

Namhee Jeong

List of Publications by Year in descending order

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17
papers

695
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840776

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849
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#	ARTICLE	IF	CITATIONS
1	Chromosomal features revealed by comparison of genetic maps of Glycine max and Glycine soja. <i>Genomics</i> , 2020, 112, 1481-1489.	2.9	8
2	Korean soybean core collection: Genotypic and phenotypic diversity population structure and genome-wide association study. <i>PLoS ONE</i> , 2019, 14, e0224074.	2.5	56
3	Genetic diversity patterns and domestication origin of soybean. <i>Theoretical and Applied Genetics</i> , 2019, 132, 1179-1193.	3.6	44
4	A System for Distinguishing Octoploid Strawberry Cultivars Using High-Throughput SNP Genotyping. <i>Tropical Plant Biology</i> , 2017, 10, 68-76.	1.9	12
5	Molecular Genetic Analysis of a Novel Recessive White Flower Gene in Wild Soybean. <i>Crop Science</i> , 2017, 57, 3027-3034.	1.8	1
6	Identification of haplotypes at the Rsv4 genomic region in soybean associated with durable resistance to soybean mosaic virus. <i>Theoretical and Applied Genetics</i> , 2016, 129, 453-468.	3.6	37
7	Development of Marker-free TaGlu-Ax1 Transgenic Rice Harboring a Wheat High-molecular-weight Glutenin Subunit (HMW-GS) Protein. <i>Journal of Life Science</i> , 2016, 26, 1121-1129.	0.2	3
8	Development, validation and genetic analysis of a large soybean SNP genotyping array. <i>Plant Journal</i> , 2015, 81, 625-636.	5.7	125
9	Multiple genes confer resistance to soybean mosaic virus in the soybean cultivar Hwangkeum. <i>Plant Genetic Resources: Characterisation and Utilisation</i> , 2014, 12, S41-S44.	0.8	7
10	Population Structure and Domestication Revealed by High-Depth Resequencing of Korean Cultivated and Wild Soybean Genomes. <i>DNA Research</i> , 2014, 21, 153-167.	3.4	92
11	Dynamic genetic features of chromosomes revealed by comparison of soybean genetic and sequence-based physical maps. <i>Theoretical and Applied Genetics</i> , 2013, 126, 1103-1119.	3.6	21
12	<i>Sh1</i> is a Key Regulator of Leaflet Shape and Number of Seeds per Pod in Soybean. <i>Plant Cell</i> , 2013, 24, 4807-4818.	6.6	90
13	Fine genetic mapping of the genomic region controlling leaflet shape and number of seeds per pod in the soybean. <i>Theoretical and Applied Genetics</i> , 2011, 122, 865-874.	3.6	52
14	Glycine max non-nodulation locus <i>rj1</i> : a recombinogenic region encompassing a SNP in a lysine motif receptor-like kinase (<i>GmNFR1L</i>). <i>Theoretical and Applied Genetics</i> , 2011, 122, 875-884.	3.6	7
15	Novel major quantitative trait loci regulating the content of isoflavone in soybean seeds. <i>Genes and Genomics</i> , 2011, 33, 685-692.	1.4	38
16	Genetic Analysis of Genes Controlling Natural Variation of Seed Coat and Flower Colors in Soybean. <i>Journal of Heredity</i> , 2010, 101, 757-768.	2.4	84
17	Genome structure in soybean revealed by a genomewide genetic map constructed from a single population. <i>Genomics</i> , 2008, 92, 52-59.	2.9	18