

Environmental stress activation of plant long-terminal

Functional Plant Biology

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Retrotransposon-based molecular markers for assessment of genomic diversity. <i>Functional Plant Biology</i> , 2014, 41, 781.	1.1	13
2	DKRE1â€”The first full-length Ty1-copia-like retrotransposon in persimmon: Isolation, characteristic and potential involvement in occurrence of bud mutations. <i>Scientia Horticulturae</i> , 2015, 184, 149-159.	1.7	5
3	Introgression of bacterial wilt resistance from <i>Solanum melongena</i> to <i>S. tuberosum</i> through asymmetric protoplast fusion. <i>Plant Cell, Tissue and Organ Culture</i> , 2016, 125, 433-443.	1.2	15
4	A highly specific microRNA-mediated mechanism silences LTR retrotransposons of strawberry. <i>Plant Journal</i> , 2016, 85, 70-82.	2.8	31
5	Partial sequencing reveals the transposable element composition of <i>Coffea</i> genomes and provides evidence for distinct evolutionary stories. <i>Molecular Genetics and Genomics</i> , 2016, 291, 1979-1990.	1.0	16
6	Useful parasites: the evolutionary biology and biotechnology applications of transposable elements. <i>Journal of Genetics</i> , 2016, 95, 1039-1052.	0.4	4
7	Rapid amplification of four retrotransposon families promoted speciation and genome size expansion in the genus <i>Panax</i> . <i>Scientific Reports</i> , 2017, 7, 9045.	1.6	24
8	Heterochromatic and cytomolecular diversification in the <i>Caesalpinia</i> group (Leguminosae): Relationships between phylogenetic and cytogeographical data. <i>Perspectives in Plant Ecology, Evolution and Systematics</i> , 2017, 29, 51-63.	1.1	30
9	Genome-wide analysis of salinity-stress induced DNA methylation alterations in cotton (<i>Gossypium</i>)	0.3	11
10	Short-term mutagenicity test by using IRAP molecular marker in rice grown under herbicide treatment. <i>Biotechnology and Biotechnological Equipment</i> , 2018, 32, 923-928.	0.5	11
11	Retrotransposons in Plant Genomes: Structure, Identification, and Classification through Bioinformatics and Machine Learning. <i>International Journal of Molecular Sciences</i> , 2019, 20, 3837.	1.8	56
12	Karyological traits related to phylogenetic signal and environmental conditions within the <i>Hymenaea</i> clade (Leguminosae, Detarioideae). <i>Perspectives in Plant Ecology, Evolution and Systematics</i> , 2019, 39, 125462.	1.1	10
13	LTR-TEs abundance, timing and mobility in <i>Solanum commersonii</i> and <i>S. tuberosum</i> genomes following cold-stress conditions. <i>Planta</i> , 2019, 250, 1781-1787.	1.6	25
14	Physiological, epigenetic and genetic regulation in some olive cultivars under salt stress. <i>Scientific Reports</i> , 2019, 9, 1093.	1.6	64
15	Zinc priming and foliar application enhances photoprotection mechanisms in drought-stressed wheat plants during anthesis. <i>Plant Physiology and Biochemistry</i> , 2019, 140, 27-42.	2.8	26
16	Epigenetic changes and their relationship to somaclonal variation: a need to monitor the micropropagation of plantation crops. <i>Functional Plant Biology</i> , 2020, 47, 508.	1.1	24
17	Inter-retrotransposon amplified polymorphism markers revealed long terminal repeat retrotransposon insertion polymorphism in flax cultivated on the experimental fields around Chernobyl. <i>Journal of Environmental Science and Health - Part A Toxic/Hazardous Substances and Environmental Engineering</i> , 2020, 55, 957-963.	0.9	9
18	Evaluation of wheat (<i>Triticum aestivum</i> L.) salt stress tolerance using physiological parameters and retrotransposon-based markers. <i>Genetic Resources and Crop Evolution</i> , 2021, 68, 227-242.	0.8	59

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19	Activation of plant LTR-retrotransposons under in vitro culture stress. <i>Visnik Ukrains Kogo Tovaristva Genetikiv i Selekcioneriv</i> , 2021, 18, 58-69.	0.4	0
20	InpactorDB: A Classified Lineage-Level Plant LTR Retrotransposon Reference Library for Free-Alignment Methods Based on Machine Learning. <i>Genes</i> , 2021, 12, 190.	1.0	14
21	Wheat omics: Classical breeding to new breeding technologies. <i>Saudi Journal of Biological Sciences</i> , 2021, 28, 1433-1444.	1.8	12
22	Genome-wide characterization of LTR retrotransposons in the non-model deep-sea annelid <i>Lamellibrachia luymesi</i> . <i>BMC Genomics</i> , 2021, 22, 466.	1.2	5
23	Complex Networks of Prion-Like Proteins Reveal Cross Talk Between Stress and Memory Pathways in Plants. <i>Frontiers in Plant Science</i> , 2021, 12, 707286.	1.7	13
24	Aluminum-Induced Changes on DNA Damage, DNA Methylation and LTR Retrotransposon Polymorphism in Maize. <i>Arabian Journal for Science and Engineering</i> , 2018, 43, 123-131.	1.7	34
26	Bridging the Gap Between Environmental Adversity and Neuropsychiatric Disorders: The Role of Transposable Elements. <i>Frontiers in Genetics</i> , 0, 13, .	1.1	6
28	The role of LTR retrotransposons in plant genetic engineering: how to control their transposition in the genome. <i>Plant Cell Reports</i> , 2023, 42, 3-15.	2.8	4
29	The mechanisms underpinning lateral gene transfer between grasses. <i>Plants People Planet</i> , 2023, 5, 672-682.	1.6	3