

Thomas Van Leeuwen

List of Publications by Year in descending order

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156
papers

10,987
citations

36691

53
h-index

42259

96
g-index

166
all docs

166
docs citations

166
times ranked

7033
citing authors

#	ARTICLE	IF	CITATIONS
1	Multiple TaqMan qPCR and droplet digital PCR (ddPCR) diagnostics for pesticide resistance monitoring and management, in the major agricultural pest <i>Tetranychus urticae</i> . Pest Management Science, 2022, 78, 263-273.	1.7	15
2	Biochemical and molecular mechanisms of acaricide resistance in <i>Dermanyssus gallinae</i> populations from Turkey. Pesticide Biochemistry and Physiology, 2022, 180, 104985.	1.6	8
3	Combination of target site mutation and associated CYPs confers high-level resistance to pyridaben in <i>Tetranychus urticae</i> . Pesticide Biochemistry and Physiology, 2022, 181, 105000.	1.6	12
4	Selectivity and molecular stress responses to classical and botanical acaricides in the predatory mite <i>Phytoseiulus persimilis</i> (Acari: Phytoseiidae) (<i>Athias</i> Henriot) (Acari: Phytoseiidae). Pest Management Science, 2022, 78, 881-895.	1.7	13
5	Variation of diazinon and amitraz susceptibility of <i>Hyalomma marginatum</i> (Acari: Ixodidae) in the Rabat-Sale-Kenitra region of Morocco. Ticks and Tick-borne Diseases, 2022, 13, 101883.	1.1	2
6	Pirimicarb resistance and associated mechanisms in field-collected and selected populations of <i>Neoseiulus californicus</i> . Pesticide Biochemistry and Physiology, 2022, 180, 104984.	1.6	7
7	Over-expression in cis of the midgut P450 CYP392A16 contributes to abamectin resistance in <i>Tetranychus urticae</i> . Insect Biochemistry and Molecular Biology, 2022, 142, 103709.	1.2	13
8	Structural and functional characterization of \hat{I}^2 -cyanoalanine synthase from <i>Tetranychus urticae</i> . Insect Biochemistry and Molecular Biology, 2022, 142, 103722.	1.2	2
9	Cover Image, Volume 78, Issue 3. Pest Management Science, 2022, 78, .	1.7	0
10	QTL mapping suggests that both cytochrome P450-mediated detoxification and target-site resistance are involved in fenbutatin oxide resistance in <i>Tetranychus urticae</i> . Insect Biochemistry and Molecular Biology, 2022, 145, 103757.	1.2	13
11	A H258Y mutation in subunit B of the succinate dehydrogenase complex of the spider mite <i>Tetranychus urticae</i> confers resistance to cyenopyrafen and pyflubumide, but likely reinforces cyflumetofen binding and toxicity. Insect Biochemistry and Molecular Biology, 2022, 144, 103761.	1.2	7
12	Interactions With Plant Defences Isolate Sympatric Populations of an Herbivorous Mite. Frontiers in Ecology and Evolution, 2022, 10, .	1.1	3
13	Biochemical and insecticidal effects of plant essential oils on insecticide resistant and susceptible populations of <i>Musca domestica</i> L. point to a potential cross-resistance risk. Pesticide Biochemistry and Physiology, 2022, 184, 105115.	1.6	8
14	The H92R substitution in PSST is a reliable diagnostic biomarker for predicting resistance to mitochondrial electron transport inhibitors of complex I in European populations of <i>Tetranychus urticae</i> . Pest Management Science, 2022, 78, 3644-3653.	1.7	10
15	A mutation in chitin synthase I associated with etoxazole resistance in the citrus red mite <i>Panonychus citri</i> (Acari: Tetranychidae) and its uneven geographical distribution in Japan. Pest Management Science, 2022, 78, 4028-4036.	1.7	3
16	Intradiol ring cleavage dioxygenases from herbivorous spider mites as a new detoxification enzyme family in animals. BMC Biology, 2022, 20, .	1.7	14
17	Incidence of spiromesifen resistance and resistance mechanisms in <i>Tetranychus urticae</i> populations collected from strawberry production areas in Turkey. Crop Protection, 2022, 160, 106049.	1.0	7
18	Fenpyroximate resistance in Iranian populations of the European red mite <i>Panonychus ulmi</i> (Acari: Tj ETQq0 0 0 rgBT/Overlock 10 Tf 50	0.7	7

