

Magda-Viola Hanke

List of Publications by Year in descending order

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Version: 2024-02-01

113
papers

2,930
citations

147801

31
h-index

206112

48
g-index

120
all docs

120
docs citations

120
times ranked

2257
citing authors

#	ARTICLE	IF	CITATIONS
1	Insights into the susceptibility of raspberries to <i>Drosophila suzukii</i> oviposition. <i>Journal of Applied Entomology</i> , 2021, 145, 182-190.	1.8	7
2	Toward Systematic Understanding of Flower Bud Induction in Apple: A Multi-Omics Approach. <i>Frontiers in Plant Science</i> , 2021, 12, 604810.	3.6	12
3	Transcriptional profile of AvrRpt2EA-mediated resistance and susceptibility response to <i>Erwinia amylovora</i> in apple. <i>Scientific Reports</i> , 2021, 11, 8685.	3.3	4
4	No Evidence of Unexpected Transgenic Insertions in T1190 “A Transgenic Apple Used in Rapid Cycle Breeding” Following Whole Genome Sequencing. <i>Frontiers in Plant Science</i> , 2021, 12, 715737.	3.6	2
5	Self-incompatibility of raspberry cultivars assessed by SSR markers. <i>Scientia Horticulturae</i> , 2021, 288, 110384.	3.6	2
6	The MADS-Box Gene MdDAM1 Controls Growth Cessation and Bud Dormancy in Apple. <i>Frontiers in Plant Science</i> , 2020, 11, 1003.	3.6	38
7	Introgressing blue mold resistance into elite apple germplasm by rapid cycle breeding and foreground and background DNA-informed selection. <i>Tree Genetics and Genomes</i> , 2020, 16, 1.	1.6	16
8	SSR fingerprinting of raspberry cultivars traded in Germany clearly showed that certainty about the genotype authenticity is a prerequisite for any horticultural experiment. <i>European Journal of Horticultural Science</i> , 2020, 85, 79-85.	0.7	8
9	Physiological, biochemical and genetic responses of Caucasian tea (<i>Camellia sinensis</i> (L.) Kuntze) genotypes under cold and frost stress. <i>PeerJ</i> , 2020, 8, e9787.	2.0	18
10	A comparison of genetic stability in tea [<i>Camellia sinensis</i> (L.) Kuntze] plantlets derived from callus with plantlets from long-term in vitro propagation. <i>Plant Cell, Tissue and Organ Culture</i> , 2019, 138, 467-474.	2.3	17
11	Evaluation of <i>Malus</i> genetic resources for tolerance to apple replant disease (ARD). <i>Scientia Horticulturae</i> , 2019, 256, 108517.	3.6	38
12	High crop load and low temperature delay the onset of bud initiation in apple. <i>Scientific Reports</i> , 2019, 9, 17986.	3.3	13
13	Mapping of fire blight resistance in <i>Malus robusta</i> 5 flowers following artificial inoculation. <i>BMC Plant Biology</i> , 2019, 19, 532.	3.6	24
14	Evaluation of <i>Malus</i> gene bank resources with German strains of <i>Marssonina coronaria</i> using a greenhouse-based screening method. <i>European Journal of Plant Pathology</i> , 2019, 153, 743-757.	1.7	7
15	Genes Involved in Stress Response and Especially in Phytoalexin Biosynthesis Are Upregulated in Four <i>Malus</i> Genotypes in Response to Apple Replant Disease. <i>Frontiers in Plant Science</i> , 2019, 10, 1724.	3.6	27
16	Inoculation of <i>Malus</i> genotypes with a set of <i>Erwinia amylovora</i> strains indicates a gene-for-gene relationship between the effector gene <i>eop1</i> and both <i>Malus floribunda</i> 821 and <i>Malus</i> “Evereste”™. <i>Plant Pathology</i> , 2018, 67, 938-947.	2.4	22
17	Generation of advanced fire blight-resistant apple (<i>Malus domestica</i>) selections of the fifth generation within 7 years of applying the early flowering approach. <i>Planta</i> , 2018, 247, 1475-1488.	3.2	38
18	Apple cultivar Regia possessing both <i>Rvi2</i> and <i>Rvi4</i> resistance genes is the source of a new race of <i>Venturia inaequalis</i> . <i>European Journal of Plant Pathology</i> , 2018, 151, 533-539.	1.7	8

