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Fast and sensitive silver staining of DNA in polyacrylamide gels

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2277	DNA amplification fingerprinting: A strategy for genome analysis. 1991 , 9, 294-307		73
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550	Genetic Evidence for Diversity of Spiralling Whitefly, <i>Aleurodicus dispersus</i> (Hemiptera: Aleyrodidae) Populations in India. 2014 , 97, 1115-1122		13
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544	Identification of QTL associated with flower and runner production in octoploid strawberry (<i>Fragaria</i> × <i>nanassa</i>). 2015 , 5, 107-116		8
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510	Genetic analysis and molecular mapping of resistance gene to <i>Phakopsora pachyrhizi</i> in soybean germplasm SX6907. <i>Theoretical and Applied Genetics</i> , 2015 , 128, 733-43	6	10
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464	Development and characterization of microsatellite markers from an enriched genomic library of <i>Saccharina japonica</i> . 2015 , 27, 479-487		13
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436	Diversity and genetic structure of cassava landraces and their wild relatives (<i>Manihot</i> spp.) in Colombia revealed by simple sequence repeats. 2016 , 14, 200-210		5
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408	Extensive length variation in the ribosomal DNA intergenic spacer of yellow perch (<i>Perca flavescens</i>). 2016 , 59, 149-58		3

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400	Rich haplotypes of Viviparous-1 in <i>Triticum aestivum</i> subsp. <i>spelta</i> with different abscisic acid sensitivities. 2017 , 97, 497-504		4
399	Reevaluation of Two Quantitative Trait Loci for Type II Resistance to Fusarium Head Blight in Wheat Germplasm PI 672538. 2017 , 107, 92-99		13
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390	The effect of four microbial products on production performance and water quality in integrated culture of freshwater pearl mussel and fishes. 2017 , 48, 4897-4909		7

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388	De novo assembly of transcriptome from <i>Rhododendron latoucheae</i> Franch. using Illumina sequencing and development of new EST-SSR markers for genetic diversity analysis in <i>Rhododendron</i> . 2017 , 13, 1	27
387	Genetic diversity analysis of Iranian melon (<i>Cucumis melo</i> L.) accessions using ISSR markers. 2017 , 37-44	1
386	Assessment of genetic diversity on snake melon (<i>Cucumis melo</i> var. <i>flexuosus</i>) using ISSR markers in Iran. 2017 , 45-50	2
385	Genetic analysis and molecular mapping of stripe rust resistance genes in Chinese native wheat (<i>Triticum aestivum</i>) Lankao 5. 2017 , 46, 213-221	2
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372	Activity, biomass and composition of microbial communities and their degradation pathways in exposed propazine soil. 2017 , 145, 398-407	10

371	Assessment of genetic diversity in cowpea (<i>Vigna unguiculata</i> L.) germplasm using morphological and molecular characterisation. 2017 , 3, 1327092		11
370	SSR-enriched genetic linkage maps of bermudagrass (<i>Cynodon dactylon</i> Transvaalensis), and their comparison with allied plant genomes. <i>Theoretical and Applied Genetics</i> , 2017 , 130, 819-839	6	7
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