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19	Identification and characterization of striking multiple insecticide resistance in a <i>Tetranychus urticae</i> field population from Greece. <i>Pest Management Science</i> , 2021, 77, 666-676.	1.7	23
20	Is the emerging mite pest <i>Aculops lycopersici</i> controllable? Global and genome-based insights in its biology and management. <i>Pest Management Science</i> , 2021, 77, 2635-2644.	1.7	21
21	Reduced proinsecticide activation by cytochrome P450 confers coumaphos resistance in the major bee parasite <i>Varroa destructor</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	42
22	Short term transcriptional responses of P450s to phytochemicals in insects and mites. <i>Current Opinion in Insect Science</i> , 2021, 43, 117-127.	2.2	35
23	Genetic analysis and screening of pyrethroid resistance mutations in <i>Varroa destructor</i> populations from Turkey. <i>Experimental and Applied Acarology</i> , 2021, 84, 433-444.	0.7	9
24	High-resolution genetic mapping reveals cis-regulatory and copy number variation in loci associated with cytochrome P450-mediated detoxification in a generalist arthropod pest. <i>PLoS Genetics</i> , 2021, 17, e1009422.	1.5	26
25	The role of detoxification enzymes in the susceptibility of <i>Brevipalpus californicus</i> exposed to acaricide and insecticide mixtures. <i>Pesticide Biochemistry and Physiology</i> , 2021, 175, 104855.	1.6	7
26	The phenylpropanoid pathway inhibitor piperonylic acid induces broad-spectrum pest and disease resistance in plants. <i>Plant, Cell and Environment</i> , 2021, 44, 3122-3139.	2.8	31
27	Adaptive divergence and post-zygotic barriers to gene flow between sympatric populations of a herbivorous mite. <i>Communications Biology</i> , 2021, 4, 853.	2.0	12
28	The genome of the extremophile <i>Artemia</i> provides insight into strategies to cope with extreme environments. <i>BMC Genomics</i> , 2021, 22, 635.	1.2	20
29	Editorial: Invertebrate UDP-Glycosyltransferases: Nomenclature, Diversity and Functions. <i>Frontiers in Physiology</i> , 2021, 12, 748290.	1.3	3
30	Comparing the efficiency of RNAi after feeding and injection of dsRNA in spider mites. <i>Pesticide Biochemistry and Physiology</i> , 2021, 179, 104966.	1.6	9
31	Untangling a Gordian knot: the role of a <i>GluCl3 I321T</i> mutation in abamectin resistance in <i>Tetranychus urticae</i> . <i>Pest Management Science</i> , 2021, 77, 1581-1593.	1.7	29
32	The G126S substitution in mitochondrially encoded cytochrome b does not confer bifenthrin resistance in the spider mite <i>Tetranychus urticae</i> . <i>Experimental and Applied Acarology</i> , 2021, 85, 161-172.	0.7	7
33	Pyrethroid target-site resistance mutations in populations of the honey bee parasite <i>Varroa destructor</i> (Acari: Varroidae) from Flanders, Belgium. <i>Experimental and Applied Acarology</i> , 2021, 85, 205-221.	0.7	8
34	Ticks and Tick-Borne Pathogens Abound in the Cattle Population of the Rabat-Sale Kenitra Region, Morocco. <i>Pathogens</i> , 2021, 10, 1594.	1.2	7
35	Overexpression of an alternative allele of carboxyl/choline esterase 4 (CCE04) of <i>Tetranychus urticae</i> is associated with high levels of resistance to the keto-enol acaricide spiroticlofen. <i>Pest Management Science</i> , 2020, 76, 1142-1153.	1.7	29
36	Identification and geographical distribution of pyrethroid resistance mutations in the poultry red mite <i>Dermanyssus gallinae</i> . <i>Pest Management Science</i> , 2020, 76, 125-133.	1.7	33

