

Pankaj Kumar

List of Publications by Year in descending order

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15
papers

238
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1307594

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1125743

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16
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230
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#	ARTICLE	IF	CITATIONS
1	Development of EMS-induced mutation population for amylose and resistant starch variation in bread wheat (<i>Triticum aestivum</i>) and identification of candidate genes responsible for amylose variation. <i>BMC Plant Biology</i> , 2016, 16, 217.	3.6	54
2	Comparative Analysis of Phenolic Compound Characterization and Their Biosynthesis Genes between Two Diverse Bread Wheat (<i>Triticum aestivum</i>) Varieties Differing for Chapatti (Unleavened Flat Bread) Quality. <i>Frontiers in Plant Science</i> , 2016, 7, 1870.	3.6	48
3	Expression patterns of genes involved in starch biosynthesis during seed development in bread wheat (<i>Triticum aestivum</i>). <i>Molecular Breeding</i> , 2015, 35, 1.	2.1	32
4	Pivotal role of bZIPs in amylose biosynthesis by genome survey and transcriptome analysis in wheat (<i>Triticum aestivum</i> L.) mutants. <i>Scientific Reports</i> , 2018, 8, 17240.	3.3	30
5	Large-scale identification and characterization of phenolic compounds and their marker-trait association in wheat. <i>Euphytica</i> , 2020, 216, 1.	1.2	12
6	Novel intron length polymorphic (ILP) markers from starch biosynthesis genes reveal genetic relationships in Indian wheat varieties and related species. <i>Molecular Biology Reports</i> , 2020, 47, 3485-3500.	2.3	12
7	Genome-wide identification and expression profiling of basic leucine zipper transcription factors following abiotic stresses in potato (<i>Solanum tuberosum</i> L.). <i>PLoS ONE</i> , 2021, 16, e0247864.	2.5	9
8	Development and characterization of bZIP transcription factor based SSRs in wheat. <i>Gene</i> , 2020, 756, 144912.	2.2	8
9	Genome-wide analysis of RING-type E3 ligase family identifies potential candidates regulating high amylose starch biosynthesis in wheat (<i>Triticum aestivum</i> L.). <i>Scientific Reports</i> , 2021, 11, 11461.	3.3	8
10	Marker-trait association identified candidate starch biosynthesis pathway genes for starch and amylose-lipid complex gelatinization in wheat (<i>Triticum aestivum</i> L.). <i>Euphytica</i> , 2020, 216, 1.	1.2	7
11	Enhancement of chlorogenic content of the eggplant fruit with eggplant hydroxycinnamoyl CoA-quinase transferase gene via novel agroinfiltration protocol. <i>Pharmacognosy Magazine</i> , 2020, 16, 450.	0.6	6
12	Understanding the regulatory relationship of abscisic acid and bZIP transcription factors towards amylose biosynthesis in wheat. <i>Molecular Biology Reports</i> , 2021, 48, 2473-2483.	2.3	5
13	Unraveling novel and rare mutations for alpha-amylase and key transcription factors in EMS-induced wheat mutants for amylose by TILLING. <i>Molecular Biology Reports</i> , 2022, , 1.	2.3	2
14	Resistant starch: biosynthesis, regulatory pathways, and engineering via CRISPR system. , 2021, , 303-317.		1
15	<i>OsMATE6</i> gene putatively involved in host defense response toward susceptibility against <i>Rhizoctonia solani</i> in rice. <i>Journal of Plant Interactions</i> , 2022, 17, 744-755.	2.1	0