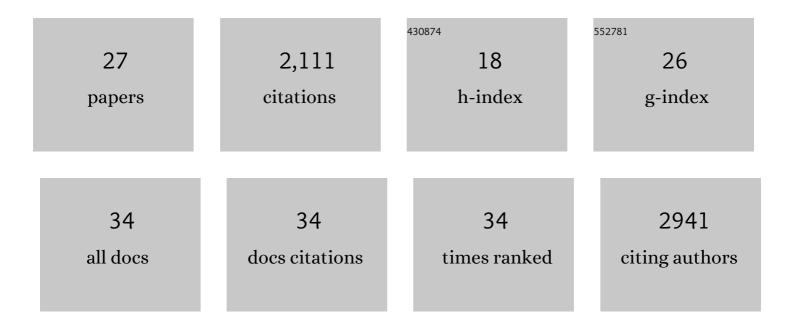
Anita Sil

List of Publications by Year in descending order

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ΔΝΙΤΑ ΟΠ

#	Article	IF	CITATIONS
1	Amplification-free detection of SARS-CoV-2 with CRISPR-Cas13a and mobile phone microscopy. Cell, 2021, 184, 323-333.e9.	28.9	613
2	Genetic Screens Identify Host Factors for SARS-CoV-2 and Common Cold Coronaviruses. Cell, 2021, 184, 106-119.e14.	28.9	320
3	Temperature-induced switch to the pathogenic yeast form of <i>Histoplasma capsulatum</i> requires Ryp1, a conserved transcriptional regulator. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 4880-4885.	7.1	179
4	Histoplasma Requires SID1, a Member of an Iron-Regulated Siderophore Gene Cluster, for Host Colonization. PLoS Pathogens, 2008, 4, e1000044.	4.7	131
5	A Temperature-Responsive Network Links Cell Shape and Virulence Traits in a Primary Fungal Pathogen. PLoS Biology, 2013, 11, e1001614.	5.6	115
6	Identifying Phase-specific Genes in the Fungal PathogenHistoplasma capsulatumUsing a Genomic Shotgun Microarray. Molecular Biology of the Cell, 2003, 14, 2314-2326.	2.1	111
7	Conserved factors Ryp2 and Ryp3 control cell morphology and infectious spore formation in the fungal pathogen <i>Histoplasma capsulatum</i> . Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 14573-14578.	7.1	108
8	Thermally Dimorphic Human Fungal Pathogens—Polyphyletic Pathogens with a Convergent Pathogenicity Trait. Cold Spring Harbor Perspectives in Medicine, 2015, 5, a019794.	6.2	103
9	N-acetylglucosamine (GlcNAc) Triggers a Rapid, Temperature-Responsive Morphogenetic Program in Thermally Dimorphic Fungi. PLoS Genetics, 2013, 9, e1003799.	3.5	58
10	Comparative Transcriptomics of Infectious Spores from the Fungal Pathogen Histoplasma capsulatum Reveals a Core Set of Transcripts That Specify Infectious and Pathogenic States. Eukaryotic Cell, 2013, 12, 828-852.	3.4	45
11	Genome-Wide Reprogramming of Transcript Architecture by Temperature Specifies the Developmental States of the Human Pathogen Histoplasma. PLoS Genetics, 2015, 11, e1005395.	3.5	35
12	Macrophage cell death and transcriptional response are actively triggered by the fungal virulence factor <scp>C</scp> bp1 during <scp><i>H</i></scp> . <i>capsulatum</i> infection. Molecular Microbiology, 2015, 98, 910-929.	2.5	34
13	CD103+ Conventional Dendritic Cells Are Critical for TLR7/9-Dependent Host Defense against Histoplasma capsulatum, an Endemic Fungal Pathogen of Humans. PLoS Pathogens, 2016, 12, e1005749.	4.7	32
14	The 3-Hydroxy-Methylglutaryl Coenzyme A Lyase <i>HCL1</i> Is Required for Macrophage Colonization by Human Fungal Pathogen Histoplasma capsulatum. Infection and Immunity, 2013, 81, 411-420.	2.2	30
15	COVID-19–associated Lung Microvascular Endotheliopathy: A "From the Bench―Perspective. American Journal of Respiratory and Critical Care Medicine, 2022, 206, 961-972.	5.6	30
16	The transcription factor CHOP, an effector of the integrated stress response, is required for host sensitivity to the fungal intracellular pathogen Histoplasma capsulatum. PLoS Pathogens, 2017, 13, e1006589.	4.7	26
17	Opposing signaling pathways regulate morphology in response to temperature in the fungal pathogen Histoplasma capsulatum. PLoS Biology, 2019, 17, e3000168.	5.6	22
18	The protein kinase Pho85 is required for asymmetric accumulation of the Ash1 protein in Saccharomyces cerevisiae. Molecular Microbiology, 2001, 42, 345-353.	2.5	21

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19	MyD88-Dependent Signaling Drives Host Survival and Early Cytokine Production during Histoplasma capsulatum Infection. Infection and Immunity, 2015, 83, 1265-1275.	2.2	18
20	Recurrent Loss of abaA, a Master Regulator of Asexual Development in Filamentous Fungi, Correlates with Changes in Genomic and Morphological Traits. Genome Biology and Evolution, 2020, 12, 1119-1130.	2.5	16
21	Molecular regulation of Histoplasma dimorphism. Current Opinion in Microbiology, 2019, 52, 151-157.	5.1	11
22	The WOPR family protein Ryp1 is a key regulator of gene expression, development, and virulence in the thermally dimorphic fungal pathogen Coccidioides posadasii. PLoS Pathogens, 2022, 18, e1009832.	4.7	9
23	Sensing the heat and the host: Virulence determinants of Histoplasma capsulatum. Virulence, 2019, 10, 793-800.	4.4	8
24	Protomer alignment modulates specificity of RNA substrate recognition by Ire1. ELife, 2021, 10, .	6.0	7
25	Chromosome-Level Genome Assembly of a Human Fungal Pathogen Reveals Synteny among Geographically Distinct Species. MBio, 2022, 13, e0257421.	4.1	7
26	Cbp1, a fungal virulence factor under positive selection, forms an effector complex that drives macrophage lysis. PLoS Pathogens, 2022, 18, e1010417.	4.7	4
27	Future of Functional Genomics of Histoplasma capsulatum. , 0, , 611-625.		1