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37	Identification and characterization of new mutations in mitochondrial cytochrome b that confer resistance to bifenazate and acequinocyl in the spider mite <i>Tetranychus urticae</i> . <i>Pest Management Science</i> , 2020, 76, 1154-1163.	1.7	39
38	Molecular and genetic analysis of resistance to MET-I acaricides in Iranian populations of the citrus red mite <i>Panonychus citri</i> . <i>Pesticide Biochemistry and Physiology</i> , 2020, 164, 73-84.	1.6	21
39	Costs and benefits of multiple mating in a species with first-male sperm precedence. <i>Journal of Animal Ecology</i> , 2020, 89, 1045-1054.	1.3	16
40	Acaricide resistance status and identification of resistance mutations in populations of the two-spotted spider mite <i>Tetranychus urticae</i> from Ethiopia. <i>Experimental and Applied Acarology</i> , 2020, 82, 475-491.	0.7	16
41	Genome-enabled insights into the biology of thrips as crop pests. <i>BMC Biology</i> , 2020, 18, 142.	1.7	54
42	Diversity and evolution of the P450 family in arthropods. <i>Insect Biochemistry and Molecular Biology</i> , 2020, 127, 103490.	1.2	109
43	Cover Image, Volume 76, Issue 8. <i>Pest Management Science</i> , 2020, 76, .	1.7	0
44	QTL mapping using microsatellite linkage reveals target-site mutations associated with high levels of resistance against three mitochondrial complex II inhibitors in <i>Tetranychus urticae</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2020, 123, 103410.	1.2	36
45	Using CRISPR/Cas9 genome modification to understand the genetic basis of insecticide resistance: <i>Drosophila</i> and beyond. <i>Pesticide Biochemistry and Physiology</i> , 2020, 167, 104595.	1.6	36
46	Geographical distribution and molecular insights into abamectin and milbemectin cross-resistance in European field populations of <i>Tetranychus urticae</i> . <i>Pest Management Science</i> , 2020, 76, 2569-2581.	1.7	47
47	Resistance risk assessment of the novel complex II inhibitor pyflubumide in the polyphagous pest <i>Tetranychus urticae</i> . <i>Journal of Pest Science</i> , 2020, 93, 1085-1096.	1.9	18
48	Targeted mutagenesis using CRISPR-Cas9 in the chelicerate herbivore <i>Tetranychus urticae</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2020, 120, 103347.	1.2	49
49	Identifying drivers of spatio-temporal dynamics in barley yellow dwarf virus epidemiology as a critical factor in disease control. <i>Pest Management Science</i> , 2020, 76, 2548-2556.	1.7	11
50	Improving the compatibility of pesticides and predatory mites: recent findings on physiological and ecological selectivity. <i>Current Opinion in Insect Science</i> , 2020, 39, 63-68.	2.2	29
51	Significance and interpretation of molecular diagnostics for insecticide resistance management of agricultural pests. <i>Current Opinion in Insect Science</i> , 2020, 39, 69-76.	2.2	64
52	Metabolic mechanisms of resistance to spiroticlofen and spiromesifen in Iranian populations of <i>Panonychus ulmi</i> . <i>Crop Protection</i> , 2020, 134, 105166.	1.0	21
53	Identification and functional characterization of a novel acetyl-CoA carboxylase mutation associated with ketoenol resistance in <i>Bemisia tabaci</i> . <i>Pesticide Biochemistry and Physiology</i> , 2020, 166, 104583.	1.6	28
54	Genome streamlining in a minute herbivore that manipulates its host plant. <i>ELife</i> , 2020, 9, .	2.8	33

