models are applicable to growth data, not to stationary-phase data. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 814-815.	7.1	6
21	Peptide Self-Assembly Is Linked to Antibacterial, but Not Antifungal, Activity of Histatin 5 Derivatives. <i>MSphere</i> , 2020, 5, .	2.9	5
22	Combining Miconazole and Domiphen Bromide Results in Excess of Reactive Oxygen Species and Killing of Biofilm Cells. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 617214.	3.7	10
23	Anidulafungin Susceptibility Testing of <i>Candida glabrata</i> Isolates from Blood Cultures by the MALDI Biotyper Antibiotic (Antifungal) Susceptibility Test Rapid Assay. <i>Antimicrobial Agents and Chemotherapy</i> , 2019, 63, .	3.2	17
24	Chromatin Profiling of the Repetitive and Nonrepetitive Genomes of the Human Fungal Pathogen <i>Candida albicans</i> . <i>MBio</i> , 2019, 10, .	4.1	19
25	Predicting microbial growth in a mixed culture from growth curve data. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 14698-14707.	7.1	102
26	<i>Candida auris</i> Identification and Rapid Antifungal Susceptibility Testing Against Echinocandins by MALDI-TOF MS. <i>Frontiers in Cellular and Infection Microbiology</i> , 2019, 9, 20.	3.9	48
27	Selection of <i>Candida albicans</i> trisomy during oropharyngeal infection results in a commensal-like phenotype. <i>PLoS Genetics</i> , 2019, 15, e1008137.	3.5	43
28	Aneuploidy Enables Cross-Adaptation to Unrelated Drugs. <i>Molecular Biology and Evolution</i> , 2019, 36, 1768-1782.	8.9	75
29	Autonomously Replicating Linear Plasmids That Facilitate the Analysis of Replication Origin Function in <i>Candida albicans</i> . <i>MSphere</i> , 2019, 4, .	2.9	14
30	Dynamic ploidy changes drive fluconazole resistance in human cryptococcal meningitis. <i>Journal of Clinical Investigation</i> , 2019, 129, 999-1014.	8.2	112
31	Localizing Antifungal Drugs to the Correct Organelle Can Markedly Enhance their Efficacy. <i>Angewandte Chemie</i> , 2018, 130, 6338-6343.	2.0	10
32	Localizing Antifungal Drugs to the Correct Organelle Can Markedly Enhance their Efficacy. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 6230-6235.	13.8	29
33	Maize Transposable Elements <i>Ac/Ds</i> as Insertion Mutagenesis Tools in <i>Candida albicans</i> . <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 1139-1145.	1.8	22
34	Gene Essentiality Analyzed by <i>In Vivo</i> Transposon Mutagenesis and Machine Learning in a Stable Haploid Isolate of <i>Candida albicans</i> . <i>MBio</i> , 2018, 9, .	4.1	110
35	Antifungal tolerance is a subpopulation effect distinct from resistance and is associated with persistent candidemia. <i>Nature Communications</i> , 2018, 9, 2470.	12.8	175
36	Living Bacteria in Thermoresponsive Gel for Treating Fungal Infections. <i>Advanced Functional Materials</i> , 2018, 28, 1801581.	14.9	29

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37	Methodologies for in vitro and in vivo evaluation of efficacy of antifungal and antibiofilm agents and surface coatings against fungal biofilms. <i>Microbial Cell</i> , 2018, 5, 300-326.	3.2	81
38	Rapid Phenotypic and Genotypic Diversification After Exposure to the Oral Host Niche in <i>Candida albicans</i> . <i>Genetics</i> , 2018, 209, 725-741.	2.9	82
39	<i>Candida albicans</i> Dispersed Cells Are Developmentally Distinct from Biofilm and Planktonic Cells. <i>MBio</i> , 2018, 9, .	4.1	69
