

# Ben Vosman

## List of Publications by Year in descending order

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133  
papers

9,081  
citations

34016

52  
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46693

89  
g-index

135  
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135  
docs citations

135  
times ranked

8883  
citing authors

#	ARTICLE	IF	CITATIONS
1	WRKY Gene Family Drives Dormancy Release in Onion Bulbs. <i>Cells</i> , 2022, 11, 1100.	1.8	6
2	A novel non-trichome based whitefly resistance QTL in <i>Solanum galapagense</i> . <i>Euphytica</i> , 2021, 217, 1.	0.6	7
3	Fine mapping of a thrips resistance QTL in <i>Capsicum</i> and the role of diterpene glycosides in the underlying mechanism. <i>Theoretical and Applied Genetics</i> , 2021, 134, 1557-1573.	1.8	5
4	Insights from the first genome assembly of Onion ( <i>Allium cepa</i> ). <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	0.8	32
5	Aphid populations showing differential levels of virulence on <i>Capsicum</i> accessions. <i>Insect Science</i> , 2020, 27, 336-348.	1.5	10
6	Aphid resistance in <i>Capsicum</i> maps to a locus containing LRR-RLK gene analogues. <i>Theoretical and Applied Genetics</i> , 2020, 133, 227-237.	1.8	15
7	The effect of a thrips resistance QTL in different <i>Capsicum</i> backgrounds. <i>Euphytica</i> , 2020, 216, 1.	0.6	3
8	The ability to manipulate ROS metabolism in pepper may affect aphid virulence. <i>Horticulture Research</i> , 2020, 7, 6.	2.9	10
9	Genetic variation in phytochemicals in leaves of pepper ( <i>Capsicum</i> ) in relation to thrips resistance. <i>Arthropod-Plant Interactions</i> , 2019, 13, 1-9.	0.5	18
10	The effect of plant development on thrips resistance in <i>Capsicum</i> . <i>Arthropod-Plant Interactions</i> , 2019, 13, 11-18.	0.5	9
11	QTL mapping of insect resistance components of <i>Solanum galapagense</i> . <i>Theoretical and Applied Genetics</i> , 2019, 132, 531-541.	1.8	37
12	Broad spectrum insect resistance and metabolites in close relatives of the cultivated tomato. <i>Euphytica</i> , 2018, 214, 46.	0.6	40
13	Combining QTL mapping with transcriptome and metabolome profiling reveals a possible role for ABA signaling in resistance against the cabbage whitefly in cabbage. <i>PLoS ONE</i> , 2018, 13, e0206103.	1.1	13
14	Reduced phloem uptake of <i>Myzus persicae</i> on an aphid resistant pepper accession. <i>BMC Plant Biology</i> , 2018, 18, 138.	1.6	46
15	Genetic architecture of plant stress resistance: multi-trait genome-wide association mapping. <i>New Phytologist</i> , 2017, 213, 1346-1362.	3.5	144
16	Antibiosis resistance against larval cabbage root fly, <i>Delia radicum</i> , in wild Brassica-species. <i>Euphytica</i> , 2016, 211, 139-155.	0.6	18
17	Quantitative resistance against <i>Bemisia tabaci</i> in <i>Solanum pennellii</i> : Genetics and metabolomics. <i>Journal of Integrative Plant Biology</i> , 2016, 58, 397-412.	4.1	19
18	Defence against vertebrate herbivores trades off into architectural and low nutrient strategies amongst savanna Fabaceae species. <i>Oikos</i> , 2016, 125, 126-136.	1.2	32