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55	Trait mapping in diverse arthropods by bulked segregant analysis. <i>Current Opinion in Insect Science</i> , 2019, 36, 57-65.	2.2	32
56	Convergent evolution of cytochrome P450s underlies independent origins of keto-carotenoid pigmentation in animals. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2019, 286, 20191039.	1.2	28
57	Characterization of abamectin resistance in Iranian populations of European red mite, <i>Panonychus ulmi</i> Koch (Acari: Tetranychidae). <i>Crop Protection</i> , 2019, 125, 104903.	1.0	21
58	Resistance incidence and presence of resistance mutations in populations of <i>Tetranychus urticae</i> from vegetable crops in Turkey. <i>Experimental and Applied Acarology</i> , 2019, 78, 343-360.	0.7	30
59	High-resolution QTL mapping in <i>Tetranychus urticae</i> reveals acaricide-specific responses and common target-site resistance after selection by different MET-I acaricides. <i>Insect Biochemistry and Molecular Biology</i> , 2019, 110, 19-33.	1.2	62
60	Point mutations in the voltage-gated sodium channel gene associated with pyrethroid resistance in Iranian populations of the European red mite <i>Panonychus ulmi</i> . <i>Pesticide Biochemistry and Physiology</i> , 2019, 157, 80-87.	1.6	16
61	Long-Term Population Studies Uncover the Genome Structure and Genetic Basis of Xenobiotic and Host Plant Adaptation in the Herbivore <i>Tetranychus urticae</i> . <i>Genetics</i> , 2019, 211, 1409-1427.	1.2	70
62	Substrate specificity and promiscuity of horizontally transferred UDP-glycosyltransferases in the generalist herbivore <i>Tetranychus urticae</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2019, 109, 116-127.	1.2	38
63	Genome-wide gene expression profiling reveals that cuticle alterations and P450 detoxification are associated with deltamethrin and DDT resistance in <i>Anopheles arabiensis</i> populations from Ethiopia. <i>Pest Management Science</i> , 2019, 75, 1808-1818.	1.7	42
64	Structural and functional characterization of an intradiol ring-cleavage dioxygenase from the polyphagous spider mite herbivore <i>Tetranychus urticae</i> Koch. <i>Insect Biochemistry and Molecular Biology</i> , 2019, 107, 19-30.	1.2	6
65	A massive incorporation of microbial genes into the genome of <i>Tetranychus urticae</i> , a polyphagous arthropod herbivore. <i>Insect Molecular Biology</i> , 2018, 27, 333-351.	1.0	40
66	Does host plant adaptation lead to pesticide resistance in generalist herbivores?. <i>Current Opinion in Insect Science</i> , 2018, 26, 25-33.	2.2	74
67	The role of glutathione S-transferases (GSTs) in insecticide resistance in crop pests and disease vectors. <i>Current Opinion in Insect Science</i> , 2018, 27, 97-102.	2.2	197
68	Fitness costs of key point mutations that underlie acaricide target-site resistance in the two-spotted spider mite <i>Tetranychus urticae</i> . <i>Evolutionary Applications</i> , 2018, 11, 1540-1553.	1.5	40
69	Molecular characterization of pyrethroid resistance in the olive fruit fly <i>Bactrocera oleae</i> . <i>Pesticide Biochemistry and Physiology</i> , 2018, 148, 1-7.	1.6	16
70	A Gene Family Coding for Salivary Proteins (SHOT) of the Polyphagous Spider Mite <i>Tetranychus urticae</i> Exhibits Fast Host-Dependent Transcriptional Plasticity. <i>Molecular Plant-Microbe Interactions</i> , 2018, 31, 112-124.	1.4	29
71	Kin competition accelerates experimental range expansion in an arthropod herbivore. <i>Ecology Letters</i> , 2018, 21, 225-234.	3.0	46
72	Draft Genome Assembly of the Poultry Red Mite, <i>Dermanyssus gallinae</i> . <i>Microbiology Resource Announcements</i> , 2018, 7, .	0.3	26

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73	Transcriptomic Plasticity in the Arthropod Generalist <i>Tetranychus urticae</i> Upon Long-Term Acclimation to Different Host Plants. <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 3865-3879.	0.8	36

