## Yong-Bao Pan

List of Publications by Year in descending order

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567281 552781 34 764 15 26 citations h-index g-index papers 34 34 34 509 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Sugarcane microsatellites for the assessment of genetic diversity in sugarcane germplasm. Plant Science, 2003, 165, 181-189.	3.6	133
2	Highly polymorphic microsatellite dna markers for sugarcane germplasm evaluation and variety identity testing. Sugar Tech, 2006, 8, 246-256.	1.8	88
3	Genetic analysis of the sugarcane (Saccharum spp.) cultivar â€~LCP 85-384'. I. Linkage mapping using AFLP, SSR, and TRAP markers. Theoretical and Applied Genetics, 2011, 123, 77-93.	3.6	61
4	Genetic diversity and population structure analysis of Saccharum and Erianthus genera using microsatellite (SSR) markers. Scientific Reports, 2019, 9, 395.	3.3	47
5	Genetic Analysis of Diversity within a Chinese Local Sugarcane Germplasm Based on Start Codon Targeted Polymorphism. BioMed Research International, 2014, 2014, 1-10.	1.9	44
6	Biplot evaluation of test environments and identification of mega-environment for sugarcane cultivars in China. Scientific Reports, 2015, 5, 15505.	3.3	34
7	Databasing Molecular Identities of Sugarcane (Saccharum spp.) Clones Constructed with Microsatellite (SSR) DNA Markers. American Journal of Plant Sciences, 2010, 01, 87-94.	0.8	31
8	Genetic Diversity Analysis of Sugarcane Germplasm Based on Fluorescence-Labeled Simple Sequence Repeat Markers and a Capillary Electrophoresis-based Genotyping Platform. Sugar Tech, 2016, 18, 380-390.	1.8	30
9	Registration of â€~Ho 02â€113' Sugarcane. Journal of Plant Registrations, 2013, 7, 51-57.	0.5	30
10	Highly Polymorphic Microsatellite DNA Markers for Sugarcane Germplasm Evaluation and Variety Identity Testing. Sugar Tech, 2011, 13, 129-136.	1.8	28
11	Microsatellite (Simple Sequence Repeat) Marker–based Paternity Analysis of a Sevenâ€Parent Sugarcane Polycross. Crop Science, 2010, 50, 1401-1408.	1.8	23
12	Sugarcane Mosaic Disease: Characteristics, Identification and Control. Microorganisms, 2021, 9, 1984.	3.6	22
13	Rational regional distribution of sugarcane cultivars in China. Scientific Reports, 2015, 5, 15721.	3.3	21
14	Development and Use of Single Sequence Repeats (SSRs) Markers for Sugarcane Breeding and Genetic Studies. Agronomy, 2018, 8, 260.	3.0	21
15	SSR Marker-Assisted Management of Parental Germplasm in Sugarcane (Saccharum spp. hybrids) Breeding Programs. Agronomy, 2019, 9, 449.	3.0	20
16	Allelic divergence and cultivar-specific SSR alleles revealed by capillary electrophoresis using fluorescence-labeled SSR markers in sugarcane. Genome, 2014, 57, 363-372.	2.0	18
17	Registration of †Ho 00â€961†Sugarcane. Journal of Plant Registrations, 2011, 5, 332-338.	0.5	17
18	Genome-Wide Analysis of the Trehalose-6-Phosphate Synthase (TPS) Gene Family and Expression Profiling of ScTPS Genes in Sugarcane. Agronomy, 2020, 10, 969.	3.0	16

#	Article	IF	CITATIONS
19	Genetic Analysis of Resistance Gene Analogues from a Sugarcane Cultivar Resistant to Red Rot Disease. Journal of Phytopathology, 2015, 163, 755-763.	1.0	10
20	Development and Integration of an SSR-Based Molecular Identity Database into Sugarcane Breeding Program. Agronomy, 2016, 6, 28.	3.0	9
21	Identification of quantitative trait loci controlling sucrose content based on an enriched genetic linkage map of sugarcane (Saccharum spp. hybrids) cultivar †LCP 85-384'. Euphytica, 2016, 207, 527-549.	1.2	9
22	Development of an RAPD-based SCAR marker for smut disease resistance in commercial sugarcane cultivars of Pakistan. Crop Protection, 2017, 94, 166-172.	2.1	9
23	Cultivar Evaluation and Essential Test Locations Identification for Sugarcane Breeding in China. Scientific World Journal, The, 2014, 2014, 1-10.	2.1	7
24	Genetic Structure and Diversity of Parental Cultivars Involved in China Mainland Sugarcane Breeding Programs as Inferred from DNA Microsatellites. Journal of Integrative Agriculture, 2012, 11, 1794-1803.	3.5	6
25	Independently Segregating Simple Sequence Repeats (SSR) Alleles in Polyploid Sugarcane. Sugar Tech, 2015, 17, 235-242.	1.8	6
26	SSR-Based Genetic Identity of Sugarcane Clones and its Potential Application in Breeding and Variety Extension. Sugar Tech, 2020, 22, 367-378.	1.8	6
27	A Comparative Study of Three Detection Techniques for <i>Leifsonia xyli </i> Subsp. <i>xyli </i> , the Causal Pathogen of Sugarcane Ratoon Stunting Disease. BioMed Research International, 2018, 2018, 1-11.	1.9	4
28	Genotype-Specific Microsatellite (SSR) Markers for the Sugarcane Germplasm from the Karst Region of Guizhou, China. American Journal of Plant Sciences, 2016, 07, 2209-2220.	0.8	4
29	Population Structure and Genetic Diversity Analysis in Sugarcane (Saccharum spp. hybrids) and Six Related Saccharum Species. Agronomy, 2022, 12, 412.	3.0	4
30	Identification of quantitative trait loci (QTL) controlling fibre content of sugarcane ( Saccharum) Tj ETQq0 0 0 rgB	T/Qverloc	ck <sub>3</sub> 10 Tf 50 3
31	Biotechnology: Impact on Sugarcane Agriculture and Industry. Sugar Tech, 2012, 14, 1-2.	1.8	2
32	Molecular Cloning and Functional Analysis of an Ethylene Receptor Gene from Sugarcane (Saccharum) Tj ETQq0 0	)	)verlock 10 T
33	Application of Molecular Markers in Sugarcane Germplasm Innovation and Breeding: New Germplasm with Cytoplasm from Saccharum spontaneum. Sugar Tech, 2018, 20, 375-375.	1.8	0
34	Registration of â€~Ho 05â€961' sugarcane. Journal of Plant Registrations, 0, , .	0.5	0