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19	SNP-markers in <i>Allium</i> species to facilitate introgression breeding in onion. <i>BMC Plant Biology</i> , 2016, 16, 187.	1.6	40
20	Managing the Colorado potato beetle; the need for resistance breeding. <i>Euphytica</i> , 2015, 204, 487-501.	0.6	32
21	Host plant resistance towards the cabbage whitefly in <i>Brassica oleracea</i> and its wild relatives. <i>Euphytica</i> , 2015, 202, 297-306.	0.6	21
22	Parasitism overrides herbivore identity allowing hyperparasitoids to locate their parasitoid host using herbivore-induced plant volatiles. <i>Molecular Ecology</i> , 2015, 24, 2886-2899.	2.0	40
23	QTL mapping of thrips resistance in pepper. <i>Theoretical and Applied Genetics</i> , 2015, 128, 1945-1956.	1.8	20
24	Novel Genes Affecting the Interaction between the Cabbage Whitefly and <i>Arabidopsis</i> Uncovered by Genome-Wide Association Mapping. <i>PLoS ONE</i> , 2015, 10, e0145124.	1.1	9
25	Response of <i>Solanum tuberosum</i> to <i>Myzus persicae</i> infestation at different stages of foliage maturity. <i>Insect Science</i> , 2014, 21, 727-740.	1.5	8
26	Normal adult survival but reduced <i>Bemisia tabaci</i> oviposition rate on tomato lines carrying an introgression from <i>S. habrochaites</i> . <i>BMC Genetics</i> , 2014, 15, 142.	2.7	17
27	Constitutive overexpression of the pollen specific gene SKS13 in leaves reduces aphid performance on <i>Arabidopsis thaliana</i> . <i>BMC Plant Biology</i> , 2014, 14, 217.	1.6	10
28	Mining the Genus <i>Solanum</i> for Increasing Disease Resistance. , 2014, , 27-46.		10
29	Comparative analysis of <i>Solanum stoloniferum</i> responses to probing by the green peach aphid <i>Myzus persicae</i> and the potato aphid <i>Macrosiphum euphorbiae</i> . <i>Insect Science</i> , 2013, 20, 207-227.	1.5	30
30	Tomato breeding in the genomics era: insights from a SNP array. <i>BMC Genomics</i> , 2013, 14, 354.	1.2	86
31	Identification and QTL mapping of whitefly resistance components in <i>Solanum galapagense</i> . <i>Theoretical and Applied Genetics</i> , 2013, 126, 1487-1501.	1.8	66
32	Differences in insect resistance between tomato species endemic to the Galapagos Islands. <i>BMC Evolutionary Biology</i> , 2013, 13, 175.	3.2	45
33	The <i>Bemisia tabaci</i> species complex: Additions from different parts of the world. <i>Insect Science</i> , 2013, 20, 723-733.	1.5	94
34	Large subclonal variation in <i>Phytophthora infestans</i> populations associated with Ecuadorian potato landraces. <i>Plant Pathology</i> , 2013, 62, 1081-1088.	1.2	31
35	QualitySNPng: a user-friendly SNP detection and visualization tool. <i>Nucleic Acids Research</i> , 2013, 41, W587-W590.	6.5	28
36	Overexpression of IRM1 Enhances Resistance to Aphids in <i>Arabidopsis thaliana</i> . <i>PLoS ONE</i> , 2013, 8, e70914.	1.1	24

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37	Nutrients, technological properties and genetic relationships among twenty cowpea landraces cultivated in West Africa. <i>International Journal of Food Science and Technology</i> , 2012, 47, 2636-2647.	1.3	12
38	Resistance to <i>Bemisia tabaci</i> in tomato wild relatives. <i>Euphytica</i> , 2012, 187, 31-45.	0.6	93
39	High throughput phenotyping for aphid resistance in large plant collections. <i>Plant Methods</i> , 2012, 8, 33.	1.9	23
40	Phloem-specific resistance in <i>Brassica oleracea</i> against the whitefly <i>Aleyrodes proletella</i> . <i>Entomologia Experimentalis Et Applicata</i> , 2012, 142, 153-164.	0.7	23
41	Resistance factors in pepper inhibit larval development of thrips ( <i>Frankliniella</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 T	0.7	38
42	Identification of silverleaf whitefly resistance in pepper. <i>Plant Breeding</i> , 2011, 130, 708-714.	1.0	34
43	Exploiting natural variation to identify insect-resistance genes. <i>Plant Biotechnology Journal</i> , 2011, 9, 819-825.	4.1	95
44	Transcriptional responses of <i>Brassica nigra</i> to feeding by specialist insects of different feeding guilds. <i>Insect Science</i> , 2011, 18, 259-272.	1.5	30
45	Screening of pepper accessions for resistance against two thrips species ( <i>Frankliniella occidentalis</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 T	0.6	46
46	In vitro screening and QTL analysis for drought tolerance in diploid potato. <i>Euphytica</i> , 2011, 181, 357-369.	0.6	51
47	Construction of an integrated microsatellite and key morphological characteristic database of potato varieties on the EU common catalogue. <i>Euphytica</i> , 2011, 182, 239.	0.6	53
48	SolRgene: an online database to explore disease resistance genes in tuber-bearing <i>Solanum</i> species. <i>BMC Plant Biology</i> , 2011, 11, 116.	1.6	38
49	Genotype calling in tetraploid species from bi-allelic marker data using mixture models. <i>BMC Bioinformatics</i> , 2011, 12, 172.	1.2	175
50	What's in a name; Genetic structure in <i>Solanum</i> section <i>Petota</i> studied using population-genetic tools. <i>BMC Evolutionary Biology</i> , 2011, 11, 42.	3.2	38
51	Diversity, Distribution, and Evolution of <i>Solanum bulbocastanum</i> Late Blight Resistance Genes. <i>Molecular Plant-Microbe Interactions</i> , 2010, 23, 1206-1216.	1.4	69
52	Development and evaluation of robust molecular markers linked to disease resistance in tomato for distinctness, uniformity and stability testing. <i>Theoretical and Applied Genetics</i> , 2010, 120, 655-664.	1.8	84
53	A novel approach to locate <i>Phytophthora infestans</i> resistance genes on the potato genetic map. <i>Theoretical and Applied Genetics</i> , 2010, 120, 785-796.	1.8	49
54	A pipeline for high throughput detection and mapping of SNPs from EST databases. <i>Molecular Breeding</i> , 2010, 26, 65-75.	1.0	51