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19	Towards map-based cloning of FB_Mfu10: identification of a receptor-like kinase candidate gene underlying the <i>Malus fusca</i> fire blight resistance locus on linkage group 10. <i>Molecular Breeding</i> , 2018, 38, 106.	2.1	28
20	A Single Effector Protein, AvrRpt2_{EA}, from <i>Erwinia amylovora</i> Can Cause Fire Blight Disease Symptoms and Induces a Salicylic Acid-Dependent Defense Response. <i>Molecular Plant-Microbe Interactions</i> , 2018, 31, 1179-1191.	2.6	19
21	Evaluation of <i>Rubus</i> genetic resources on their resistance to cane disease. <i>Genetic Resources and Crop Evolution</i> , 2018, 65, 1979-1993.	1.6	4
22	SSR fingerprinting of a German <i>Rubus</i> collection and pedigree based evaluation on trueness-to-type. <i>Genetic Resources and Crop Evolution</i> , 2017, 64, 189-203.	1.6	18
23	To what extent do wild apples in Kazakhstan retain their genetic integrity?. <i>Tree Genetics and Genomes</i> , 2017, 13, 1.	1.6	26
24	Spatial and Temporal Localization of Flavonoid Metabolites in Strawberry Fruit (<i>Fragaria</i> Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	8.2	41
25	Cryopreservation of fruit germplasm. <i>In Vitro Cellular and Developmental Biology - Plant</i> , 2017, 53, 372-381.	2.1	21
26	Fire blight resistance of <i>Malus arnoldiana</i> is controlled by a quantitative trait locus located at the distal end of linkage group 12. <i>European Journal of Plant Pathology</i> , 2017, 148, 1011-1018.	1.7	32
27	Evaluation of a MdMYB10/GFP43 fusion gene for its suitability to act as reporter gene in promoter studies in <i>Fragaria vesca</i> L. â€”RÄ¼genâ€™. <i>Plant Cell, Tissue and Organ Culture</i> , 2017, 130, 345-356.	2.3	4
28	Himbeere und Brombeere (<i>Rubus</i> spp.)., 2017, , 353-384.		1
29	<sc>TERMINAL FLOWER</sc> 1</i> is a breeding target for a novel everbearing trait and tailored flowering responses in cultivated strawberry (<i>Fragaria</i> — <i>ananassa</i> Duch.). <i>Plant Biotechnology Journal</i> , 2016, 14, 1852-1861.	8.3	52
30	Molecular and flow cytometric evaluation of pear (<i>Pyrus</i> L.) genetic resources of the German and Romanian national fruit collections. <i>Genetic Resources and Crop Evolution</i> , 2016, 63, 1023-1033.	1.6	19
31	Efficient heat-shock removal of the selectable marker gene in genetically modified grapevine. <i>Plant Cell, Tissue and Organ Culture</i> , 2016, 124, 471-481.	2.3	37
32	Transgenic apple plants overexpressing the chalcone 3-hydroxylase gene of <i>Cosmos sulphureus</i> show increased levels of 3-hydroxyphloridzin and reduced susceptibility to apple scab and fire blight. <i>Planta</i> , 2016, 243, 1213-1224.	3.2	35
33	Homologs of the FB_MR5 fire blight resistance gene of <i>Malus robusta</i> 5 are present in other <i>Malus</i> wild species accessions. <i>Tree Genetics and Genomes</i> , 2016, 12, 1.	1.6	5
34	Resistance and systemic dispersal of <i>Xanthomonas fragariae</i> in strawberry germplasm (<i>Fragaria</i> L.). <i>Plant Pathology</i> , 2015, 64, 71-80.	2.4	20
35	Integration of <i>BpMADS</i> 4 on various linkage groups improves the utilization of the rapid cycle breeding system in apple. <i>Plant Biotechnology Journal</i> , 2015, 13, 246-258.	8.3	20
36	The fire blight resistance QTL of <i>Malus fusca</i> (Mfu10) is affected but not broken down by the highly virulent Canadian <i>Erwinia amylovora</i> strain E2002A. <i>European Journal of Plant Pathology</i> , 2015, 141, 631-635.	1.7	16

