

John G Vontas

List of Publications by Year in descending order

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226
papers

14,856
citations

23500

58
h-index

26548

107
g-index

245
all docs

245
docs citations

245
times ranked

9173
citing authors

#	ARTICLE	IF	CITATIONS
1	Acaricide resistance mechanisms in the two-spotted spider mite <i>Tetranychus urticae</i> and other important Acari: A review. <i>Insect Biochemistry and Molecular Biology</i> , 2010, 40, 563-572.	1.2	626
2	Contemporary status of insecticide resistance in the major <i>Aedes</i> vectors of arboviruses infecting humans. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005625.	1.3	504
3	Highly evolvable malaria vectors: The genomes of 16 <i>Anopheles</i> mosquitoes. <i>Science</i> , 2015, 347, 1258522.	6.0	492
4	Glutathione S-transferases as antioxidant defence agents confer pyrethroid resistance in <i>Nilaparvata lugens</i> . <i>Biochemical Journal</i> , 2001, 357, 65-72.	1.7	437
5	Over-expression of cytochrome P450 CYP6CM1 is associated with high resistance to imidacloprid in the B and Q biotypes of <i>Bemisia tabaci</i> (Hemiptera: Aleyrodidae). <i>Insect Biochemistry and Molecular Biology</i> , 2008, 38, 634-644.	1.2	349
6	A link between host plant adaptation and pesticide resistance in the polyphagous spider mite <i>Tetranychus urticae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E113-22.	3.3	347
7	Insecticide resistance in the major dengue vectors <i>Aedes albopictus</i> and <i>Aedes aegypti</i> . <i>Pesticide Biochemistry and Physiology</i> , 2012, 104, 126-131.	1.6	292
8	Glutathione S-transferases as antioxidant defence agents confer pyrethroid resistance in <i>Nilaparvata lugens</i> . <i>Biochemical Journal</i> , 2001, 357, 65.	1.7	284
9	The <i>Anopheles gambiae</i> detoxification chip: A highly specific microarray to study metabolic-based insecticide resistance in malaria vectors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 4080-4084.	3.3	282
10	Cytochrome P450 associated with insecticide resistance catalyzes cuticular hydrocarbon production in <i>Anopheles gambiae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 9268-9273.	3.3	279
11	Detection of knockdown resistance (<i>kdr</i>) mutations in <i>Anopheles gambiae</i> : a comparison of two new high-throughput assays with existing methods. <i>Malaria Journal</i> , 2007, 6, 111.	0.8	273
12	An Overview of Insecticide Resistance. <i>Science</i> , 2002, 298, 96-97.	6.0	269
13	Insect cuticle: a critical determinant of insecticide resistance. <i>Current Opinion in Insect Science</i> , 2018, 27, 68-74.	2.2	264
14	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
15	Cross-induction of detoxification genes by environmental xenobiotics and insecticides in the mosquito <i>Aedes aegypti</i> : Impact on larval tolerance to chemical insecticides. <i>Insect Biochemistry and Molecular Biology</i> , 2008, 38, 540-551.	1.2	246
16	Structural model and functional characterization of the <i>Bemisia tabaci</i> CYP6CM1vQ, a cytochrome P450 associated with high levels of imidacloprid resistance. <i>Insect Biochemistry and Molecular Biology</i> , 2009, 39, 697-706.	1.2	204
17	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
18	Gene expression in insecticide resistant and susceptible <i>Anopheles gambiae</i> strains constitutively or after insecticide exposure. <i>Insect Molecular Biology</i> , 2005, 14, 509-521.	1.0	183

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19	Heterologous expression of four glutathione transferase genes genetically linked to a major insecticide-resistance locus from the malaria vector <i>Anopheles gambiae</i> . <i>Biochemical Journal</i> , 2003, 373, 957-963.	1.7	166
20	Alternative strategies for mosquito-borne arbovirus control. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0006822.	1.3	165
21	Gene Amplification, ABC Transporters and Cytochrome P450s: Unraveling the Molecular Basis of Pyrethroid Resistance in the Dengue Vector, <i>Aedes aegypti</i> . <i>PLoS Neglected Tropical Diseases</i> , 2012, 6, e1692.	1.3	163
22	Management of insecticide resistance in the major <i>Aedes</i> vectors of arboviruses: Advances and challenges. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007615.	1.3	162
23	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. <i>Insect Biochemistry and Molecular Biology</i> , 2012, 42, 455-465.	1.2	161
24	Purification, molecular cloning and heterologous expression of a glutathione S-transferase involved in insecticide resistance from the rice brown planthopper, <i>Nilaparvata lugens</i> . <i>Biochemical Journal</i> , 2002, 362, 329-337.	1.7	158
25	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
26	The Role of Cytochrome P450s in Insect Toxicology and Resistance. <i>Annual Review of Entomology</i> , 2022, 67, 105-124.	5.7	149
27	Insecticide resistance in the tomato pinworm <i>Tuta absoluta</i> : patterns, spread, mechanisms, management and outlook. <i>Journal of Pest Science</i> , 2019, 92, 1329-1342.	1.9	147
28	Resistance-associated point mutations of organophosphate insensitive acetylcholinesterase, in the olive fruit fly <i>Bactrocera oleae</i> . <i>Insect Molecular Biology</i> , 2002, 11, 329-336.	1.0	144
29	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
30	PCR-based detection of <i>Plasmodium</i> in <i>Anopheles</i> mosquitoes: a comparison of a new high-throughput assay with existing methods. <i>Malaria Journal</i> , 2008, 7, 177.	0.8	129
31	A sensory appendage protein protects malaria vectors from pyrethroids. <i>Nature</i> , 2020, 577, 376-380.	13.7	129
32	Control of the olive fruit fly using genetics-enhanced sterile insect technique. <i>BMC Biology</i> , 2012, 10, 51.	1.7	128
33	Transcriptional analysis of insecticide resistance in <i>Anopheles stephensi</i> using cross-species microarray hybridization. <i>Insect Molecular Biology</i> , 2007, 16, 315-324.	1.0	124
34	Current status of insecticide resistance in Q biotype <i>Bemisia tabaci</i> populations from Crete. <i>Pest Management Science</i> , 2009, 65, 313-322.	1.7	123
35	Ryanodine receptor point mutations confer diamide insecticide resistance in tomato leafminer, <i>Tuta absoluta</i> (Lepidoptera: Gelechiidae). <i>Insect Biochemistry and Molecular Biology</i> , 2017, 80, 11-20.	1.2	122
36	Contributions of cuticle permeability and enzyme detoxification to pyrethroid resistance in the major malaria vector <i>Anopheles gambiae</i> . <i>Scientific Reports</i> , 2017, 7, 11091.	1.6	117

