

Tadayuki Wako

List of Publications by Year in descending order

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papers

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citations

840776

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839539

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21
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21
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166
citing authors

#	ARTICLE	IF	CITATIONS
1	Construction of a high-density linkage map and graphical representation of the arrangement of transcriptome-based unigene markers on the chromosomes of onion, <i>Allium cepa</i> L. <i>BMC Genomics</i> , 2021, 22, 481.	2.8	13
2	QTL analysis of morphological traits and pseudostem pigmentation in bunching onion (<i>Allium</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 702	1.2	6
3	Mapping of quantitative trait loci for bolting time in bunching onion (<i>Allium fistulosum</i> L.). <i>Euphytica</i> , 2016, 209, 537-546.	1.2	7
4	Development of transcriptome shotgun assembly-derived markers in bunching onion (<i>Allium</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 622	2.1	27
5	Screening and incorporation of rust resistance from <i>Allium cepa</i> into bunching onion (<i>Allium fistulosum</i>) via alien chromosome addition. <i>Genome</i> , 2015, 58, 135-142.	2.0	7
6	Supplementation with Japanese bunching onion (<i>Allium fistulosum</i> L.) expressing a single alien chromosome from shallot increases the antioxidant activity of Kamaboko fish jelly paste in vitro. <i>Biomedical Reports</i> , 2013, 1, 355-358.	2.0	2
7	QTL analysis for pseudostem pungency in bunching onion (<i>Allium fistulosum</i>). <i>Molecular Breeding</i> , 2012, 30, 1689-1698.	2.1	7
8	Direct determination of the chromosomal location of bunching onion and bulb onion markers using bunching onion-shallot monosomic additions and allotriploid-bunching onion single alien deletions. <i>Theoretical and Applied Genetics</i> , 2011, 122, 501-510.	3.6	16
9	Development of microsatellite markers in cultivated and wild species of sections <i>Cepa</i> and <i>Phyllodolon</i> in <i>Allium</i> . <i>Euphytica</i> , 2010, 173, 321-328.	1.2	22
10	Inheritance mode of male sterility in bunching onion (<i>Allium fistulosum</i> L.) accessions. <i>Euphytica</i> , 2010, 173, 357-367.	1.2	15
11	Classification and identification of bunching onion (<i>Allium fistulosum</i>) varieties based on SSR markers. <i>Breeding Science</i> , 2010, 60, 139-152.	1.9	12
12	Mapping of Quantitative Trait Loci Controlling Seedling Growth in Bunching Onion (<i>Allium</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 302 To	0.8	7
13	SSR-tagged breeding scheme for allogamous crops: a trial in bunching onion (<i>Allium fistulosum</i>). <i>Euphytica</i> , 2009, 169, 327-334.	1.2	9
14	Molecular and biochemical identification of alien chromosome additions in shallot (<i>Allium cepa</i> L.) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 Genetic Systems, 2009, 84, 43-55.	0.7	11
15	Construction of SSR-based chromosome map in bunching onion (<i>Allium fistulosum</i>). <i>Theoretical and Applied Genetics</i> , 2008, 117, 1213-1223.	3.6	39
16	Detection of Textural Difference between Cultivars of Bunching Onion using the Device for Acoustic Measurement of Food Texture. <i>Japanese Society for Horticultural Science</i> , 2008, 77, 440-446.	0.8	3
17	Isolation of 1,796 SSR clones from SSR-enriched DNA libraries of bunching onion (<i>Allium fistulosum</i>). <i>Euphytica</i> , 2007, 157, 83-94.	1.2	28
18	Considerable Heterogeneity in Commercial F1 Varieties of Bunching Onion (<i>Allium fistulosum</i>) and Proposal of Breeding Scheme for Conferring Variety Traceability Using SSR Markers. <i>Breeding Science</i> , 2006, 56, 321-326.	1.9	22

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19	Chromosomal Locations of Microsatellites in Onion. Hortscience: A Publication of the American Society for Horticultural Science, 2006, 41, 315-318.	1.0	13
20	Genetic mapping of AFLP markers in Japanese bunching onion (<i>Allium fistulosum</i>). Euphytica, 2005, 144, 255-263.	1.2	34
21	Development of Microsatellite Markers in Bunching Onion (<i>Allium fistulosum</i> L.). Breeding Science, 2004, 54, 361-365.	1.9	30