

# Phylogenetic analysis of organellar DNA sequences in th

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Chromosome assortment in <i>Saccharum</i> . <i>Theoretical and Applied Genetics</i> , 1994, 89-89, 959-963.	3.6	29
2	Genetic analysis of agronomic traits in a cross between sugarcane ( <i>Saccharum officinarum</i> L.) and its presumed progenitor ( <i>S. robustum</i> Brandes & Jesw. ex Grassl). <i>Molecular Breeding</i> , 1995, 1, 355-363.	2.1	38
3	Ribosomal DNA variations in <i>Erianthus</i> , a wild sugarcane relative (Andropogoneae-Saccharinae). <i>Theoretical and Applied Genetics</i> , 1996, 92, 733-743.	3.6	37
4	Comparative mapping of Andropogoneae: <i>Saccharum</i> L. (sugarcane) and its relation to sorghum and maize. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1997, 94, 14261-14266.	7.1	119
5	Genetic diversity of North American and Old World <i>Saccharum</i> assessed by RAPD analysis. <i>Genetic Resources and Crop Evolution</i> , 1997, 44, 235-240.	1.6	18
6	Title is missing!. <i>Euphytica</i> , 1997, 93, 283-292.	1.2	52
7	Isolation and characterisation of repeated DNA sequences from <i>Erianthus</i> spp. (Saccharinae: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 502	2.0	9
8	cpDNA and mtDNA Primers in Plants. , 1998, , 256-261.		12
9	Isolation and characterization of a satellite DNA family in the <i>Saccharum</i> complex. <i>Genome</i> , 1998, 41, 854-864.	2.0	57
10	Genetics, phylogenetics, and comparative genetics of <i>Saccharum</i> L., a polysomic polyploid Poales: Andropogoneae. <i>Biotechnology Annual Review</i> , 1998, 4, 285-319.	2.1	0
11	Analysis of genetic diversity and phylogeny in <i>Saccharum</i> and related genera using RAPD markers. <i>Genetic Resources and Crop Evolution</i> , 1999, 46, 73-79.	1.6	95
12	Detection of episomal banana streak badnavirus by IC-PCR. <i>Journal of Virological Methods</i> , 1999, 79, 1-8.	2.1	72
13	Molecular Characterization of the Sugarcane Variability for Genetic Improvement. <i>Developments in Plant Genetics and Breeding</i> , 2000, 5, 49-61.	0.6	0
14	Distribution and sequence analysis of the centromere-associated repetitive element CEN38 of <i>Sorghum bicolor</i> (Poaceae). <i>American Journal of Botany</i> , 2000, 87, 1757-1764.	1.7	35
15	Comparative Genomics of Plant Chromosomes. <i>Plant Cell</i> , 2000, 12, 1523.	6.6	1
16	Comparative Genomics of Plant Chromosomes. <i>Plant Cell</i> , 2000, 12, 1523-1539.	6.6	301
18	Genomics and plant breeding. <i>Biotechnology Annual Review</i> , 2001, 7, 195-238.	2.1	7
19	A search for markers of sugarcane evolution. <i>Genetics and Molecular Biology</i> , 2001, 24, 169-174.	1.3	5

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20	The use of ribosomal ITS to determine phylogenetic relationships within Sorghum. <i>Plant Systematics and Evolution</i> , 2001, 230, 97-110.	0.9	19
21	Construction of a <i>Saccharum</i> Consensus Genetic Map from Two Interspecific Crosses. <i>Crop Science</i> , 2002, 42, 570-583.	1.8	40
22	Phylogenetics of <i>Miscanthus</i> , <i>Saccharum</i> and related genera (Saccharinae, Andropogoneae, Poaceae) based on DNA sequences from ITS nuclear ribosomal DNA and plastid trnL intron and trnL-F intergenic spacers. <i>Journal of Plant Research</i> , 2002, 115, 381-392.	2.4	246
23	Genetic diversity among wild sugarcane germplasm from Laos revealed with markers. <i>Euphytica</i> , 2002, 123, 121-130.	1.2	14
24	Oligoclonal interspecific origin of 'North Indian' and 'Chinese' sugarcanes. <i>Chromosome Research</i> , 2002, 10, 253-262.	2.2	101
25	Title is missing!. <i>Euphytica</i> , 2002, 127, 219-225.	1.2	52
26	Regeneration of citrus sinensis (+) clausena lansium intergeneric triploid and tetraploid somatic hybrids and their identification by molecular markers. <i>In Vitro Cellular and Developmental Biology - Plant</i> , 2003, 39, 360-364.	2.1	23
27	Molecular characterization of cytoplasmic and nuclear genomes in phenotypically abnormal Valencia orange ( <i>Citrus sinensis</i> ) + Meiwa kumquat ( <i>Fortunella crassifolia</i> ) intergeneric somatic hybrids. <i>Plant Cell Reports</i> , 2003, 21, 445-451.	5.6	53
28	Evaluation of maize microsatellite markers for genetic diversity analysis and fingerprinting in sugarcane. <i>Genome</i> , 2003, 46, 394-403.	2.0	73
29	Sugarcane microsatellites for the assessment of genetic diversity in sugarcane germplasm. <i>Plant Science</i> , 2003, 165, 181-189.	3.6	133
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35	Species-specific accumulation of interspersed sequences in genus <i>Saccharum</i> . <i>Genes and Genetic Systems</i> , 2004, 79, 361-365.	0.7	4
36	Issues of Fertility or Potential for Fertility in Oats, Olives, the Vigna Group, Ryegrass Species, Safflower, and Sugarcane. , 2005, , 231-255.		8
37	Very close relationship of the chloroplast genomes among <i>Saccharum</i> species. <i>Theoretical and Applied Genetics</i> , 2005, 110, 1523-1529.	3.6	29

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39	Morphological and molecular analysis of common millet ( <i>P. miliaceum</i> ) cultivars compared to an aDNA sample from the 15th century (Hungary). Euphytica, 2005, 146, 77-85.	1.2	34
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50	Identification of QTL for sugar-related traits in a sweet grain sorghum ( <i>Sorghum bicolor</i> L. Moench) recombinant inbred population. Molecular Breeding, 2008, 22, 367-384.	2.1	138
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