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55	Characterisation of sugar beet ( <i>Beta vulgaris</i> L. ssp. <i>vulgaris</i> ) varieties using microsatellite markers. <i>BMC Genetics</i> , 2010, 11, 41.	2.7	51
56	Development of microsatellite markers for identifying Brazilian <i>Coffea arabica</i> varieties. <i>Genetics and Molecular Biology</i> , 2010, 33, 507-514.	0.6	20
57	Intraspecific variation in herbivore community composition and transcriptional profiles in field-grown <i>Brassica oleracea</i> cultivars. <i>Journal of Experimental Botany</i> , 2010, 61, 807-819.	2.4	29
58	Microsatellite Markers in and around Rice Genes: Applications in Variety Identification and DUS Testing. <i>Crop Science</i> , 2009, 49, 880-886.	0.8	23
59	Genetic mapping and annotation of genomic microsatellites isolated from globe artichoke. <i>Theoretical and Applied Genetics</i> , 2009, 118, 1573-1587.	1.8	38
60	Darwin's wind hypothesis: does it work for plant dispersal in fragmented habitats?. <i>New Phytologist</i> , 2009, 183, 667-677.	3.5	59
61	Potato Cultivar Genome Analysis. <i>Methods in Molecular Biology</i> , 2009, 508, 295-308.	0.4	14
62	FISH mapping and molecular organization of the major repetitive sequences of tomato. <i>Chromosome Research</i> , 2008, 16, 919-933.	1.0	69
63	Natural hybridisation between <i>Populus nigra</i> L. and <i>P. x canadensis</i> Moench. Hybrid offspring competes for niches along the Rhine river in the Netherlands. <i>Tree Genetics and Genomes</i> , 2008, 4, 663-675.	0.6	62
64	Allele mining in <i>Solanum</i> : conserved homologues of <i>Rpi-blb1</i> are identified in <i>Solanum stoloniferum</i> . <i>Theoretical and Applied Genetics</i> , 2008, 116, 933-943.	1.8	117
65	The utility of NBS profiling for plant systematics: a first study in tuber-bearing <i>Solanum</i> species. <i>Plant Systematics and Evolution</i> , 2008, 276, 137-148.	0.3	12
66	HaploSNPer: a web-based allele and SNP detection tool. <i>BMC Genetics</i> , 2008, 9, 23.	2.7	37
67	Responses of <i>Brassica oleracea</i> cultivars to infestation by the aphid <i>Brevicoryne brassicae</i> : an ecological and molecular approach. <i>Plant, Cell and Environment</i> , 2008, 31, 1592-1605.	2.8	63
68	Large-scale identification of polymorphic microsatellites using an in silico approach. <i>BMC Bioinformatics</i> , 2008, 9, 374.	1.2	65
69	AFLP analysis reveals a lack of phylogenetic structure within <i>Solanum</i> section <i>Petota</i> . <i>BMC Evolutionary Biology</i> , 2008, 8, 145.	3.2	52
70	Effector Genomics Accelerates Discovery and Functional Profiling of Potato Disease Resistance and <i>Phytophthora Infestans</i> Avirulence Genes. <i>PLoS ONE</i> , 2008, 3, e2875.	1.1	361
71	AFLP markers as a tool to reconstruct complex relationships: A case study in <i>Rosa</i> ( <i>Rosaceae</i> ). <i>American Journal of Botany</i> , 2008, 95, 353-366.	0.8	143
72	Structure of the genetic diversity in black poplar ( <i>Populus nigra</i> L.) populations across European river systems: Consequences for conservation and restoration. <i>Forest Ecology and Management</i> , 2008, 255, 1388-1399.	1.4	116

