

Kaustuv Sanyal

List of Publications by Year in descending order

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Version: 2024-02-01

66
papers

2,323
citations

257450

24
h-index

254184

43
g-index

102
all docs

102
docs citations

102
times ranked

2311
citing authors

#	ARTICLE	IF	CITATIONS
1	Analysis of the Genome and Transcriptome of <i>Cryptococcus neoformans</i> var. <i>grubii</i> Reveals Complex RNA Expression and Microevolution Leading to Virulence Attenuation. <i>PLoS Genetics</i> , 2014, 10, e1004261.	3.5	336
2	Centromeric DNA sequences in the pathogenic yeast <i>Candida albicans</i> are all different and unique. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 11374-11379.	7.1	178
3	Broad Spectrum Antibacterial and Antifungal Polymeric Paint Materials: Synthesis, Structure–Activity Relationship, and Membrane-Active Mode of Action. <i>ACS Applied Materials & Interfaces</i> , 2015, 7, 1804-1815.	8.0	134
4	RNAi is a critical determinant of centromere evolution in closely related fungi. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 3108-3113.	7.1	112
5	Formation of functional centromeric chromatin is specified epigenetically in <i>Candida albicans</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 14877-14882.	7.1	91
6	Rapid evolution of Cse4p-rich centromeric DNA sequences in closely related pathogenic yeasts, <i>Candida albicans</i> and <i>Candida dubliniensis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 19797-19802.	7.1	81
7	Early Diverging Fungus <i>Mucor circinelloides</i> Lacks Centromeric Histone CENP-A and Displays a Mosaic of Point and Regional Centromeres. <i>Current Biology</i> , 2019, 29, 3791-3802.e6.	3.9	77
8	Efficient neocentromere formation is suppressed by gene conversion to maintain centromere function at native physical chromosomal loci in <i>Candida albicans</i> . <i>Genome Research</i> , 2013, 23, 638-652.	5.5	76
9	Chitosan Derivatives Active against Multidrug-Resistant Bacteria and Pathogenic Fungi: <i>In Vivo</i> Evaluation as Topical Antimicrobials. <i>Molecular Pharmaceutics</i> , 2016, 13, 3578-3589.	4.6	71
10	Fungal genome and mating system transitions facilitated by chromosomal translocations involving intercentromeric recombination. <i>PLoS Biology</i> , 2017, 15, e2002527.	5.6	67
11	The CENP-A homolog CaCse4p in the pathogenic yeast <i>Candida albicans</i> is a centromere protein essential for chromosome transmission. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 12969-12974.	7.1	66
12	Repeat-Associated Fission Yeast-Like Regional Centromeres in the Ascomycetous Budding Yeast <i>Candida tropicalis</i> . <i>PLoS Genetics</i> , 2016, 12, e1005839.	3.5	56
13	A Coordinated Interdependent Protein Circuitry Stabilizes the Kinetochores Ensemble to Protect CENP-A in the Human Pathogenic Yeast <i>Candida albicans</i> . <i>PLoS Genetics</i> , 2012, 8, e1002661.	3.5	47
14	Proteogenomics produces comprehensive and highly accurate protein-coding gene annotation in a complete genome assembly of <i>Malassezia sympodialis</i> . <i>Nucleic Acids Research</i> , 2017, 45, gkx006.	14.5	47
15	Loss of centromere function drives karyotype evolution in closely related <i>Malassezia</i> species. <i>ELife</i> , 2020, 9, .	6.0	45
16	Ordered Kinetochores Assembly in the Human-Pathogenic Basidiomycetous Yeast <i>Cryptococcus neoformans</i> . <i>MBio</i> , 2013, 4, e00614-13.	4.1	42
17	The Essentiality of the Fungus-Specific Dam1 Complex Is Correlated with a One-Kinetochores-One-Microtubule Interaction Present throughout the Cell Cycle, Independent of the Nature of a Centromere. <i>Eukaryotic Cell</i> , 2011, 10, 1295-1305.	3.4	41
18	Diversity in Requirement of Genetic and Epigenetic Factors for Centromere Function in Fungi. <i>Eukaryotic Cell</i> , 2011, 10, 1384-1395.	3.4	39