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37	Improving resistance of different apple cultivars using the Rvi6 scab resistance gene in a cisgenic approach based on the Flp/FRT recombinase system. <i>Molecular Breeding</i> , 2015, 35, 1.	2.1	44
38	Heat mediated silencing of MdTFL1 genes in apple (<i>Malus domestica</i>). <i>Plant Cell, Tissue and Organ Culture</i> , 2015, 123, 511-521.	2.3	10
39	Evaluation of strawberry (<i>Fragaria</i> L.) genetic resources for resistance to <i>Botrytis cinerea</i> . <i>Plant Pathology</i> , 2015, 64, 396-405.	2.4	23
40	Identification of a major quantitative trait locus for resistance to fire blight in the wild apple species <i>Malus fusca</i> . <i>Molecular Breeding</i> , 2014, 34, 407-419.	2.1	50
41	Engineering fire blight resistance into the apple cultivar 'Gala' using the <i>Fb_MR5</i> <i>CC-NBS-LRR</i> resistance gene of <i>Malus robusta</i> 5. <i>Plant Biotechnology Journal</i> , 2014, 12, 728-733.	8.3	70
42	Premature and ectopic anthocyanin formation by silencing of anthocyanidin reductase in strawberry (<i>Fragaria ananassa</i>). <i>New Phytologist</i> , 2014, 201, 440-451.	7.3	57
43	QTL mapping of fire blight resistance in <i>Malus robusta</i> 5 after inoculation with different strains of <i>Erwinia amylovora</i> . <i>Molecular Breeding</i> , 2014, 34, 217-230.	2.1	31
44	A diallel crossing approach aimed on selection for ripening time and yield in breeding of new strawberry (<i>Fragaria ananassa</i> Duch.) cultivars. <i>Plant Breeding</i> , 2014, 133, 115-120.	1.9	14
45	Phenotypic and genetic analysis of the German <i>Malus</i> Germplasm Collection in terms of type 1 and type 2 red-fleshed apples. <i>Gene</i> , 2014, 544, 198-207.	2.2	15
46	DEVELOPMENT OF APPLE PRE-BREEDING GENOTYPES HIGHLY RESISTANT TO FIRE BLIGHT BY EARLY FLOWERING. <i>Acta Horticulturae</i> , 2014, , 55-64.	0.2	1
47	COMPARATIVE MAPPING OF FIRE BLIGHT RESISTANCE IN MALUS. <i>Acta Horticulturae</i> , 2014, , 47-51.	0.2	6
48	BIPHENYLS AND DIBENZOFURANS - FIRE BLIGHT-INDUCED PHYTOALEXINS OF PEAR. <i>Acta Horticulturae</i> , 2014, , 181-185.	0.2	0
49	INVESTIGATION ON FIRE BLIGHT RESISTANCE IN THE CROSS POPULATION 'IDARED' <i>Malus robusta</i> 5 WITH DIFFERENT ERWINIA AMYLOVORA STRAINS. <i>Acta Horticulturae</i> , 2014, , 277-280.	0.2	1
50	EVIDENCE OF A MAJOR QTL FOR FIRE BLIGHT RESISTANCE IN THE APPLE WILD SPECIES MALUS FUSCA. <i>Acta Horticulturae</i> , 2014, , 289-293.	0.2	14
51	FB-MR5 IS AN APPLE GENE PROVIDING RESISTANCE TO FIRE BLIGHT. <i>Acta Horticulturae</i> , 2014, , 273-276.	0.2	1
52	FRUIT GENETIC RESOURCES MANAGEMENT: COLLECTION, CONSERVATION, EVALUATION AND UTILIZATION IN GERMANY. <i>Acta Horticulturae</i> , 2014, , 231-234.	0.2	4
53	THE "GERMAN NATIONAL FRUIT GENE BANK", A FIRST REVIEW FIVE YEARS AFTER LAUNCHING. <i>Acta Horticulturae</i> , 2014, , 227-230.	0.2	0
54	Phytoalexin formation in fire blight-infected apple. <i>Trees - Structure and Function</i> , 2013, 27, 477-484.	1.9	27