40	Functional diversification accompanies gene family expansion of MED2 homologs in <i>Candida albicans</i> . <i>PLoS Genetics</i> , 2018, 14, e1007326.	3.5	25
41	Haplotyping a Non-meiotic Diploid Fungal Pathogen Using Induced Aneuploidies and SNP/CGH Microarray Analysis. <i>Methods in Molecular Biology</i> , 2017, 1551, 131-146.	0.9	1
42	Ploidy tug-of-war: Evolutionary and genetic environments influence the rate of ploidy drive in a human fungal pathogen. <i>Evolution; International Journal of Organic Evolution</i> , 2017, 71, 1025-1038.	2.3	42
43	Real-Time Imaging of the Azole Class of Antifungal Drugs in Live <i>Candida</i> Cells. <i>ACS Chemical Biology</i> , 2017, 12, 1769-1777.	3.4	57
44	Phenotypic and genotypic characteristics of <i>Candida albicans</i> isolates from bloodstream and mucosal infections. <i>Mycoses</i> , 2017, 60, 534-545.	4.0	12
45	Generation of Fluorescent Protein Fusions in <i>Candida</i> Species. <i>Journal of Visualized Experiments</i> , 2017, , .	0.3	6
46	Adaptive Mistranslation Accelerates the Evolution of Fluconazole Resistance and Induces Major Genomic and Gene Expression Alterations in <i>Candida albicans</i> . <i>MSphere</i> , 2017, 2, .	2.9	29
47	Multidrug-Resistant <i>Candida haemulonii</i> and <i>C. auris</i> , Tel Aviv, Israel. <i>Emerging Infectious Diseases</i> , 2017, 23, .	4.3	224
48	Assessment of <i>Candida auris</i> Response to Antifungal Drugs Using Time-Kill Assays and an Animal Model. <i>Open Forum Infectious Diseases</i> , 2017, 4, S73-S73.	0.9	2
49	<i>Candida haemulonii</i> and <i>Candida auris</i> : Emerging Multidrug-Resistant Species With Distinct Virulence and Epidemiological Characteristics. <i>Open Forum Infectious Diseases</i> , 2016, 3, .	0.9	2
50	<i>Candida albicans</i> repetitive elements display epigenetic diversity and plasticity. <i>Scientific Reports</i> , 2016, 6, 22989.	3.3	35
51	Ploidy dynamics and evolvability in fungi. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2016, 371, 20150461.	4.0	46
52	Phenotypic Consequences of a Spontaneous Loss of Heterozygosity in a Common Laboratory Strain of <i>Candida albicans</i> . <i>Genetics</i> , 2016, 203, 1161-1176.	2.9	28
53	Sir2 regulates stability of repetitive domains differentially in the human fungal pathogen <i>Candida albicans</i> . <i>Nucleic Acids Research</i> , 2016, 44, gkw594.	14.5	29
54	Heteroresistance to Fluconazole Is a Continuously Distributed Phenotype among <i>Candida glabrata</i> Clinical Strains Associated with <i>In Vivo</i> Persistence. <i>MBio</i> , 2016, 7, .	4.1	61

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55	High frame-rate resolution of cell division during <i>Candida albicans</i> filamentation. <i>Fungal Genetics and Biology</i> , 2016, 88, 54-58.	2.1	12
56	Ploidy plasticity: a rapid and reversible strategy for adaptation to stress. <i>FEMS Yeast Research</i> , 2016, 16, fow020.	2.3	86
57	diskImageR: quantification of resistance and tolerance to antimicrobial drugs using disk diffusion assays. <i>Microbiology (United Kingdom)</i> , 2016, 162, 1059-1068.	1.8	58
58	Neocentromeres Provide Chromosome Segregation Accuracy and Centromere Clustering to Multiple Loci along a <i>Candida albicans</i> Chromosome. <i>PLoS Genetics</i> , 2016, 12, e1006317.	3.5	34
