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Retrotransposon-based molecular markers for assessment of genomic diversity

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#	Paper	IF	Citations
13	DKRE1IThe first full-length Ty1-copia-like retrotransposon in persimmon: Isolation, characteristic and potential involvement in occurrence of bud mutations. <i>Scientia Horticulturae</i> , <b>2015</b> , 184, 149-159	4.1	5
12	: Emerging Regulatory Roles for Transposons in Plant Stress Response. <i>Frontiers in Plant Science</i> , <b>2016</b> , 7, 1448	6.2	72
11	Genetic diversity in populations of the medicinal plant Leonurus cardiaca L. revealed by inter-primer binding site (iPBS) markers. <i>Genetic Resources and Crop Evolution</i> , <b>2017</b> , 64, 479-492	2	10
10	Genetic finger printing of salt- and drought-tolerant cotton cultivars (Gossypium hirsutum) by IRAP-REMAP and SRAP molecular markers. <i>Plant Gene</i> , <b>2018</b> , 14, 12-19	3.1	1
9	Genetic fingerprinting of diploid and tetraploid cotton cultivars by retrotransposon-based markers. <i>Nucleus (India)</i> , <b>2018</b> , 61, 137-143	1.7	2
8	Application of ITAP-PCR Techniques to Assess the Genetic Variability of Selected Cultivars of Winter Triticale (Triticosecale Wittmack). <i>Notulae Botanicae Horti Agrobotanici Cluj-Napoca</i> , <b>2019</b> , 47, 947-953	1.2	
7	Evaluation of wheat (Triticum aestivum L.) salt stress tolerance using physiological parameters and retrotransposon-based markers. <i>Genetic Resources and Crop Evolution</i> , <b>2021</b> , 68, 227-242	2	27
6	Retrotransposon-based genetic variation and population structure of Impatiens macrovexilla Y. L. Chen in natural habitats and the implications for breeding. <i>Scientia Horticulturae</i> , <b>2021</b> , 276, 109753	4.1	3
5	Genetic stability analysis of tissue culture derived date palm cv. Dedhi plants using IRAP markers. <i>Acta Ecologica Sinica</i> , <b>2021</b> , 42, 76-76	2.7	1
4	Draft genome sequence of the pulse crop blackgram [Vigna mungo (L.) Hepper] reveals potential R-genes. <i>Scientific Reports</i> , <b>2021</b> , 11, 11247	4.9	6
3	Genetic analysis and population structure of wild and cultivated wishbone flower (Lind.) lines related to specific floral color. <i>PeerJ</i> , <b>2021</b> , 9, e11702	3.1	
2	Deciphering the genetic diversity and population structure of Turkish bread wheat germplasm using iPBS-retrotransposons markers. <i>Molecular Biology Reports</i> , <b>2021</b> , 48, 6739-6748	2.8	2
1	Draft genome sequence of the pulse crop blackgram [Vigna mungo (L.) Hepper] reveals potential R-genes.		6