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91	Resistance mutation conserved between insects and mites unravels the benzoylurea insecticide mode of action on chitin biosynthesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 14692-14697.	3.3	144
92	Complex Evolutionary Dynamics of Massively Expanded Chemosensory Receptor Families in an Extreme Generalist Chelicerate Herbivore. <i>Genome Biology and Evolution</i> , 2016, 8, 3323-3339.	1.1	42
93	Horizontal Gene Transfer Contributes to the Evolution of Arthropod Herbivory. <i>Genome Biology and Evolution</i> , 2016, 8, 1785-1801.	1.1	155
94	The Salivary Protein Repertoire of the Polyphagous Spider Mite <i>Tetranychus urticae</i> : A Quest for Effectors. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 3594-3613.	2.5	113
95	Salivary proteins of spider mites suppress defenses in <i>Nicotiana benthamiana</i> and promote mite reproduction. <i>Plant Journal</i> , 2016, 86, 119-131.	2.8	149
96	Comparative genome-wide transcriptome analysis of <i>Vitis vinifera</i> responses to adapted and non-adapted strains of two-spotted spider mite, <i>Tetranychus urticae</i> . <i>BMC Genomics</i> , 2016, 17, 74.	1.2	53
97	The Molecular Evolution of Xenobiotic Metabolism and Resistance in Chelicerate Mites. <i>Annual Review of Entomology</i> , 2016, 61, 475-498.	5.7	227
98	Molecular analysis of cyenopyrafen resistance in the two-spotted spider mite <i>Tetranychus urticae</i> . <i>Pest Management Science</i> , 2016, 72, 103-112.	1.7	60
99	Incidence and characterization of resistance to pyrethroid and organophosphorus insecticides in <i>Thrips tabaci</i> (Thysanoptera: Thripidae) in onion fields in Isfahan, Iran. <i>Pesticide Biochemistry and Physiology</i> , 2016, 129, 28-35.	1.6	19
100	Mutations in chitin synthase-1 (CHS-1) confer resistance to a range of structurally diverse acaricides and insecticides. , 2016, , .		0
101	Adaptation of a polyphagous herbivore to a novel host plant extensively shapes the transcriptome of herbivore and host. <i>Molecular Ecology</i> , 2015, 24, 4647-4663.	2.0	131
102	Feeding History Affects Intraguild Interactions between <i>Harmonia axyridis</i> (Coleoptera: Coccinellidae) and <i>Episyrphus balteatus</i> (Diptera: Syrphidae). <i>PLoS ONE</i> , 2015, 10, e0128518.	1.1	9
103	The economic importance of acaricides in the control of phytophagous mites and an update on recent acaricide mode of action research. <i>Pesticide Biochemistry and Physiology</i> , 2015, 121, 12-21.	1.6	238
104	Mechanisms and ecological consequences of plant defence induction and suppression in herbivore communities. <i>Annals of Botany</i> , 2015, 115, 1015-1051.	1.4	244
105	Transcriptome profiling of a spiroadiclofen susceptible and resistant strain of the European red mite <i>Panonychus ulmi</i> using strand-specific RNA-seq. <i>BMC Genomics</i> , 2015, 16, 974.	1.2	54
106	Tomato Whole Genome Transcriptional Response to <i>Tetranychus urticae</i> Identifies Divergence of Spider Mite-Induced Responses Between Tomato and <i>Arabidopsis</i> . <i>Molecular Plant-Microbe Interactions</i> , 2015, 28, 343-361.	1.4	90
107	Functional characterization of glutathione S-transferases associated with insecticide resistance in <i>Tetranychus urticae</i> . <i>Pesticide Biochemistry and Physiology</i> , 2015, 121, 53-60.	1.6	69
108	Genotype to phenotype, the molecular and physiological dimensions of resistance in arthropods. <i>Pesticide Biochemistry and Physiology</i> , 2015, 121, 61-77.	1.6	237

