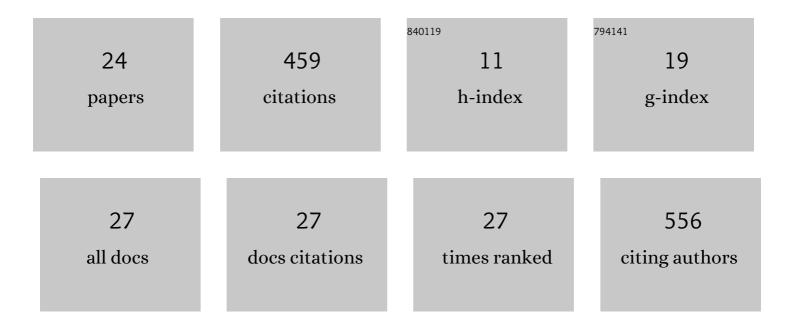
Renesh Bedre

List of Publications by Year in descending order

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RENECH REDDE

#	Article	IF	CITATIONS
1	Transcriptome analysis of smooth cordgrass (Spartina alterniflora Loisel), a monocot halophyte, reveals candidate genes involved in its adaptation to salinity. BMC Genomics, 2016, 17, 657.	1.2	55
2	Genetic Mapping of Quantitative Trait Loci for Grain Yield under Drought in Rice under Controlled Greenhouse Conditions. Frontiers in Chemistry, 2017, 5, 129.	1.8	43
3	Identification of cold-responsive genes in energycane for their use in genetic diversity analysis and future functional marker development. Plant Science, 2013, 211, 122-131.	1.7	42
4	Salt Adaptation Mechanisms of Halophytes: Improvement of Salt Tolerance in Crop Plants. , 2015, , 243-279.		36
5	Plant hairy roots enable high throughput identification of antimicrobials against Candidatus Liberibacter spp Nature Communications, 2020, 11, 5802.	5.8	36
6	Global Awakening of Cryptic Biosynthetic Gene Clusters in <i>Burkholderia thailandensis</i> . ACS Chemical Biology, 2017, 12, 3012-3021.	1.6	35
7	Genome-Wide Transcriptome Analysis of Cotton (Gossypium hirsutum L.) Identifies Candidate Gene Signatures in Response to Aflatoxin Producing Fungus Aspergillus flavus. PLoS ONE, 2015, 10, e0138025.	1.1	30
8	An actinâ€depolymerizing factor from the halophyte smooth cordgrass, <i>Spartina alterniflora</i> (<i>Sa<scp>ADF</scp>2</i>), is superior to its rice homolog (<i>Os<scp>ADF</scp>2</i>) in conferring drought and salt tolerance when constitutively overexpressed in rice. Plant Biotechnology Journal, 2019, 17, 188-205.	4.1	25
9	Genome-wide alternative splicing landscapes modulated by biotrophic sugarcane smut pathogen. Scientific Reports, 2019, 9, 8876.	1.6	24
10	Sequencing and expression analysis of salt-responsive miRNAs and target genes in the halophyte smooth cordgrass (Spartina alternifolia Loisel). Molecular Biology Reports, 2015, 42, 1341-1350.	1.0	19
11	GenFam: A web application and database for gene familyâ€based classification and functional enrichment analysis. Plant Direct, 2019, 3, e00191.	0.8	16
12	<i>Brachypodium</i> Phenylalanine Ammonia Lyase (PAL) Promotes Antiviral Defenses against <i>Panicum mosaic virus</i> and Its Satellites. MBio, 2021, 12, .	1.8	16
13	Developing Growthâ€Associated Molecular Markers Via Highâ€Throughput Phenotyping in Spinach. Plant Genome, 2019, 12, 190027.	1.6	15
14	Singleâ€marker and haplotypeâ€based association analysis of anthracnose (<i>Colletotrichum) Tj ETQq0 0 0 rgB</i>	T /Overloc	k 10 Tf 50 2
15	Identifying cotton (Gossypium hirsutum L.) genes induced in response to Aspergillus flavus infection. Physiological and Molecular Plant Pathology, 2012, 80, 35-40.	1.3	9
16	New Era in Plant Alternative Splicing Analysis Enabled by Advances in High-Throughput Sequencing (HTS) Technologies. Frontiers in Plant Science, 2019, 10, 740.	1.7	9
17	Genetic Diversity and Population Structure of Tomato (<i>Solanum) Tj ETQq1 1 0.784314 rgBT /C American Journal of Plant Sciences, 2019, 10, 1154-1180.</i>)verlock 1 0.3	0 Tf 50 107 8
18	Antiviral therapy with nucleotide/nucleoside analogues in chronic hepatitis B: A meta-analysis of prospective randomized trials. Indian Journal of Gastroenterology, 2016, 35, 75-82.	0.7	6

#	Article	IF	CITATIONS
19	Minor alleles are associated with white rust (Albugo occidentalis) susceptibility in spinach (Spinacia) Tj ETQq1 1 ().784314 ı 2.9	rg&T /Overlo
20	Genomic Approaches to Analyze Alternative Splicing, A Key Regulator of Transcriptome and Proteome Diversity in Brachypodium distachyon. Methods in Molecular Biology, 2018, 1667, 73-85.	0.4	4
21	Phenotypic Diversity and Association Mapping of Ascorbic Acid Content in Spinach. Frontiers in Genetics, 2021, 12, 752313.	1.1	4
22	Identification of candidate resistance genes of cotton against Aspergillus flavus infection using a comparative transcriptomics approach. Physiology and Molecular Biology of Plants, 2018, 24, 513-519.	1.4	3
23	HTSQualC is a flexible and one-step quality control software for high-throughput sequencing data analysis. Scientific Reports, 2021, 11, 18725.	1.6	3
24	Efficacy of Sucralfate in Reflux Disease in comparison to H2 – Receptor Antagonist: A Meta-Analysis of Randomized Trials. Biology Engineering Medicine and Science Reports, 2016, 2, 18-22.	0.5	0