Hans de Jong

List of Publications by Year in descending order

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218592 206029 2,505 51 26 48 h-index citations g-index papers 68 68 68 3079 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	2D morphometric analysis of Arabidopsis thaliana nuclei reveals characteristic profiles of different cell types and accessions. Chromosome Research, 2022, 30, 5-24.	1.0	7
2	Is partial desynapsis in cauliflower (Brassica oleracea L. var. botrytis) pollen mother cells linked to aneuploidy in the crop?. Euphytica, 2022, 218, .	0.6	2
3	Male meiosis and pollen morphology in diploid Indonesian wild bananas and cultivars. Nucleus (India), 2021, 64, 181-191.	0.9	7
4	Meiotic recombination profiling of interspecific hybrid F1 tomato pollen by linked read sequencing. Plant Journal, 2020, 102, 480-492.	2.8	14
5	Meiotic crossover reduction by virusâ€induced gene silencing enables the efficient generation of chromosome substitution lines and reverse breeding in <i>Arabidopsis thaliana</i> . Plant Journal, 2020, 104, 1437-1452.	2.8	6
6	Genetic mapping of Fusarium wilt resistance in a wild banana Musa acuminata ssp. malaccensis accession. Theoretical and Applied Genetics, 2020, 133, 3409-3418.	1.8	35
7	Cytogenetics of structural rearrangements in Musa hybrids and cultivars. Burleigh Dodds Series in Agricultural Science, 2020, , 31-58.	0.1	1
8	Comparative analysis of repetitive sequences among species from the potato and the tomato clades. Annals of Botany, 2019, 123, 521-532.	1.4	36
9	Intact DNA purified from flow-sorted nuclei unlocks the potential of next-generation genome mapping and assembly in Solanum species. MethodsX, 2018, 5, 328-336.	0.7	3
10	Introgressive Hybridization in Potato Revealed by Novel Cytogenetic and Genomic Technologies. American Journal of Potato Research, 2018, 95, 607-621.	0.5	13
11	Genetic Dissection of Morphometric Traits Reveals That Phytochrome B Affects Nucleus Size and Heterochromatin Organization in <i>Arabidopsis thaliana</i> 2519-2531.	0.8	14
12	Optimization of Cell Spreading and Image Quality for the Study of Chromosomes in Plant Tissues. Methods in Molecular Biology, 2017, 1669, 141-158.	0.4	6
13	Two reported cytotypes of the emergent orchid model species Erycina pusilla are two different species. Euphytica, 2017, 213, 1.	0.6	4
14	Pairing analysis and in situ Hybridisation reveal autopolyploid-like behaviour in Solanum commersoniiÂ×ÂS. tuberosum (potato) interspecific hybrids. Euphytica, 2017, 213, 1.	0.6	19
15	Collinearity between potato (Solanum tuberosum L.) and wild relatives assessed by comparative cytogenetic mapping. Genome, 2017, 60, 228-240.	0.9	11
16	Molecular, genetic and evolutionary analysis of a paracentric inversion in <i>Arabidopsis thaliana</i> . Plant Journal, 2016, 88, 159-178.	2.8	81
17	Meiotic analysis and FISH with rDNA and rice BAC probes of the Thai KPS 01-01-25 sugarcane cultivar. Plant Systematics and Evolution, 2016, 302, 305-317.	0.3	10
18	Cnidaria: fast, reference-free clustering of raw and assembled genome and transcriptome NGS data. BMC Bioinformatics, 2015, 16, 352.	1.2	11