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109	Empirically simulated spatial sorting points at fast epigenetic changes in dispersal behaviour. <i>Evolutionary Ecology</i> , 2015, 29, 299-310.	0.5	23
110	Functional characterization of the <i>Tetranychus urticae</i> CYP392A11, a cytochrome P450 that hydroxylates the METI acaricides cyenopyrafen and fenpyroximate. <i>Insect Biochemistry and Molecular Biology</i> , 2015, 65, 91-99.	1.2	72
111	Genome sequence of the Asian Tiger mosquito, <i>Aedes albopictus</i> , reveals insights into its biology, genetics, and evolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E5907-15.	3.3	251
112	Intraguild predation by <i>Armonia axyridis</i> (Coleoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 627 Td (<sc>H</sc> armonia axyridis) (<sc>C</sc>oleoptera:) samples. <i>Entomological Science</i> , 2015, 18, 130-133.	0.3	40
113	Reciprocal Responses in the Interaction between <i>Arabidopsis</i> and the Cell-Content-Feeding Chelicerate Herbivore Spider Mite <i>Tetranychus urticae</i> . <i>Plant Physiology</i> , 2014, 164, 384-399.	2.3	151
114	Cross-resistance risk of the novel complex <i>Tetranychus urticae</i> inhibitors cyenopyrafen and cyflumetofen in resistant strains of the two-spotted spider mite <i>Tetranychus urticae</i> . <i>Pest Management Science</i> , 2014, 70, 365-368.	1.7	59
115	Abamectin is metabolized by CYP392A16, a cytochrome P450 associated with high levels of acaricide resistance in <i>Tetranychus urticae</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2014, 46, 43-53.	1.2	155
116	Bacterial origin of a diverse family of UDP-glycosyltransferase genes in the <i>Tetranychus urticae</i> genome. <i>Insect Biochemistry and Molecular Biology</i> , 2014, 50, 43-57.	1.2	59
117	The ABC gene family in arthropods: Comparative genomics and role in insecticide transport and resistance. <i>Insect Biochemistry and Molecular Biology</i> , 2014, 45, 89-110.	1.2	462
118	Fitness maximization by dispersal: evidence from an invasion experiment. <i>Ecology</i> , 2014, 95, 3104-3111.	1.5	38
119	The cyclic keto-enol insecticide spirotetramat inhibits insect and spider mite acetyl-CoA carboxylases by interfering with the carboxyltransferase partial reaction. <i>Insect Biochemistry and Molecular Biology</i> , 2014, 55, 1-8.	1.2	76
120	High resolution genetic mapping uncovers chitin synthase-1 as the target-site of the structurally diverse mite growth inhibitors clofentezine, hexythiazox and etoxazole in <i>Tetranychus urticae</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2014, 51, 52-61.	1.2	83
121	A gene horizontally transferred from bacteria protects arthropods from host plant cyanide poisoning. <i>ELife</i> , 2014, 3, e02365.	2.8	135
122	Application of Two-spotted Spider Mite <i>Tetranychus urticae</i> for Plant-pest Interaction Studies. <i>Journal of Visualized Experiments</i> , 2014, , .	0.2	43
123	Spider mite control and resistance management: does a genome help?. <i>Pest Management Science</i> , 2013, 69, 156-159.	1.7	50
124	A burst of ABC genes in the genome of the polyphagous spider mite <i>Tetranychus urticae</i> . <i>BMC Genomics</i> , 2013, 14, 317.	1.2	118
125	Genome wide gene-expression analysis of facultative reproductive diapause in the two-spotted spider mite <i>Tetranychus urticae</i> . <i>BMC Genomics</i> , 2013, 14, 815.	1.2	92
126	Molecular analysis of resistance to acaricidal spirocyclic tetrone acids in <i>Tetranychus urticae</i> : CYP392E10 metabolizes spirodiclofen, but not its corresponding enol. <i>Insect Biochemistry and Molecular Biology</i> , 2013, 43, 544-554.	1.2	107