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55	The Fast-track breeding approach can be improved by heat-induced expression of the FLOWERING LOCUS T genes from poplar (<i>Populus trichocarpa</i>) in apple (<i>Malus × domestica</i> Borkh.). <i>Plant Cell, Tissue and Organ Culture</i> , 2013, 115, 127-137.	2.3	52
56	Assessment of phenotypic variation of <i>Malus orientalis</i> in the North Caucasus region. <i>Genetic Resources and Crop Evolution</i> , 2013, 60, 1463-1477.	1.6	23
57	Studies on heat shock induction and transgene expression in order to optimize the Flp/FRT recombinase system in apple (<i>Malus × domestica</i> Borkh.). <i>Plant Cell, Tissue and Organ Culture</i> , 2013, 115, 457-467.	2.3	13
58	Gene-for-gene relationship in the host-pathogen system <i>Malus × domestica</i> – <i>Erwinia amylovora</i> . <i>New Phytologist</i> , 2013, 197, 1262-1275.	7.3	88
59	BREEDING OF RESISTANT STRAWBERRY CULTIVARS FOR ORGANIC FRUIT PRODUCTION - PRELIMINARY RESULTS WITH <i>BOTRYTIS CINEREA</i> . <i>Acta Horticulturae</i> , 2013, , 87-90.	0.2	4
60	CISGENIC APPROACH FOR IMPROVED DISEASE RESISTANCE IN APPLE. <i>Acta Horticulturae</i> , 2013, , 117-121.	0.2	2
61	PRELIMINARY RESULTS TO ESTABLISH A SPEED-BREED PROGRAM BASED ON HEAT-INDUCED PRECOCIOUS FLOWERING OF APPLE PLANTS CONTAINING THE FLOWERING LOCUS T GENE FROM POPLAR (<i>POPULUS</i>) Tj ETQq10120.784314 rgBT		
62	QTL MAPPING FOR RESISTANCE TO FIRE BLIGHT USING SEVERAL <i>ERWINIA AMYLOVORA</i> STRAINS RESULTING IN DIFFERENT HOST-PATHOGEN INTERACTIONS. <i>Acta Horticulturae</i> , 2013, , 509-512.	0.2	0
63	RNA-Mediated Gene Silencing Signals Are Not Graft Transmissible from the Rootstock to the Scion in Greenhouse-Grown Apple Plants <i>Malus</i> sp.. <i>International Journal of Molecular Sciences</i> , 2012, 13, 9992-10009.	4.1	28
64	Differential Expression of Biphenyl Synthase Gene Family Members in Fire-Blight-Infected Apple ‘Holsteiner Cox’™. <i>Plant Physiology</i> , 2012, 158, 864-875.	4.8	42
65	BIOTECHNOLOGICAL APPROACHES TO SHORTEN THE JUVENILE PERIOD IN FRUIT TREES. <i>Acta Horticulturae</i> , 2012, , 309-314.	0.2	1
66	Functional Genomics of Flowering Time in Trees. , 2012, , 39-69.		5
67	The MdTFL1 gene of apple (<i>Malus × domestica</i> Borkh.) reduces vegetative growth and generation time. <i>Tree Physiology</i> , 2012, 32, 1288-1301.	3.1	91
68	The role of Schmidt ‘Antonovka’™ in apple scab resistance breeding. <i>Tree Genetics and Genomes</i> , 2012, 8, 627-642.	1.6	14
69	Use of a transgenic early flowering approach in apple (<i>Malus × domestica</i> Borkh.) to introgress fire blight resistance from cultivar Evereste. <i>Molecular Breeding</i> , 2012, 30, 857-874.	2.1	39
70	Heat-shock-mediated elimination of the nptII marker gene in transgenic apple (<i>Malus × domestica</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	2.2	32
71	Chitinase activities, scab resistance, mycorrhization rates and biomass of own-rooted and grafted transgenic apple. <i>Genetics and Molecular Biology</i> , 2012, 35, 466-473.	1.3	9
72	GENETIC CONTROL OF FLOWER DEVELOPMENT IN APPLE AND THE UTILISATION OF TRANSGENIC EARLY FLOWERING APPLE PLANTS IN BREEDING. <i>Acta Horticulturae</i> , 2012, , 29-34.	0.2	0