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19	Karyotype evolution in apomictic <i>Boechera</i> and the origin of the aberrant chromosomes. Plant Journal, 2015, 82, 785-793.	2.8	42
20	Introgression browser: highâ€ŧhroughput wholeâ€genome <scp>SNP</scp> visualization. Plant Journal, 2015, 82, 174-182.	2.8	17
21	Homologues of potato chromosome 5 show variable collinearity in the euchromatin, but dramatic absence of sequence similarity in the pericentromeric heterochromatin. BMC Genomics, 2015, 16, 374.	1.2	15
22	Exploring genetic variation in the tomato (<i>Solanum</i> section <i>Lycopersicon</i>) clade by wholeâ€genome sequencing. Plant Journal, 2014, 80, 136-148.	2.8	397
23	Fluorescence <i>In Situ</i> Hybridization and Optical Mapping to Correct Scaffold Arrangement in the Tomato Genome. G3: Genes, Genomes, Genetics, 2014, 4, 1395-1405.	0.8	81
24	Hybrid recreation by reverse breeding in Arabidopsis thaliana. Nature Protocols, 2014, 9, 761-772.	5.5	37
25	<i>DELAY OF GERMINATION $1 < i>$ mediates a conserved coat-dormancy mechanism for the temperature- and gibberellin-dependent control of seed germination. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E3571-80.</i>	3.3	175
26	Chromosomal organizations of major repeat families on potato (Solanum tuberosum) and further exploring in its sequenced genome. Molecular Genetics and Genomics, 2014, 289, 1307-1319.	1.0	16
27	Fine mapping of the tomato yellow leaf curl virus resistance gene Ty-2 on chromosome 11 of tomato. Molecular Breeding, 2014, 34, 749-760.	1.0	95
28	Epigenetic changes and transposon reactivation in Thai rice hybrids. Molecular Breeding, 2013, 31, 815-827.	1.0	12
29	Chromosome evolution in <i>Solanum</i> traced by crossâ€species BACâ€FISH. New Phytologist, 2012, 195, 688-698.	3.5	64
30	Structural homology in the Solanaceae: analysis of genomic regions in support of synteny studies in tomato, potato and pepper. Plant Journal, 2012, 71, 602-614.	2.8	40
31	From nucleosome to chromosome: a dynamic organization of genetic information. Plant Journal, 2011, 66, 4-17.	2.8	83
32	Chromosomal rearrangements between tomato and <i>Solanum chilense</i> hamper mapping and breeding of the TYLCV resistance gene <i>Tyâ€1</i> Plant Journal, 2011, 68, 1093-1103.	2.8	96
33	Comparison of the chromosome maps around a resistance hot spot on chromosome 5 of potato and tomato using BAC-FISH painting. Genome, 2010, 53, 103-110.	0.9	7
34	A Snapshot of the Emerging Tomato Genome Sequence. Plant Genome, 2009, 2, .	1.6	73
35	Assignment of genetic linkage maps to diploid Solanum tuberosum pachytene chromosomes by BAC-FISH technology. Chromosome Research, 2009, 17, 899-915.	1.0	44
36	The potential of high-resolution BAC-FISH in banana breeding. Euphytica, 2009, 166, 431-443.	0.6	25

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37	FISH mapping and molecular organization of the major repetitive sequences of tomato. Chromosome Research, 2008, 16, 919-933.	1.0	69
38	Map- vs. homology-based cloning for the recessive gene ol-2 conferring resistance to tomato powdery mildew. Euphytica, 2008, 162, 91-98.	0.6	24
39	A new wholeâ€mount DNA quantification method and the analysis of nuclear DNA content in the stemâ€cell niche of Arabidopsis roots. Plant Journal, 2008, 55, 886-894.	2.8	8
40	Highâ€resolution chromosome mapping of BACs using multiâ€colour FISH and pooledâ€BAC FISH as a backbone for sequencing tomato chromosomeâ€f6. Plant Journal, 2008, 56, 627-637.	2.8	82
41	Managing meiotic recombination in plant breeding. Trends in Plant Science, 2008, 13, 640-646.	4.3	132
42	Cross-Species Bacterial Artificial Chromosome–Fluorescence in Situ Hybridization Painting of the Tomato and Potato Chromosome 6 Reveals Undescribed Chromosomal Rearrangements. Genetics, 2008, 180, 1319-1328.	1.2	78
43	Fluorescence In Situ Hybridization on Medicago truncatula Chromosomes. , 2008, , 371-383.		0
44	Characterization of the centromere and peri-centromere retrotransposons in Brassica rapa and their distribution in related Brassica species. Plant Journal, 2007, 49, 173-183.	2.8	116
45	Use of the SSLP-based method for detection of rare apomictic events in a sexual AtSERK1 transgenic Arabidopsis population. Sexual Plant Reproduction, 2006, 19, 73-82.	2.2	3
46	Molecular cytogenetics and DNA sequence analysis of an apomixis-linked BAC in Paspalum simplex reveal a non pericentromere location and partial microcolinearity with rice. Theoretical and Applied Genetics, 2006, 112, 1179-1191.	1.8	90
47	NOR activity and repeat sequences of the paternal sex ratio chromosome of the parasitoid wasp Trichogramma kaykai. Chromosoma, 2005, 114, 410-419.	1.0	39
48	Characterization of rDNAs and tandem repeats in the heterochromatin of Brassica rapa. Molecules and Cells, 2005, 19, 436-44.	1.0	70
49	Cytogenetic tools for Arabidopsis thaliana. Chromosome Research, 2003, 11, 183-194.	1.0	64
50	Visualizing DNA domains and sequences by microscopy: a fifty-year history of molecular cytogenetics. Genome, 2003, 46, 943-946.	0.9	30
51	FISH studies reveal the molecular and chromosomal organization of individual telomere domains in tomato. Plant Journal, 1998, 13, 507-517.	2.8	97