59	Parasexual Ploidy Reduction Drives Population Heterogeneity Through Random and Transient Aneuploidy in <i>Candida albicans</i> . <i>Genetics</i> , 2015, 200, 781-794.	2.9	98
60	Real-Time Evolution of a Subtelomeric Gene Family in <i>Candida albicans</i> . <i>Genetics</i> , 2015, 200, 907-919.	2.9	36
61	Targeting the Adaptability of Heterogeneous Aneuploids. <i>Cell</i> , 2015, 160, 771-784.	28.9	115
62	<i>Candida albicans</i> Morphology and Dendritic Cell Subsets Determine T Helper Cell Differentiation. <i>Immunity</i> , 2015, 42, 356-366.	14.3	182
63	Telomeric ORFs in <i>Candida albicans</i> : Does Mediator Tail Wag the Yeast?. <i>PLoS Pathogens</i> , 2015, 11, e1004614.	4.7	12
64	Polyloid Titan Cells Produce Haploid and Aneuploid Progeny To Promote Stress Adaptation. <i>MBio</i> , 2015, 6, e01340-15.	4.1	135
65	Shift and adapt: the costs and benefits of karyotype variations. <i>Current Opinion in Microbiology</i> , 2015, 26, 130-136.	5.1	37
66	Physical limits on kinesin-5-mediated chromosome congression in the smallest mitotic spindles. <i>Molecular Biology of the Cell</i> , 2015, 26, 3999-4014.	2.1	11
67	Genetic and phenotypic intra-species variation in <i>Candida albicans</i> . <i>Genome Research</i> , 2015, 25, 413-425.	5.5	305
68	The evolution of drug resistance in clinical isolates of <i>Candida albicans</i> . <i>ELife</i> , 2015, 4, e00662.	6.0	268
69	A Tetraploid Intermediate Precedes Aneuploid Formation in Yeasts Exposed to Fluconazole. <i>PLoS Biology</i> , 2014, 12, e1001815.	5.6	147
70	Silencing Is Noisy: Population and Cell Level Noise in Telomere-Adjacent Genes Is Dependent on Telomere Position and Sir2. <i>PLoS Genetics</i> , 2014, 10, e1004436.	3.5	30
71	Telomeric ORFs (TLOs) in <i>Candida</i> spp. Encode Mediator Subunits That Regulate Distinct Virulence Traits. <i>PLoS Genetics</i> , 2014, 10, e1004658.	3.5	36
72	Origin Replication Complex Binding, Nucleosome Depletion Patterns, and a Primary Sequence Motif Can Predict Origins of Replication in a Genome with Epigenetic Centromeres. <i>MBio</i> , 2014, 5, e01703-14.	4.1	21

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73	YMAP: a pipeline for visualization of copy number variation and loss of heterozygosity in eukaryotic pathogens. <i>Genome Medicine</i> , 2014, 6, 100.	8.2	95
74	Rapid Mechanisms for Generating Genome Diversity: Whole Ploidy Shifts, Aneuploidy, and Loss of Heterozygosity. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2014, 4, a019604-a019604.	6.2	106
75	Y MAP : a pipeline for visualization of copy number variation and loss of heterozygosity in eukaryotic pathogens. <i>Genome Medicine</i> , 2014, 6, 100.	8.2	33
76	The "obligate diploid" <i>Candida albicans</i> forms mating-competent haploids. <i>Nature</i> , 2013, 494, 55-59.	27.8	246
77	Shuttle vectors for facile gap repair cloning and integration into a neutral locus in <i>Candida albicans</i> . <i>Microbiology (United Kingdom)</i> , 2013, 159, 565-579.	1.8	74
78	Monopolin recruits condensin to organize centromere DNA and repetitive DNA sequences. <i>Molecular Biology of the Cell</i> , 2013, 24, 2807-2819.	2.1	22
79	The Three Clades of the Telomere-Associated <i>TLO</i> Gene Family of <i>Candida albicans</i> Have Different Splicing, Localization, and Expression Features. <i>Eukaryotic Cell</i> , 2012, 11, 1268-1275.	3.4	38