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19	Rad51â€™Rad52 Mediated Maintenance of Centromeric Chromatin in <i>Candida albicans</i> . <i>PLoS Genetics</i> , 2014, 10, e1004344.	3.5	37
20	Mode of Action of a Designed Antimicrobial Peptide: High Potency against <i>Cryptococcus neoformans</i> . <i>Biophysical Journal</i> , 2016, 111, 1724-1737.	0.5	37
21	Fluconazole-Induced Ploidy Change in <i>Cryptococcus neoformans</i> Results from the Uncoupling of Cell Growth and Nuclear Division. <i>MSphere</i> , 2017, 2, .	2.9	35
22	<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
23	Spatial inter-centromeric interactions facilitated the emergence of evolutionary new centromeres. <i>ELife</i> , 2020, 9, .	6.0	31
24	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
25	Long transposon-rich centromeres in an oomycete reveal divergence of centromere features in <i>Stramenopila-Alveolata-Rhizaria</i> lineages. <i>PLoS Genetics</i> , 2020, 16, e1008646.	3.5	29
26	Impact of <i>FKS1</i> Genotype on Echinocandin <i>In Vitro</i> Susceptibility in <i>Candida auris</i> and <i>In Vivo</i> Response in a Murine Model of Infection. <i>Antimicrobial Agents and Chemotherapy</i> , 2022, 66, AAC0165221.	3.2	29
27	Dual-Function Polymerâ€™Silver Nanocomposites for Rapid Killing of Microbes and Inhibiting Biofilms. <i>ACS Biomaterials Science and Engineering</i> , 2019, 5, 81-91.	5.2	26
28	The IML3/MCM19 gene of <i>Saccharomyces cerevisiae</i> is required for a kinetochore-related process during chromosome segregation. <i>Molecular Genetics and Genomics</i> , 2001, 265, 249-257.	2.1	25
29	A comprehensive model to predict mitotic division in budding yeasts. <i>Molecular Biology of the Cell</i> , 2015, 26, 3954-3965.	2.1	25
30	Aryl-alkyl-lysines: Membrane-Active Fungicides That Act against Biofilms of <i>Candida albicans</i> . <i>ACS Infectious Diseases</i> , 2017, 3, 293-301.	3.8	25
31	Functional characterization of the <i>Saccharomyces cerevisiae</i> protein Chl1 reveals the role of sister chromatid cohesion in the maintenance of spindle length during S-phase arrest. <i>BMC Genetics</i> , 2011, 12, 83.	2.7	23
32	The <i>Candida albicans</i> biofilm gene circuit modulated at the chromatin level by a recent molecular histone innovation. <i>PLoS Biology</i> , 2019, 17, e3000422.	5.6	22
33	Implications of the Evolutionary Trajectory of Centromeres in the Fungal Kingdom. <i>Annual Review of Microbiology</i> , 2020, 74, 835-853.	7.3	22
34	The MCM16 gene of the yeast <i>Saccharomyces cerevisiae</i> is required for chromosome segregation. <i>Molecular Genetics and Genomics</i> , 1998, 260, 242-250.	2.4	19
35	Epigenetic determinants of phenotypic plasticity in <i>Candida albicans</i> . <i>Fungal Biology Reviews</i> , 2018, 32, 10-19.	4.7	19
36	Spatio-temporal regulation of nuclear division by Aurora B kinase Ipl1 in <i>Cryptococcus neoformans</i> . <i>PLoS Genetics</i> , 2019, 15, e1007959.	3.5	19

