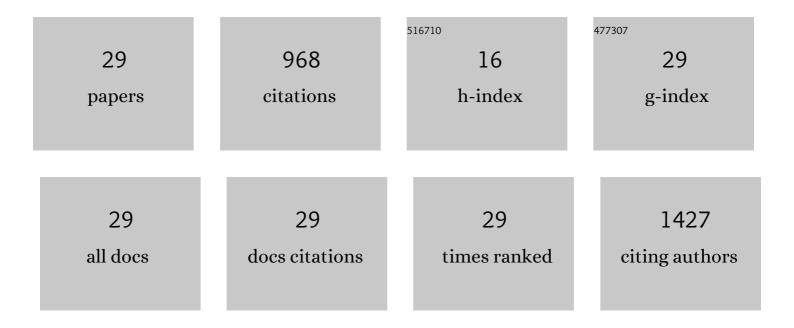
Mohit Kumar Swarnkar

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5761727/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Deciphering key regulators involved in epilepsyâ€induced cardiac damage through whole transcriptome and proteome analysis in a rat model. Epilepsia, 2021, 62, 504-516.	5.1	17
2	Comparative transcriptome analysis of Rheum australe, an endangered medicinal herb, growing in its natural habitat and those grown in controlled growth chambers. Scientific Reports, 2021, 11, 3702.	3.3	11
3	Prickle morphogenesis in rose is coupled with secondary metabolite accumulation and governed by canonical MBW transcriptional complex. Plant Direct, 2021, 5, e00325.	1.9	13
4	De novo transcriptome provides insights into the growth behaviour and resveratrol and trans-stilbenes biosynthesis in Dactylorhiza hatagirea - An endangered alpine terrestrial orchid of western Himalaya. Scientific Reports, 2019, 9, 13133.	3.3	20
5	Transcriptome and Co-Expression Network Analyses Identify Key Genes Regulating Nitrogen Use Efficiency in Brassica juncea L Scientific Reports, 2018, 8, 7451.	3.3	37
6	Chilling Affects Phytohormone and Post-Embryonic Development Pathways during Bud Break and Fruit Set in Apple (Malus domestica Borkh.). Scientific Reports, 2017, 7, 42593.	3.3	22
7	Paenibacillus ihbetae sp. nov., a cold-adapted antimicrobial producing bacterium isolated from high altitude Suraj Tal Lake in the Indian trans-Himalayas. Systematic and Applied Microbiology, 2017, 40, 430-439.	2.8	14
8	Molecular dissection of transcriptional reprogramming of steviol glycosides synthesis in leaf tissue during developmental phase transitions in Stevia rebaudiana Bert. Scientific Reports, 2017, 7, 11835.	3.3	51
9	Genome assembly of Chryseobacterium sp. strain IHBB 10212 from glacier top-surface soil in the Indian trans-Himalayas with potential for hydrolytic enzymes. Genomics Data, 2017, 13, 46-49.	1.3	9
10	Complete genome sequence of a low-temperature active and alkaline-stable endoglucanase-producing Paenibacillus sp. strain IHB B 3084 from the Indian Trans-Himalayas. Journal of Biotechnology, 2016, 230, 1-2.	3.8	11
11	Transcriptome Analysis Reveals Candidate Genes involved in Blister Blight defense in Tea (Camellia) Tj ETQq1 1	0.784314	rgBT_/Overloc
12	Complete genome sequence of Arthrobacter alpinus ERGS4:06, a yellow pigmented bacterium tolerant to cold and radiations isolated from Sikkim Himalaya. Journal of Biotechnology, 2016, 220, 86-87.	3.8	30
13	First complete genome sequence of a species in the genus Microterricola, an extremophilic cold active enzyme producing bacterial strain ERGS5:02 isolated from Sikkim Himalaya. Journal of Biotechnology, 2016, 222, 17-18.	3.8	13
14	Comprehensive transcriptome analysis of Crocus sativus for discovery and expression of genes involved in apocarotenoid biosynthesis. BMC Genomics, 2015, 16, 698.	2.8	81
15	Complete Genome Sequencing of Protease-Producing Novel <i>Arthrobacter</i> sp. Strain IHBB 11108 Using PacBio Single-Molecule Real-Time Sequencing Technology. Genome Announcements, 2015, 3, .	0.8	23
16	Draft Genome Sequence of a Cellulase-Producing Psychrotrophic Paenibacillus Strain, IHB B 3415, Isolated from the Cold Environment of the Western Himalayas, India. Genome Announcements, 2015, 3, .	0.8	8
17	Complete Genome Sequence of Potential Probiotic <i>Lactobacillus</i> sp. HFC8, Isolated from Human Gut Using PacBio SMRT Sequencing. Genome Announcements, 2015, 3, .	0.8	14
18	Genome Sequence of a Potential Probiotic Strain, Lactobacillus fermentum HFB3, Isolated from a Human Gut. Genome Announcements, 2015, 3, .	0.8	3

#	Article	IF	CITATIONS
19	Expression of SOD and APX genes positively regulates secondary cell wall biosynthesis and promotes plant growth and yield in Arabidopsis under salt stress. Plant Molecular Biology, 2015, 87, 615-631.	3.9	183
20	Complete genome sequence of Arthrobacter sp. ERGS1:01, a putative novel bacterium with prospective cold active industrial enzymes, isolated from East Rathong glacier in India. Journal of Biotechnology, 2015, 214, 139-140.	3.8	37
21	Complete Genome Sequence of Paenibacillus sp. Strain IHBB 10380 Using PacBio Single-Molecule Real-Time Sequencing Technology. Genome Announcements, 2015, 3, .	0.8	4
22	Complete Genome Sequence of the Rhizobacterium Pseudomonas trivialis Strain IHBB745 with Multiple Plant Growth-Promoting Activities and Tolerance to Desiccation and Alkalinity. Genome Announcements, 2015, 3, .	0.8	8
23	Genome Assembly of Chryseobacterium polytrichastri ERMR1:04, a Psychrotolerant Bacterium with Cold Active Proteases, Isolated from East Rathong Glacier in India. Genome Announcements, 2015, 3, .	0.8	18
24	Draft Genome Sequence of Psychrotrophic <i>Acinetobacter</i> sp. Strain MN12 (MTCC 10786), Which Produces a Low-Temperature-Active and Alkaline-Stable Peptidase. Genome Announcements, 2014, 2, .	0.8	2
25	Transcriptome sequencing of rhizome tissue of Sinopodophyllum hexandrum at two temperatures. BMC Genomics, 2014, 15, 871.	2.8	35
26	Comprehensive transcriptomic study on horse gram (Macrotyloma uniflorum): De novo assembly, functional characterization and comparative analysis in relation to drought stress. BMC Genomics, 2013, 14, 647.	2.8	71
27	A Sequence-Based Variation Map of Zebrafish. Zebrafish, 2013, 10, 15-20.	1.1	40
28	De Novo Transcriptome Sequencing and Analysis for Venturia inaequalis, the Devastating Apple Scab Pathogen. PLoS ONE, 2013, 8, e53937.	2.5	32
29	Protein Model Discrimination Using Mutational Sensitivity Derived from Deep Sequencing. Structure, 2012, 20, 371-381.	3.3	83