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73	Reconstruction of fig wasp mating structure: how many mothers share a fig?. <i>Ecological Entomology</i> , 2007, 32, 485-491.	1.1	14
74	A Mixed-Model Approach to Association Mapping Using Pedigree Information With an Illustration of Resistance to <i>Phytophthora infestans</i> in Potato. <i>Genetics</i> , 2007, 175, 879-889.	1.2	205
75	Assignment Tests for Variety Identification Compared to Genetic Similarity-Based Methods Using Experimental Datasets from Different Marker Systems in Sugar Beet. <i>Crop Science</i> , 2007, 47, 1964-1974.	0.8	14
76	Infection of potato plants with potato leafroll virus changes attraction and feeding behaviour of <i>Myzus persicae</i> . <i>Entomologia Experimentalis Et Applicata</i> , 2007, 125, 135-144.	0.7	97
77	Genotypic variation in genome-wide transcription profiles induced by insect feeding: <i>Brassica oleracea</i> vs <i>Pieris rapae</i> interactions. <i>BMC Genomics</i> , 2007, 8, 239.	1.2	75
78	QTL identification for early blight resistance ( <i>Alternaria solani</i> ) in a <i>Solanum lycopersicum</i> x <i>S. arcanum</i> cross. <i>Theoretical and Applied Genetics</i> , 2007, 114, 439-450.	1.8	42
79	The distribution of genetic diversity in a <i>Brassica oleracea</i> gene bank collection related to the effects on diversity of regeneration, as measured with AFLPs. <i>Theoretical and Applied Genetics</i> , 2007, 114, 777-786.	1.8	46
80	Genetic population differentiation and connectivity among fragmented Moor frog ( <i>Rana arvalis</i> ) populations in The Netherlands. <i>Landscape Ecology</i> , 2007, 22, 1489-1500.	1.9	84
81	Plant translational genomics: from model species to crops. <i>Molecular Breeding</i> , 2007, 20, 1-13.	1.0	39
82	Unique genomic configuration revealed by microsatellite DNA in polyploid dogroses, <i>Rosa</i> sect. <i>Caninae</i> . <i>Journal of Evolutionary Biology</i> , 2006, 19, 635-648.	0.8	59
83	Location of resistance factors in the leaves of potato and wild tuber-bearing <i>Solanum</i> species to the aphid <i>Myzus persicae</i> . <i>Entomologia Experimentalis Et Applicata</i> , 2006, 121, 145-157.	0.7	171
84	Linked vs. unlinked markers: multilocus microsatellite haplotype-sharing as a tool to estimate gene flow and introgression. <i>Molecular Ecology</i> , 2006, 16, 243-256.	2.0	40
85	QualitySNP: a pipeline for detecting single nucleotide polymorphisms and insertions/deletions in EST data from diploid and polyploid species. <i>BMC Bioinformatics</i> , 2006, 7, 438.	1.2	127
86	Alpha-gliadin genes from the A, B, and D genomes of wheat contain different sets of celiac disease epitopes. <i>BMC Genomics</i> , 2006, 7, 1.	1.2	445
87	Genetic Structure in Populations of an Ancient Woodland Sedge, <i>Carex sylvatica</i> Hudson, at a Regional and Local Scale. <i>Plant Biology</i> , 2005, 7, 387-396.	1.8	15
88	Development and characterization of microsatellite markers for two dioecious <i>Ficus</i> species. <i>Molecular Ecology Notes</i> , 2005, 5, 355-357.	1.7	34
89	Development and mapping of a public reference set of SSR markers in <i>Lolium perenne</i> L.. <i>Molecular Ecology Notes</i> , 2005, 5, 951-957.	1.7	52
90	Pollinating fig wasps: genetic consequences of island recolonization. <i>Journal of Evolutionary Biology</i> , 2005, 18, 1234-1243.	0.8	40