80	<i>Candida albicans</i> . <i>Current Biology</i> , 2012, 22, R620-R622.	3.9	53
81	Cell-Cycle-Coupled Structural Oscillation of Centromeric Nucleosomes in Yeast. <i>Cell</i> , 2012, 150, 304-316.	28.9	92
82	Analysis of protein function in clinical <i>C. albicans</i> isolates. <i>Yeast</i> , 2012, 29, 303-309.	1.7	21
83	Neocentromeres and epigenetically inherited features of centromeres. <i>Chromosome Research</i> , 2012, 20, 607-619.	2.2	79
84	Does stress induce (para)sex? Implications for <i>Candida albicans</i> evolution. <i>Trends in Genetics</i> , 2012, 28, 197-203.	6.7	65
85	Flexibility of centromere and kinetochore structures. <i>Trends in Genetics</i> , 2012, 28, 204-212.	6.7	49
86	Skin-Resident Murine Dendritic Cell Subsets Promote Distinct and Opposing Antigen-Specific T Helper Cell Responses. <i>Immunity</i> , 2011, 35, 260-272.	14.3	379
87	Rad52 function prevents chromosome loss and truncation in <i>Candida albicans</i> . <i>Molecular Microbiology</i> , 2011, 79, 1462-1482.	2.5	28
88	CaMtw1, a member of the evolutionarily conserved Mis12 kinetochore protein family, is required for efficient inner kinetochore assembly in the pathogenic yeast <i>Candida albicans</i> . <i>Molecular Microbiology</i> , 2011, 80, 14-32.	2.5	30
89	The Requirement for the Dam1 Complex Is Dependent upon the Number of Kinetochore Proteins and Microtubules. <i>Current Biology</i> , 2011, 21, 889-896.	3.9	47
90	Evolutionary Dynamics of <i>Candida albicans</i> during <i>In Vitro</i> Evolution. <i>Eukaryotic Cell</i> , 2011, 10, 1413-1421.	3.4	30

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91	Stress Alters Rates and Types of Loss of Heterozygosity in <i>Candida albicans</i> . <i>MBio</i> , 2011, 2, .	4.1	196
92	High-Resolution SNP/CGH Microarrays Reveal the Accumulation of Loss of Heterozygosity in Commonly Used <i>Candida albicans</i> Strains. <i>G3: Genes, Genomes, Genetics</i> , 2011, 1, 523-530.	1.8	64
93	When abnormality is beneficial. <i>Nature</i> , 2010, 468, 183-184.	27.8	7
94	Genomic Plasticity of the Human Fungal Pathogen <i>Candida albicans</i> . <i>Eukaryotic Cell</i> , 2010, 9, 991-1008.	3.4	241
95	Epigenetically-Inherited Centromere and Neocentromere DNA Replicates Earliest in S-Phase. <i>PLoS Genetics</i> , 2010, 6, e1001068.	3.5	84
96	Low Dosage of Histone H4 Leads to Growth Defects and Morphological Changes in <i>Candida albicans</i> . <i>PLoS ONE</i> , 2010, 5, e10629.	2.5	10
97	SLA2 mutations cause SWE1-mediated cell cycle phenotypes in <i>Candida albicans</i> and <i>Saccharomyces cerevisiae</i> . <i>Microbiology (United Kingdom)</i> , 2009, 155, 3847-3859.	1.8	23
98	Acquisition of Aneuploidy Provides Increased Fitness during the Evolution of Antifungal Drug Resistance. <i>PLoS Genetics</i> , 2009, 5, e1000705.	3.5	293
99	Neocentromeres Form Efficiently at Multiple Possible Loci in <i>Candida albicans</i> . <i>PLoS Genetics</i> , 2009, 5, e1000400.	3.5	152
100	Aneuploid Chromosomes Are Highly Unstable during DNA Transformation of <i>Candida albicans</i> . <i>Eukaryotic Cell</i> , 2009, 8, 1554-1566.	3.4	77
101	Evolution in <i>Candida albicans</i> Populations During a Single Passage Through a Mouse Host. <i>Genetics</i> , 2009, 182, 799-811.	2.9	151
102	Additional cassettes for epitope and fluorescent fusion proteins in <i>Candida albicans</i> . <i>Yeast</i> , 2009, 26, 399-406.	1.7	65