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73	Formation of biphenyl and dibenzofuran phytoalexins in the transition zones of fire blight-infected stems of <i>Malus domestica</i> cv. 'Holsteiner Cox'™ and <i>Pyrus communis</i> cv. 'Conference'™. <i>Phytochemistry</i> , 2012, 77, 179-185.		57
74	Silencing of flavanone-3-hydroxylase in apple (<i>Malus domestica</i> Borkh.) leads to accumulation of flavanones, but not to reduced fire blight susceptibility. <i>Plant Physiology and Biochemistry</i> , 2012, 51, 18-25.	5.8	32
75	Substrate specificity and contribution of the glycosyltransferase UGT71A15 to phloridzin biosynthesis. <i>Trees - Structure and Function</i> , 2012, 26, 259-271.	1.9	23
76	Molecular analysis of Iranian seedless barberries via SSR. <i>Scientia Horticulturae</i> , 2011, 129, 702-709.	3.6	14
77	BIPHENYL AND DIBENZOFURAN FORMATION IN FIRE BLIGHT-INFECTED <i>MALUS DOMESTICA</i> CULTIVARS. <i>Acta Horticulturae</i> , 2011, , 547-553.	0.2	0
78	INOCULATION OF <i>MALUS ROBUSTA</i> 5 PROGENY WITH A STRAIN BREAKING RESISTANCE TO FIRE BLIGHT REVEALS A MINOR QTL ON LG5. <i>Acta Horticulturae</i> , 2011, , 357-362.	0.2	31
79	Application of a high-speed breeding technology to apple (<i>Malus domestica</i>) based on transgenic early flowering plants and marker-assisted selection. <i>New Phytologist</i> , 2011, 192, 364-377.	7.3	141
80	Note added in proof to: Over-expression of an FT-homologous gene of apple induces early flowering in annual and perennial plants. <i>Planta</i> , 2011, 233, 217-218.	3.2	8
81	Transgenic Fruit Crops in Europe. , 2011, , 125-145.		2
82	Overexpression of LEAFY in apple leads to a columnar phenotype with shorter internodes. <i>Planta</i> , 2010, 231, 251-263.	3.2	50
83	Transgenic apple plants overexpressing the Lc gene of maize show an altered growth habit and increased resistance to apple scab and fire blight. <i>Planta</i> , 2010, 231, 623-635.	3.2	46
84	Over-expression of an FT-homologous gene of apple induces early flowering in annual and perennial plants. <i>Planta</i> , 2010, 232, 1309-1324.	3.2	144
85	Fruit Crops. <i>Biotechnology in Agriculture and Forestry</i> , 2010, , 307-348.	0.2	8
86	<i>Erwinia amylovora</i> -induced defense mechanisms of two apple species that differ in susceptibility to fire blight. <i>Plant Science</i> , 2010, 179, 60-67.	3.6	41
87	FIRST RESULTS ON THE EFFECT OF INCREASED CHITINASE EXPRESSION IN TRANSGENIC APPLE TREES ON MYCORRHIZATION WITH <i>GLOMUS INTRARADICES</i> AND <i>G. MOSSEAE</i> . <i>Acta Horticulturae</i> , 2009, , 719-724.	0.2	3
88	THE SWITCH TO FLOWERING: GENES INVOLVED IN FLORAL INDUCTION OF THE APPLE CULTIVAR 'PINOVA' AND THE ROLE OF THE FLOWERING GENE MdFT. <i>Acta Horticulturae</i> , 2009, , 701-705.	0.2	5
89	METABOLIC ENGINEERING OF FLAVONOID BIOSYNTHESIS IN APPLE (<i>MALUS DOMESTICA</i> BORKH.). <i>Acta Horticulturae</i> , 2009, , 511-516.	0.2	3
90	Shift in polyphenol profile and sublethal phenotype caused by silencing of anthocyanidin synthase in apple (<i>Malus</i> sp.). <i>Planta</i> , 2009, 229, 681-692.	3.2	61