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91	Microsatellite analysis of <i>Rosa damascena</i> Mill. accessions reveals genetic similarity between genotypes used for rose oil production and old Damask rose varieties. <i>Theoretical and Applied Genetics</i> , 2005, 111, 804-809.	1.8	91
92	Past and current gene flow in the selfing, wind-dispersed species <i>Mycelis muralis</i> in western Europe. <i>Molecular Ecology</i> , 2004, 13, 1391-1407.	2.0	26
93	Microsatellite DNA marker inheritance indicates preferential pairing between two highly homologous genomes in polyploid and hemisexual dog-roses, <i>Rosa</i> L. Sect. <i>Caninae</i> DC.. <i>Heredity</i> , 2004, 92, 139-150.	1.2	85
94	Genetic differentiation and trade among populations of peach palm ( <i>Bactris gasipaes</i> Kunth) in the Peruvian Amazon—implications for genetic resource management. <i>Theoretical and Applied Genetics</i> , 2004, 108, 1564-1573.	1.8	46
95	Efficient targeting of plant disease resistance loci using NBS profiling. <i>Theoretical and Applied Genetics</i> , 2004, 109, 384-393.	1.8	129
96	Assignment of allelic configuration in polyploids using the MAC-PR (microsatellite DNA allele) Tj ETQqO 0 0 rgBT /Overlock 10 Tf 50 542	1.8	208
97	The establishment of "essential derivation"™ among rose varieties, using AFLP. <i>Theoretical and Applied Genetics</i> , 2004, 109, 1718-1725.	1.8	37
98	Ex-situ conservation of Black poplar in Europe: genetic diversity in nine gene bank collections and their value for nature development. <i>Theoretical and Applied Genetics</i> , 2004, 108, 969-981.	1.8	65
99	Title is missing!. <i>Conservation Genetics</i> , 2003, 4, 441-451.	0.8	54
100	Assessment of the uniformity of wheat and tomato varieties at DNA microsatellite loci. <i>Euphytica</i> , 2003, 132, 331-341.	0.6	43
101	Identification of cut rose ( <i>Rosa hybrida</i> ) and rootstock varieties using robust sequence tagged microsatellite site markers. <i>Theoretical and Applied Genetics</i> , 2003, 106, 277-286.	1.8	133
102	Microsatellite genotyping of carnation varieties. <i>Theoretical and Applied Genetics</i> , 2003, 106, 1191-1195.	1.8	26
103	Characterization of microsatellite markers in <i>Fagus sylvatica</i> L. and <i>Fagus orientalis</i> Lipsky. <i>Molecular Ecology Notes</i> , 2003, 3, 76-78.	1.7	81
104	Cloning and characterization of four apple MADS box genes isolated from vegetative tissue. <i>Journal of Experimental Botany</i> , 2002, 53, 1025-1036.	2.4	61
105	Biodiversity assessment using markers for ecologically important traits. <i>Trends in Ecology and Evolution</i> , 2002, 17, 577-582.	4.2	149
106	Construction and testing of a microsatellite database containing more than 500 tomato varieties. <i>Theoretical and Applied Genetics</i> , 2002, 105, 1019-1026.	1.8	112
107	Construction and analysis of a microsatellite-based database of European wheat varieties. <i>Theoretical and Applied Genetics</i> , 2002, 106, 67-73.	1.8	134
108	Use of microsatellites to evaluate genetic diversity and species relationships in the genus <i>Lycopersicon</i> . <i>Theoretical and Applied Genetics</i> , 2001, 103, 1283-1292.	1.8	96

