Satoshi Natsume

List of Publications by Year in descending order

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623734 839539 3,645 19 14 18 citations g-index h-index papers 22 22 22 3870 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Genome sequencing reveals agronomically important loci in rice using MutMap. Nature Biotechnology, 2012, 30, 174-178.	17.5	1,087
2	<scp>QTL</scp> â€seq: rapid mapping of quantitative trait loci in rice by whole genome resequencing of <scp>DNA</scp> from two bulked populations. Plant Journal, 2013, 74, 174-183.	5.7	1,065
3	MutMap+: Genetic Mapping and Mutant Identification without Crossing in Rice. PLoS ONE, 2013, 8, e68529.	2.5	267
4	MutMap accelerates breeding of a salt-tolerant rice cultivar. Nature Biotechnology, 2015, 33, 445-449.	17.5	265
5	<scp>M</scp> ut <scp>M</scp> apâ€ <scp>G</scp> ap: wholeâ€genome resequencing of mutant <scp>F</scp> 2 progeny bulk combined with <i>de novo</i> assembly of gap regions identifies the rice blast resistance gene <scp><i>Pii</i> New Phytologist, 2013, 200, 276-283.</scp>	7.3	196
6	Host specialization of the blast fungus Magnaporthe oryzae is associated with dynamic gain and loss of genes linked to transposable elements. BMC Genomics, 2016, 17, 370.	2.8	157
7	Genome sequencing of the staple food crop white Guinea yam enables the development of a molecular marker for sex determination. BMC Biology, 2017, 15, 86.	3.8	114
8	A cytochrome P450, OsDSS1, is involved in growth and drought stress responses in rice (Oryza sativa) Tj ETQq0 C))verlock 10 T
9	The Draft Genome of Hop (Humulus lupulus), an Essence for Brewing. Plant and Cell Physiology, 2015, 56, 428-441.	3.1	111
10	The rice (Oryza sativa L.) LESION MIMIC RESEMBLING, which encodes an AAA-type ATPase, is implicated in defense response. Molecular Genetics and Genomics, 2015, 290, 611-622.	2.1	66
11	Coval: Improving Alignment Quality and Variant Calling Accuracy for Next-Generation Sequencing Data. PLoS ONE, 2013, 8, e75402.	2.5	58
12	Genome analyses reveal the hybrid origin of the staple crop white Guinea yam (<i>Dioscorea) Tj ETQq0 0 0 rgBT / 2020, 117, 31987-31992.</i>	Overlock : 7.1	10 Tf 50 307 40
13	A chloroplastâ€localized protein <scp>LESION AND LAMINA BENDING</scp> affects defence and growth responses in rice. New Phytologist, 2016, 210, 1282-1297.	7.3	33
14	High-performance pipeline for MutMap and QTL-seq. PeerJ, 2022, 10, e13170.	2.0	22
15	Identification of candidate flowering and sex genes in white Guinea yam (D. rotundata Poir.) by SuperSAGE transcriptome profiling. PLoS ONE, 2019, 14, e0216912.	2.5	16
16	Population Genomics of Yams: Evolution and Domestication of Dioscorea Species. Population Genomics, 2021, , .	0.5	13
17	Screening of rice mutants with improved saccharification efficiency results in the identification of CONSTITUTIVE PHOTOMORPHOGENIC 1 and GOLD HULL AND INTERNODE 1. Planta, 2017, 246, 61-74.	3.2	5
18	Next-Generation Breeding of Rice by Whole-Genome Approaches. , 2018, , 511-522.		4

SATOSHI NATSUME

#	ŧ	Article	IF	CITATIONS
1	.9	Whole genome sequencing and future breeding of rice. Journal of Plant Biochemistry and Biotechnology, 2012, 21, 10-14.	1.7	2