103	Evolution of pathogenicity and sexual reproduction in eight <i>Candida</i> genomes. <i>Nature</i> , 2009, 459, 657-662.	27.8	963
104	Dancing genomes: fungal nuclear positioning. <i>Nature Reviews Microbiology</i> , 2009, 7, 875-886.	28.6	65
105	Efficient and rapid identification of <i>Candida albicans</i> allelic status using SNP-RFLP. <i>FEMS Yeast Research</i> , 2009, 9, 1061-1069.	2.3	32
106	Growth and development: eukaryotes. <i>Current Opinion in Microbiology</i> , 2009, 12, 589-591.	5.1	1
107	An isochromosome confers drug resistance <i>in vivo</i> by amplification of two genes, <i>ERG11</i> and <i>TAC1</i> . <i>Molecular Microbiology</i> , 2008, 68, 624-641.	2.5	280
108	Dynein-dependent nuclear dynamics affect morphogenesis in <i>Candida albicans</i> by means of the Bub2p spindle checkpoint. <i>Journal of Cell Science</i> , 2008, 121, 724-724.	2.0	3

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109	Haplotype Mapping of a Diploid Non-Meiotic Organism Using Existing and Induced Aneuploidies. <i>PLoS Genetics</i> , 2008, 4, e1.	3.5	129
110	The Parasexual Cycle in <i>Candida albicans</i> Provides an Alternative Pathway to Meiosis for the Formation of Recombinant Strains. <i>PLoS Biology</i> , 2008, 6, e110.	5.6	323
111	Dynein-dependent nuclear dynamics affect morphogenesis in <i>Candida albicans</i> by means of the Bub2p spindle checkpoint. <i>Journal of Cell Science</i> , 2008, 121, 466-476.	2.0	46
112	Molecular architecture of the kinetochore-microtubule attachment site is conserved between point and regional centromeres. <i>Journal of Cell Biology</i> , 2008, 181, 587-594.	5.2	144
113	Genotypic Evolution of Azole Resistance Mechanisms in Sequential <i>Candida albicans</i> Isolates. <i>Eukaryotic Cell</i> , 2007, 6, 1889-1904.	3.4	268
114	The pattern and evolution of yeast promoter bendability. <i>Trends in Genetics</i> , 2007, 23, 318-321.	6.7	62
115	Morphogenesis and cell cycle progression in <i>Candida albicans</i> . <i>Current Opinion in Microbiology</i> , 2006, 9, 595-601.	5.1	210
116	A Mutation in Tac1p, a Transcription Factor Regulating CDR1 and CDR2, Is Coupled With Loss of Heterozygosity at Chromosome 5 to Mediate Antifungal Resistance in <i>Candida albicans</i> . <i>Genetics</i> , 2006, 172, 2139-2156.	2.9	341
117	Transcript Profiles of <i>Candida albicans</i> Cortical Actin Patch Mutants Reflect Their Cellular Defects: Contribution of the Hog1p and Mkc1p Signaling Pathways. <i>Eukaryotic Cell</i> , 2006, 5, 1252-1265.	3.4	24
118	Aneuploidy and Isochromosome Formation in Drug-Resistant <i>Candida albicans</i> . <i>Science</i> , 2006, 313, 367-370.	12.6	630
119	Comparative genome hybridization reveals widespread aneuploidy in <i>Candida albicans</i> laboratory strains. <i>Molecular Microbiology</i> , 2005, 55, 1553-1565.	2.5	175
120	The Mitotic Cyclins Clb2p and Clb4p Affect Morphogenesis in <i>Candida albicans</i> . <i>Molecular Biology of the Cell</i> , 2005, 16, 3387-3400.	2.1	90
121	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.	3.5	124
122	A Human-Curated Annotation of the <i>Candida albicans</i> Genome. <i>PLoS Genetics</i> , 2005, 1, e1.	3.5	293
123	Microtubules in <i>Candida albicans</i> Hyphae Drive Nuclear Dynamics and Connect Cell Cycle Progression to Morphogenesis. <i>Eukaryotic Cell</i> , 2005, 4, 1697-1711.	3.4	58
