

Kamran Shah

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

345
citations

933447

10
h-index

839539

18
g-index

25
all docs

25
docs citations

25
times ranked

277
citing authors

#	ARTICLE	IF	CITATIONS
1	A mini review fermentation and preservation: role of Lactic Acid Bacteria. <i>MOJ Food Processing & Technology</i> , 2018, 6, 414-417.	0.9	38
2	Chronic cement dust load induce novel damages in foliage and buds of <i>Malus domestica</i> . <i>Scientific Reports</i> , 2020, 10, 12186.	3.3	29
3	Impact assessment of leaf pigments in selected landscape plants exposed to roadside dust. <i>Environmental Science and Pollution Research</i> , 2018, 25, 23055-23073.	5.3	28
4	Comparative RNA-Sequencing and DNA Methylation Analyses of Apple (<i>Malus domestica</i> Borkh.) Buds with Diverse Flowering Capabilities Reveal Novel Insights into the Regulatory Mechanisms of Flower Bud Formation. <i>Plant and Cell Physiology</i> , 2019, 60, 1702-1721.	3.1	27
5	Genome-Wide Identification of the MdKNOX Gene Family and Characterization of Its Transcriptional Regulation in <i>Malus domestica</i> . <i>Frontiers in Plant Science</i> , 2020, 11, 128.	3.6	24
6	Role of Species and Planting Configuration on Transpiration and Microclimate for Urban Trees. <i>Forests</i> , 2020, 11, 825.	2.1	23
7	Identification and characterization of NRT gene family reveals their critical response to nitrate regulation during adventitious root formation and development in apple rootstock. <i>Scientia Horticulturae</i> , 2021, 275, 109642.	3.6	23
8	Mediation of Flower Induction by Gibberellin and its Inhibitor Paclobutrazol: mRNA and miRNA Integration Comprises Complex Regulatory Cross-Talk in Apple. <i>Plant and Cell Physiology</i> , 2018, 59, 2288-2307.	3.1	21
9	Comparative RNA-sequencing-based transcriptome profiling of buds from profusely flowering ‘Qinguan’ and weakly flowering ‘Nagafu no. 2’ apple varieties reveals novel insights into the regulatory mechanisms underlying floral induction. <i>BMC Plant Biology</i> , 2018, 18, 370.	3.6	19
10	Expression of genes in the potential regulatory pathways controlling alternate bearing in ‘Fuji’ apple trees during flower induction. <i>Plant Physiology and Biochemistry</i> , 2018, 132, 579-589.	5.8	15
11	Cement dust induce stress and attenuates photosynthesis in <i>Arachis hypogaea</i> . <i>Environmental Science and Pollution Research</i> , 2019, 26, 19490-19501.	5.3	11
12	Selection and Validation of Reliable Reference Genes for Gene Expression Studies in Different Genotypes and TRV-Infected Fruits of Peach (<i>Prunus persica</i> L. Batsch) during Ripening. <i>Genes</i> , 2022, 13, 160.	2.4	11
13	Effects of Chronic Dust Load On Leaf Pigments of the Landscape Plant <i>Murraya Paniculata</i> . <i>Gesunde Pflanzen</i> , 2019, 71, 249-258.	3.0	10
14	Epigenomic Regulatory Mechanism in Vegetative Phase Transition of <i>Malus hupehensis</i> . <i>Journal of Agricultural and Food Chemistry</i> , 2020, 68, 4812-4829.	5.2	10
15	Nitrate application affects root morphology by altering hormonal status and gene expression patterns in B9 apple rootstock nursery plants. <i>Fruit Research</i> , 2021, 1, 1-11.	2.0	10
16	Transcriptome Analysis Reveals New Insights into MdBAK1-Mediated Plant Growth in <i>Malus domestica</i> . <i>Journal of Agricultural and Food Chemistry</i> , 2019, 67, 9757-9771.	5.2	9
17	PpePL1 and PpePL15 Are the Core Members of the Pectate Lyase Gene Family Involved in Peach Fruit Ripening and Softening. <i>Frontiers in Plant Science</i> , 2022, 13, 844055.	3.6	9
18	Genome-wide identification of Gramineae histone modification genes and their potential roles in regulating wheat and maize growth and stress responses. <i>BMC Plant Biology</i> , 2021, 21, 543.	3.6	8

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19	Nitrate Application Induces Adventitious Root Growth by Regulating Gene Expression Patterns in Apple Rootstocks. <i>Journal of Plant Growth Regulation</i> , 2022, 41, 3467-3478.	5.1	7
20	Regulation of Flowering Time by Improving Leaf Health Markers and Expansion by Salicylic Acid Treatment: A New Approach to Induce Flowering in <i>Malus domestica</i> . <i>Frontiers in Plant Science</i> , 2021, 12, 655974.	3.6	6
21	PpSAUR43, an Auxin-Responsive Gene, Is Involved in the Post-Ripening and Softening of Peaches. <i>Horticulturae</i> , 2022, 8, 379.	2.8	3
22	Identification of MdMED family, key role of MdMED81, and salicylic acid at the right time of year triggers MdMED81 to induce flowering in <i>Malus domestica</i> . <i>Scientia Horticulturae</i> , 2022, 304, 111341.	3.6	3
23	A mini review fermentation and preservation: role of Lactic Acid Bacteria. <i>MOJ Food Processing & Technology</i> , 2018, 6, 414-417.	0.9	0