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## List of Publications by Year in descending order

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16  
papers

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1040056

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#	ARTICLE	IF	CITATIONS
1	Genetic diversity of the world's largest oil palm ( <i>Elaeis guineensis</i> Jacq.) field genebank accessions using microsatellite markers. <i>Genetic Resources and Crop Evolution</i> , 2015, 62, 349-360.	1.6	47
2	Sequence characterized amplified region markers: A reliable tool for adulterant detection in turmeric powder. <i>Food Research International</i> , 2011, 44, 2889-2895.	6.2	31
3	Sodium sulphite enhances RNA isolation and sensitivity of Cucumber mosaic virus detection by RT-PCR in black pepper. <i>Journal of Virological Methods</i> , 2007, 141, 107-110.	2.1	24
4	Development, Characterization and Cross Species Amplification of Polymorphic Microsatellite Markers from Expressed Sequence Tags of Turmeric ( <i>Curcuma longa</i> L.). <i>Molecular Biotechnology</i> , 2010, 44, 140-147.	2.4	24
5	Development, characterization and utilization of genomic microsatellite markers in turmeric ( <i>Curcuma longa</i> L.). <i>Biochemical Systematics and Ecology</i> , 2010, 38, 641-646.	1.3	21
6	SCAR markers for adulterant detection in ground chilli. <i>British Food Journal</i> , 2011, 113, 656-668.	2.9	21
7	Novel polymorphic microsatellite markers from turmeric, <i>Curcuma longa</i> L. (Zingiberaceae). <i>Acta Botanica Croatica</i> , 2013, 72, 407-412.	0.7	15
8	Identification and Characterization of a Badnavirus Infecting Betel Vine and Indian Long Pepper. <i>Journal of Plant Biochemistry and Biotechnology</i> , 2008, 17, 73-76.	1.7	14
9	Genetic homogeneity in <i>Jatropha curcas</i> L. individuals as revealed by microsatellite markers: implication to breeding strategies. <i>Revista Brasileira De Botanica</i> , 2016, 39, 861-868.	1.3	9
10	RAPD, SCAR and conserved 18S rDNA markers for a red-listed and endemic medicinal plant species, <i>Knema andamanica</i> (Myristicaceae). <i>Physiology and Molecular Biology of Plants</i> , 2013, 19, 245-250.	3.1	6
11	A gene co-expression network model identifies yield-related vicinity networks in <i>Jatropha curcas</i> shoot system. <i>Scientific Reports</i> , 2018, 8, 9211.	3.3	5
12	Transcriptome analysis of reproductive tissue differentiation in <i>Jatropha curcas</i> Linn.. <i>Genomics Data</i> , 2017, 13, 11-14.	1.3	4
13	An integration of phenotypic and transcriptomic data analysis reveals yield-related hub genes in <i>Jatropha curcas</i> inflorescence. <i>PLoS ONE</i> , 2018, 13, e0203441.	2.5	4
14	Genetic resources of Asian palmyrah palm ( <i>Borassus flabellifer</i> L.): a comprehensive review on diversity, characterization and utilization. <i>Plant Genetic Resources: Characterisation and Utilisation</i> , 2020, 18, 445-453.	0.8	2
15	The 18S rDNA gene discriminates between red-listed and unexplored ethnomedicinal species of Myristicaceae restricted to humid tropics of India. <i>Genetic Resources and Crop Evolution</i> , 2014, 61, 523-535.	1.6	1
16	RNA-seq data of the <i>Jatropha curcas</i> L. shoot system. <i>Data in Brief</i> , 2018, 21, 71-74.	1.0	1