124	<i>Candida albicans</i> hyphae have a Spitzenkörper that is distinct from the polarisome found in yeast and pseudohyphae. <i>Journal of Cell Science</i> , 2005, 118, 2935-2947.	2.0	164
125	Rewiring of the Yeast Transcriptional Network Through the Evolution of Motif Usage. <i>Science</i> , 2005, 309, 938-940.	12.6	268
126	Yeast Chromatin Assembly Complex 1 Protein Excludes Nonacetylatable Forms of Histone H4 from Chromatin and the Nucleus. <i>Molecular and Cellular Biology</i> , 2004, 24, 10180-10192.	2.3	38

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127	Telomere Cap Components Influence the Rate of Senescence in Telomerase-Deficient Yeast Cells. <i>Molecular and Cellular Biology</i> , 2004, 24, 837-845.	2.3	33
128	Transcriptional profiling in <i>Candida albicans</i> reveals new adaptive responses to extracellular pH and functions for Rim101p. <i>Molecular Microbiology</i> , 2004, 54, 1335-1351.	2.5	227
129	Cassettes for the PCR-mediated construction of regulatable alleles in <i>Candida albicans</i> . <i>Yeast</i> , 2004, 21, 429-436.	1.7	57
130	The distinct morphogenic states of <i>Candida albicans</i> . <i>Trends in Microbiology</i> , 2004, 12, 317-324.	7.7	725
131	Functional conservation of Dhh1p, a cytoplasmic DExD/H-box protein present in large complexes. <i>Nucleic Acids Research</i> , 2003, 31, 4995-5002.	14.5	37
132	Molecular Genetic and Genomic Approaches to the Study of Medically Important Fungi. <i>Infection and Immunity</i> , 2003, 71, 2299-2309.	2.2	34
133	mRNAs Encoding Telomerase Components and Regulators Are Controlled by UPF Genes in <i>Saccharomyces cerevisiae</i> . <i>Eukaryotic Cell</i> , 2003, 2, 134-142.	3.4	62
134	MEC3, MEC1, and DDC2 Are Essential Components of a Telomere Checkpoint Pathway Required for Cell Cycle Arrest during Senescence in <i>Saccharomyces cerevisiae</i> . <i>Molecular Biology of the Cell</i> , 2002, 13, 2626-2638.	2.1	119
135	A Forkhead Transcription Factor Is Important for True Hyphal as well as Yeast Morphogenesis in <i>Candida albicans</i> . <i>Eukaryotic Cell</i> , 2002, 1, 787-798.	3.4	114
136	<i>Candida albicans</i> : A molecular revolution built on lessons from budding yeast. <i>Nature Reviews Genetics</i> , 2002, 3, 918-931.	16.3	482
137	Cassettes for PCR-mediated construction of green, yellow, and cyan fluorescent protein fusions in <i>Candida albicans</i> . <i>Yeast</i> , 2001, 18, 859-864.	1.7	189
138	<i>Candida albicans</i> INT1-Induced Filamentation in <i>Saccharomyces cerevisiae</i> Depends on Sla2p. <i>Molecular and Cellular Biology</i> , 2001, 21, 1272-1284.	2.3	44
139	<i>Candida albicans</i> Int1p Interacts with the Septin Ring in Yeast and Hyphal Cells. <i>Molecular Biology of the Cell</i> , 2001, 12, 3538-3549.	2.1	76
140	CAC3 (MS11) Suppression of RAS2 G19V Is Independent of Chromatin Assembly Factor I and Mediated by NPR1. <i>Molecular and Cellular Biology</i> , 2001, 21, 1784-1794.	2.3	28
141	Filamentous Growth of <i>Saccharomyces cerevisiae</i> Is Regulated by Manganese. <i>Fungal Genetics and Biology</i> , 2000, 30, 155-162.	2.1	6
142	Identification of a Novel Allele of <i>SIR3</i> Defective in the Maintenance, but Not the Establishment, of Silencing in <i>Saccharomyces cerevisiae</i> . <i>Genetics</i> , 2000, 155, 523-538.	2.9	16
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