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127	A link between host plant adaptation and pesticide resistance in the polyphagous spider mite <i>Tetranychus urticae</i> . Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E113-22.	3.3	347
128	Population bulk segregant mapping uncovers resistance mutations and the mode of action of a chitin synthesis inhibitor in arthropods. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 4407-4412.	3.3	240
129	On the mode of action of bifenthrin: New evidence for a mitochondrial target site. Pesticide Biochemistry and Physiology, 2012, 104, 88-95.	1.6	39
130	The cys-loop ligand-gated ion channel gene family of <i>Tetranychus urticae</i> : Implications for acaricide toxicology and a novel mutation associated with abamectin resistance. Insect Biochemistry and Molecular Biology, 2012, 42, 455-465.	1.2	161
131	A horizontally transferred cyanase gene in the spider mite <i>Tetranychus urticae</i> is involved in cyanate metabolism and is differentially expressed upon host plant change. Insect Biochemistry and Molecular Biology, 2012, 42, 881-889.	1.2	40
132	Resistance to acaricides in Italian strains of <i>Tetranychus urticae</i> : toxicological and enzymatic assays. Experimental and Applied Acarology, 2012, 57, 53-64.	0.7	36
133	The genome of <i>Tetranychus urticae</i> reveals herbivorous pest adaptations. Nature, 2011, 479, 487-492.	13.7	897
134	Parallel evolution of cytochrome <i>b</i> mediated bifenthrin resistance in the citrus red mite <i>Panonychus citri</i> . Insect Molecular Biology, 2011, 20, 135-140.	1.0	51
135	Acaricide resistance and resistance mechanisms in <i>Tetranychus urticae</i> populations from rose greenhouses in the Netherlands. Pest Management Science, 2011, 67, 1424-1433.	1.7	108
136	The control of eriophyoid mites: state of the art and future challenges. Experimental and Applied Acarology, 2010, 51, 205-224.	0.7	70
137	Acetylcholinesterase point mutations in European strains of <i>Tetranychus urticae</i> (Acari: Tj ETQq1 1 0.784314,rgBT /Overlock 10 Tf 50 100)	1.7	87
138	Acaricide resistance mechanisms in the two-spotted spider mite <i>Tetranychus urticae</i> and other important Acari: A review. Insect Biochemistry and Molecular Biology, 2010, 40, 563-572.	1.2	626
139	Effects of spirodiclofen on reproduction in a susceptible and resistant strain of <i>Tetranychus urticae</i> (Acari: Tetranychidae). Experimental and Applied Acarology, 2009, 47, 301-309.	0.7	53
140	Susceptibility of an organophosphate resistant strain of the two-spotted spider mite (<i>Tetranychus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 and Applied Acarology, 2009, 49, 185-192.	0.7	13
141	Genetic and biochemical analysis of a laboratory-selected spirodiclofen-resistant strain of <i>Tetranychus urticae</i> Koch (Acari: Tetranychidae). Pest Management Science, 2009, 65, 358-366.	1.7	105
142	Mutations in the mitochondrial cytochrome <i>b</i> of <i>Tetranychus urticae</i> Koch (Acari: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 100) 2009, 65, 404-412.	1.7	95
143	Identification of pyrethroid resistance associated mutations in the <i>para</i> sodium channel of the two-spotted spider mite <i>Tetranychus urticae</i> (Acari: Tetranychidae). Insect Molecular Biology, 2009, 18, 583-593.	1.0	99
144	Mechanisms of Acaricide Resistance in the Two-Spotted Spider Mite <i>Tetranychus urticae</i> . , 2009, , 347-393.		66

#	ARTICLE	IF	CITATIONS
145	Resistance mechanisms to mitochondrial electron transport inhibitors in a field-collected strain of <i>Tetranychus urticae</i> Koch (Acari: Tetranychidae). <i>Bulletin of Entomological Research</i> , 2009, 99, 23-31.	0.5	107
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150	Susceptibility of the predatory stinkbug <i>Picromerus bidens</i> to selected insecticides. <i>BioControl</i> , 2007, 52, 765-774.	0.9	22
151	Complete maternal inheritance of bifenthrin resistance in <i>Tetranychus urticae</i> Koch (Acari: Tetranychidae). <i>Molecular Biology and Evolution</i> , 2006, 23, 869-877.	1.2	89
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154	Systemic Use of Spinosad to Control the Two-spotted Spider Mite (Acari: Tetranychidae) on Tomatoes Grown in Rockwool. <i>Experimental and Applied Acarology</i> , 2005, 37, 93-105.	0.7	39
155	Comparative acaricide susceptibility and detoxifying enzyme activities in field-collected resistant and susceptible strains of <i>Tetranychus urticae</i> . <i>Pest Management Science</i> , 2005, 61, 499-507.	1.7	171
156	Genetic analysis and cross-resistance spectrum of a laboratory-selected chlorfenapyr resistant strain of two-spotted spider mite (Acari: Tetranychidae). <i>Experimental and Applied Acarology</i> , 2004, 32, 249-261.	0.7	147