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109	Trinucleotide repeat microsatellite markers for black poplar ( <i>Populus nigra</i> L.). <i>Molecular Ecology Notes</i> , 2001, 1, 188-190.	1.7	137
110	Microsatellite markers for the European tree frog <i>Hyla arborea</i> . <i>Molecular Ecology</i> , 2000, 9, 1944-1946.	2.0	45
111	Heterogeneity of the internal transcribed spacer 1 (ITS1) in <i>Tulipa</i> (Liliaceae). <i>Plant Systematics and Evolution</i> , 2000, 225, 29-41.	0.3	16
112	Genetic Diversity and the Survival of Populations. <i>Plant Biology</i> , 2000, 2, 379-395.	1.8	335
113	Genetic Diversity and the Reintroduction of Meadow Species. <i>Plant Biology</i> , 2000, 2, 447-454.	1.8	34
114	Development and characterization of microsatellite markers in black poplar ( <i>Populus nigra</i> L.). <i>Theoretical and Applied Genetics</i> , 2000, 101, 317-322.	1.8	170
115	Microsatellite markers useful throughout the genus <i>Dianthus</i> . <i>Genome</i> , 2000, 43, 208-210.	0.9	13
116	Microsatellite retrieval in lettuce ( <i>Lactuca sativa</i> L.). <i>Genome</i> , 1999, 42, 139-149.	0.9	92
117	Microsatellite retrieval in lettuce ( <i>Lactuca sativa</i> L.). <i>Genome</i> , 1999, 42, 139-149.	0.9	14
118	The use of semi-automated fluorescent microsatellite analysis for tomato cultivar identification. <i>Theoretical and Applied Genetics</i> , 1998, 97, 584-590.	1.8	70
119	Molecular genetic analysis of black poplar ( <i>Populus nigra</i> L.) along Dutch rivers. <i>Molecular Ecology</i> , 1998, 7, 11-18.	2.0	160
120	Phylogenetic relationships among <i>Lactuca</i> (Asteraceae) species and related genera based on ITS-1 DNA sequences. <i>American Journal of Botany</i> , 1998, 85, 1517-1530.	0.8	80
121	Molecular characterization of GATA/GACA microsatellite repeats in tomato. <i>Genome</i> , 1997, 40, 25-33.	0.9	43
122	Molecular technologies for biodiversity evaluation: Opportunities and challenges. <i>Nature Biotechnology</i> , 1997, 15, 625-628.	9.4	147
123	Title is missing!. <i>Molecular Breeding</i> , 1997, 3, 381-390.	1.0	619
124	Use of short microsatellites from database sequences to generate polymorphisms among <i>Lycopersicon esculentum</i> cultivars and accessions of other <i>Lycopersicon</i> species. <i>Theoretical and Applied Genetics</i> , 1997, 94, 264-272.	1.8	251
125	Characterization of the extracellular lipase, LipA, of <i>Acinetobacter calcoaceticus</i> BD413 and sequence analysis of the cloned structural gene. <i>Molecular Microbiology</i> , 1995, 15, 803-818.	1.2	72
126	A gene of <i>Acinetobacter calcoaceticus</i> BD413 encodes a periplasmic peptidyl-prolyl cis-trans isomerase of the cyclophilin sub-class that is not essential for growth. <i>Biochimica Et Biophysica Acta Gene Regulatory Mechanisms</i> , 1994, 1219, 601-606.	2.4	15



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127	Direct comparison of levels of genetic variation in tomato detected by a GACA-containing microsatellite probe and by random amplified polymorphic DNA. <i>Genome</i> , 1994, 37, 375-381.	0.9	64
128	Regulation of the expression of the <i>Pseudomonas stutzeri</i> recA gene. <i>Antonie Van Leeuwenhoek</i> , 1993, 63, 55-62.	0.7	2
129	Increased stability of recombinant plasmids by Tn1000 insertion in chemostat cultures of recombinant <i>Escherichia coli</i> GT123. <i>Current Microbiology</i> , 1993, 26, 281-286.	1.0	2
130	Characterization of transformation-deficient mutants of <i>Acinetobacter calcoaceticus</i> . <i>Molecular Microbiology</i> , 1992, 6, 1747-1754.	1.2	33
131	Molecular cloning and functional characterization of a recA analog from <i>Pseudomonas stutzeri</i> and construction of a <i>P. stutzeri</i> recA mutant. <i>Antonie Van Leeuwenhoek</i> , 1991, 59, 115-123.	0.7	21
132	Cloning in <i>Escherichia coli</i> of the gene specifying the DNA-entry nuclease of <i>Bacillus subtilis</i> . <i>Gene</i> , 1987, 52, 175-183.	1.0	31
133	Integration of vector-containing <i>Bacillus subtilis</i> chromosomal DNA by a Campbell-like mechanism. <i>Molecular Genetics and Genomics</i> , 1986, 204, 524-531.	2.4	25