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91	A review on transgenic approaches to accelerate breeding of woody plants. <i>Plant Breeding</i> , 2009, 128, 217-226.	1.9	130
92	RNAi-SILENCING OF MdTFL1 INDUCES EARLY FLOWERING IN APPLE. <i>Acta Horticulturae</i> , 2009, , 633-636.	0.2	9
93	STUDIES ON MRNA EXPRESSION OF GENES INVOLVED IN FLORAL MERISTEM TRANSITION OF APPLE (MALUS) Tj ET Oq 1 1 0.784314 r	0.2	0
94	IDENTIFICATION OF CULTIVABLE BACTERIA FROM IN VITRO CULTURES OF APPLE. <i>Acta Horticulturae</i> , 2009, , 733-738.	0.2	3
95	RATIO OF HOMOZYGOUS AND HETEROZYGOUS VF GENOTYPES IN THE PROGENIES OF APPLE VFVF X VFVF CROSSES. <i>Acta Horticulturae</i> , 2009, , 819-824.	0.2	3
96	PRELIMINARY RESULTS TO ESTABLISH THE DAAO SYSTEM AS AN ALTERNATIVE SELECTION STRATEGY ON APPLE. <i>Acta Horticulturae</i> , 2009, , 267-272.	0.2	4
97	VERIFYING THE PARENTS OF THE PILLNITZER APPLE CULTIVARS. <i>Acta Horticulturae</i> , 2009, , 319-324.	0.2	6
98	SYSTEMIC ACQUIRED SILENCING OF A GUSA TRANSGENE IN APPLE. <i>Acta Horticulturae</i> , 2009, , 393-396.	0.2	1
99	Transgenic expression of a viral EPSâ€¢depolymerase is potentially useful to induce fire blight resistance in apple. <i>Annals of Applied Biology</i> , 2008, 153, 345-355.	2.5	23
100	Isolation of flowering genes and seasonal changes in their transcript levels related to flower induction and initiation in apple (<i>Malus domestica</i>). <i>Tree Physiology</i> , 2008, 28, 1459-1466.	3.1	60
101	Evaluation of the uniformity and stability of T-DNA integration and gene expression in transgenic apple plants. <i>Electronic Journal of Biotechnology</i> , 2008, 11, 0-0.	2.2	28
102	IMPROVED FIRE BLIGHT RESISTANCE IN TRANSGENIC APPLE LINES BY CONSTITUTIVE OVEREXPRESSION OF THE mbr4 GENE OF MALUS BACCATA. <i>Acta Horticulturae</i> , 2008, , 287-291.	0.2	5
103	CONFIRMATION OF THE FIRE BLIGHT QTL OF MALUS Ã— ROBUSTA 5 ON LINKAGE GROUP 3. <i>Acta Horticulturae</i> , 2008, , 297-303.	0.2	36
104	Overexpression of BpMADS4 from silver birch (<i>Betula pendula</i> Roth.) induces early-flowering in apple (<i>Malus</i> $\frac{1}{2}$ \times <i>domestica</i> Borkh.). <i>Plant Breeding</i> , 2007, 126, 137-145.	1.9	150
105	Strong evidence for a fire blight resistance gene of <i>Malus robusta</i> located on linkage group 3. <i>Plant Breeding</i> , 2007, 126, 470-475.	1.9	124
106	Maize Lc transcription factor enhances biosynthesis of anthocyanins, distinct proanthocyanidins and phenylpropanoids in apple (<i>Malus domestica</i> Borkh.). <i>Planta</i> , 2007, 226, 1243-1254.	3.2	92
107	ANALYSIS OF TISSUE UNIFORMITY IN TRANSGENIC APPLE PLANTS. <i>Acta Horticulturae</i> , 2007, , 301-306.	0.2	4
108	BPMADS4 - A MADS BOX GENE OF BIRCH INDUCES FLOWERS ON TRANSGENIC APPLE PLANTS IN VITRO. <i>Acta Horticulturae</i> , 2007, , 307-312.	0.2	2

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109	DEVELOPING MOLECULAR MARKERS FOR MARKER ASSISTED SELECTION OF FIRE BLIGHT RESISTANT APPLE SEEDLINGS. Acta Horticulturae, 2007, , 117-122.	0.2	3
110	TRANSCRIPTION PROFILING ON TRANSGENIC APPLE PLANTS AFTER OVER-EXPRESSION OF GENES, WHICH ARE INVOLVED IN THE FLOWER DEVELOPMENT. Acta Horticulturae, 2007, , 215-222.	0.2	1
111	Assessing gene flow in apple using a descendant of <i>Malus sieversii</i> var. <i>sieversii</i> f. <i>niedzwetzkyana</i> as an identifier for pollen dispersal. Environmental Biosafety Research, 2006, 5, 89-104.	1.1	14
112	An Efficient Method for Rooting and Acclimation of Micropropagated Apple Cultivars. Hortscience: A Publication of the American Society for Horticultural Science, 1998, 33, 1251-1252.	1.0	42
113	Klimawandel – Ausrichtung der Züchtung bei Obst. , 0, , .		0