Suman B Pakala

List of Publications by Year in descending order

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623734 642732 1,056 26 14 23 citations g-index h-index papers 30 30 30 1800 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	COVIDâ€19 severity from Omicron and Delta SARSâ€CoVâ€2 variants. Influenza and Other Respiratory Viruses, 2022, 16, 832-836.	3.4	60
2	Molecular signature of postmortem lung tissue from COVID-19 patients suggests distinct trajectories driving mortality. DMM Disease Models and Mechanisms, 2022, 15, .	2.4	14
3	Specific Ablation of Jak2 from CD11c+ Cells Attenuates Saltâ€Sensitive Hypertension through an ENaCâ€Dependent Mechanism. FASEB Journal, 2022, 36, .	0.5	O
4	DC ENaC-Dependent Inflammasome Activation Contributes to Salt-Sensitive Hypertension. Circulation Research, 2022, 131, 328-344.	4.5	31
5	Transcriptomic profiling of Paulownia elongata in response to heat stress. Plant Gene, 2021, 28, 100330.	2.3	О
6	Metatranscriptomics to characterize respiratory virome, microbiome, and host response directly from clinical samples. Cell Reports Methods, 2021, 1, 100091.	2.9	19
7	Generation of a Novel SARS-CoV-2 Sub-genomic RNA Due to the R203K/G204R Variant in Nucleocapsid: Homologous Recombination has Potential to Change SARS-CoV-2 at Both Protein and RNA Level. Pathogens and Immunity, 2021, 6, 27-49.	3.1	10
8	Generation of a Novel SARS-CoV-2 Sub-genomic RNA Due to the R203K/G204R Variant in Nucleocapsid: Homologous Recombination has Potential to Change SARS-CoV-2 at Both Protein and RNA Level. Pathogens and Immunity, 2021, 6, 27-49.	3.1	46
9	Practical Recommendations for Supporting a Systems Biology Cyberinfrastructure. Data Science Journal, 2020, 19, .	1.3	7
10	Distinct amino acid and lipid perturbations characterize acute versus chronic malaria. JCI Insight, 2019, 4, .	5.0	46
11	NCBI's Virus Discovery Hackathon: Engaging Research Communities to Identify Cloud Infrastructure Requirements. Genes, 2019, 10, 714.	2.4	13
12	Nearly Complete Genome Sequences of 17 Enterovirus D68 Strains from Kansas City, Missouri, 2018. Microbiology Resource Announcements, 2019, 8, .	0.6	6
13	Cross-reaction between Formosan termite (Coptotermes formosanus) proteins and cockroach allergens. PLoS ONE, 2017, 12, e0182260.	2.5	11
14	Termite Proteins Cross-React with Cockroach Allergens. Journal of Allergy and Clinical Immunology, 2016, 137, AB266.	2.9	0
15	High-Quality Genome Assembly and Annotation for Plasmodium coatneyi, Generated Using Single-Molecule Real-Time PacBio Technology. Genome Announcements, 2016, 4, .	0.8	33
16	Genetic Analysis Using an Isogenic Mating Pair of Aspergillus fumigatus Identifies Azole Resistance Genes and Lack of MAT Locus's Role in Virulence. PLoS Pathogens, 2015, 11, e1004834.	4.7	52
17	Suppression subtractive hybridization and comparative expression of a pore-forming toxin and glycosyl hydrolase genes in Rhizoctonia solani during potato sprout infection. Molecular Genetics and Genomics, 2015, 290, 877-900.	2.1	10
18	Comparative transcriptomics and metabolomics in a rhesus macaque drug administration study. Frontiers in Cell and Developmental Biology, 2014, 2, 54.	3.7	15

#	Article	IF	CITATION
19	Large-Scale Transcriptional Response to Hypoxia in Aspergillus fumigatus Observed Using RNAseq Identifies a Novel Hypoxia Regulated ncRNA. Mycopathologia, 2014, 178, 331-339.	3.1	29
20	Perturbations in small molecule synthesis uncovers an iron-responsive secondary metabolite network in Aspergillus fumigatus. Frontiers in Microbiology, 2014, 5, 530.	3.5	59
21	Draft Genome Sequence of the Plant-Pathogenic Soil Fungus Rhizoctonia solani Anastomosis Group 3 Strain Rhs1AP. Genome Announcements, 2014, 2, .	0.8	49
22	The binary protein-protein interaction landscape of Escherichia coli. Nature Biotechnology, 2014, 32, 285-290.	17.5	218
23	Mobile elements and mitochondrial genome expansion in the soil fungus and potato pathogen <i>Rhizoctonia solani</i> AG-3. FEMS Microbiology Letters, 2014, 352, 165-173.	1.8	143
24	Structural analysis of the <i><scp>R</scp>hizoctoniaÂsolani</i> agglutinin reveals a domainâ€swapping dimeric assembly. FEBS Journal, 2013, 280, 1750-1763.	4.7	19
25	Triallelic SNP-mediated genotyping of regenerated protoplasts of the heterokaryotic fungus Rhizoctonia solani. Journal of Biotechnology, 2012, 158, 144-150.	3.8	8
26	Sequencing of mitochondrial genomes of nine Aspergillus and Penicillium species identifies mobile introns and accessory genes as main sources of genome size variability. BMC Genomics, 2012, 13, 698.	2.8	131