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37	Five pillars of centromeric chromatin in fungal pathogens. PLoS Pathogens, 2018, 14, e1007150.	4.7	18
38	Cellular Dynamics and Genomic Identity of Centromeres in Cereal Blast Fungus. MBio, 2019, 10, .	4.1	18
39	Bridgin connects the outer kinetochore to centromeric chromatin. Nature Communications, 2021, 12, 146.	12.8	17
40	Sad1 Spatiotemporally Regulates Kinetochore Clustering To Ensure High-Fidelity Chromosome Segregation in the Human Fungal Pathogen <i>Cryptococcus neoformans</i> . MSphere, 2018, 3, .	2.9	14
41	The process of kinetochore assembly in yeasts. FEMS Microbiology Letters, 2013, 338, 107-117.	1.8	13
42	Cis- and Trans-chromosomal Interactions Define Pericentric Boundaries in the Absence of Conventional Heterochromatin. Genetics, 2019, 212, 1121-1132.	2.9	13
43	How Do Microbial Pathogens Make CENs?. PLoS Pathogens, 2012, 8, e1002463.	4.7	12
44	ZCF32, a fungus specific Zn(II) ₂ Cys ₆ transcription factor, is a repressor of the biofilm development in the human pathogen <i>Candida albicans</i> . Scientific Reports, 2016, 6, 31124.	3.3	11
45	Nuclear migration in budding yeasts: position before division. Current Genetics, 2019, 65, 1341-1346.	1.7	11
46	Functional and Comparative Analysis of Centromeres Reveals Clade-Specific Genome Rearrangements in <i>Candida auris</i> and a Chromosome Number Change in Related Species. MBio, 2021, 12, .	4.1	11
47	A Surprising Role for the Sch9 Protein Kinase in Chromosome Segregation in <i>Candida albicans</i> . Genetics, 2015, 199, 671-674.	2.9	10
48	Mechanics of microtubule organizing center clustering and spindle positioning in budding yeast <i>Cryptococcus neoformans</i> . Physical Review E, 2021, 104, 034402.	2.1	8
49	Sth1, the Key Subunit of the RSC Chromatin Remodeling Complex, Is Essential in Maintaining Chromosomal Integrity and Mediating High Fidelity Chromosome Segregation in the Human Fungal Pathogen <i>Candida albicans</i> . Frontiers in Microbiology, 2019, 10, 1303.	3.5	7
50	Aurora kinase Ipl1 facilitates bilobed distribution of clustered kinetochores to ensure error-free chromosome segregation in <i>Candida albicans</i> . Molecular Microbiology, 2019, 112, 569-587.	2.5	7
51	ClalD: a Rapid Method of Clade-Level Identification of the Multidrug Resistant Human Fungal Pathogen <i>Candida auris</i> . Microbiology Spectrum, 2022, 10, e0063422.	3.0	7
52	Chromatin Immunoprecipitation (ChIP) Assay in <i>Candida albicans</i> . Methods in Molecular Biology, 2016, 1356, 43-57.	0.9	6
53	Two negative regulators of biofilm development exhibit functional divergence in conferring virulence potential to <i>Candida albicans</i> . FEMS Yeast Research, 2019, 19, .	2.3	5
54	Magnetic hyperthermia adjunctive therapy for fungi: <i>in vitro</i> studies against <i>Candida albicans</i> . International Journal of Hyperthermia, 2019, 36, 544-552.	2.5	5

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55	Orc4 spatiotemporally stabilizes centromeric chromatin. <i>Genome Research</i> , 2021, 31, 607-621.	5.5	5
56	Minichromosome maintenance proteins in eukaryotic chromosome segregation. <i>BioEssays</i> , 2022, 44, e2100218.	2.5	5
57	Establishing a national fungal genetic resource to build a major cog for the bioeconomy. <i>Current Science</i> , 2015, 109, 1033.	0.8	4
58	Shugoshin ensures maintenance of the spindle assembly checkpoint response and efficient spindle disassembly. <i>Molecular Microbiology</i> , 2021, 116, 1079-1098.	2.5	3
59	Establishing a national fungal genetic resource to build a major cog for the bioeconomy. <i>Current Science</i> , 2015, 109, 1033.	0.8	2
60	A Stable Hybrid Containing Haploid Genomes of Two Obligate Diploid <i>Candida</i> Species. <i>Eukaryotic Cell</i> , 2013, 12, 1061-1071.	3.4	1
61	Early Diverging Fungus <i>Mucor circinelloides</i> Lacks Centromeric Histone CENP-A and Displays a Mosaic of Point and Regional Centromeres. <i>SSRN Electronic Journal</i> , 0, , .	0.4	1
62	Loss of nucleosome assembly protein 1 affects growth and appressorium structure in blast fungus .. <i>MicroPublication Biology</i> , 2022, 2022, .	0.1	1
63	Hypersaline fungi as a source of potentially active metabolites against pathogenic <i>Candida</i> species.. <i>Czech Mycology</i> , 2022, 74, 93-101.	0.5	1
64	Chromosome Components Important for Genome Stability in <i>Candida albicans</i> and Related Species. , 2017, , 233-251.		0
65	Vacuolar transporter Mnr2 safeguards organellar integrity in aged cells. <i>Molecular Microbiology</i> , 2021, 116, 861-876.	2.5	0
66	Identification and analysis of the origin recognition complex in the human fungal pathogen. <i>MicroPublication Biology</i> , 2021, 2021, .	0.1	0