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37	Insecticide resistance in Tephritid flies. <i>Pesticide Biochemistry and Physiology</i> , 2011, 100, 199-205.	1.6	116
38	Disruption of a horizontally transferred phytoene desaturase abolishes carotenoid accumulation and diapause in <i>Tetranychus urticae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E5871-E5880.	3.3	114
39	Molecular analysis of resistance to acaricidal spirocyclic tetrone acids in <i>Tetranychus urticae</i> : CYP392E10 metabolizes spiropdiclofen, but not its corresponding enol. <i>Insect Biochemistry and Molecular Biology</i> , 2013, 43, 544-554.	1.2	107
40	Transcription profiling of eleven cytochrome P450s potentially involved in xenobiotic metabolism in the mosquito <i>Aedes aegypti</i> . <i>Insect Molecular Biology</i> , 2010, 19, 185-193.	1.0	103
41	Identification of pyrethroid resistance associated mutations in the α sodium channel of the two-spotted spider mite <i>Tetranychus urticae</i> (Acari: Tetranychidae). <i>Insect Molecular Biology</i> , 2009, 18, 583-593.	1.0	99
42	Purification, molecular cloning and heterologous expression of a glutathione S-transferase involved in insecticide resistance from the rice brown planthopper, <i>Nilaparvata lugens</i> . <i>Biochemical Journal</i> , 2002, 362, 329.	1.7	94
43	Characterization of inhibitors and substrates of <i>Anopheles gambiae</i> CYP6Z2. <i>Insect Molecular Biology</i> , 2008, 17, 125-135.	1.0	92
44	Pymetrozine is hydroxylated by <i>CYP6CM1</i> , a cytochrome P450 conferring neonicotinoid resistance in <i>Bemisia tabaci</i> . <i>Pest Management Science</i> , 2013, 69, 457-461.	1.7	88
45	Rapid selection of a pyrethroid metabolic enzyme CYP9K1 by operational malaria control activities. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 4619-4624.	3.3	88
46	Acetylcholinesterase point mutations in European strains of <i>Tetranychus urticae</i> (Acari: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 38	1.7	87
47	A mutation in the PSST homologue of complex I (NADH:ubiquinone oxidoreductase) from <i>Tetranychus urticae</i> is associated with resistance to METI acaricides. <i>Insect Biochemistry and Molecular Biology</i> , 2017, 80, 79-90.	1.2	82
48	The relative contribution of target-site mutations in complex acaricide resistant phenotypes as assessed by marker assisted backcrossing in <i>Tetranychus urticae</i> . <i>Scientific Reports</i> , 2017, 7, 9202.	1.6	81
49	Truncated transcripts of nicotinic acetylcholine subunit gene <i>Bd1±6</i> are associated with spinosad resistance in <i>Bactrocera dorsalis</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2012, 42, 806-815.	1.2	79
50	Comparison of esterase gene amplification, gene expression and esterase activity in insecticide susceptible and resistant strains of the brown planthopper, <i>Nilaparvata lugens</i> (Stal). <i>Insect Molecular Biology</i> , 2000, 9, 655-660.	1.0	78
51	Country-level operational implementation of the Global Plan for Insecticide Resistance Management. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 9397-9402.	3.3	76
52	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
53	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
54	A glutathione-S-transferase (TuGSTd05) associated with acaricide resistance in <i>Tetranychus urticae</i> directly metabolizes the complex II inhibitor cyflumetofen. <i>Insect Biochemistry and Molecular Biology</i> , 2017, 80, 101-115.	1.2	68

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55	Global distribution and origin of target site insecticide resistance mutations in <i>Tetranychus urticae</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2014, 48, 17-28.	1.2	67
56	Mechanisms of Acaricide Resistance in the Two-Spotted Spider Mite <i>Tetranychus urticae</i> . , 2009, , 347-393.		66
57	Investigation of the contribution of RyR target-site mutations in diamide resistance by CRISPR/Cas9 genome modification in <i>Drosophila</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2017, 87, 127-135.	1.2	66
58	Genomewide transcriptional signatures of migratory flight activity in a globally invasive insect pest. <i>Molecular Ecology</i> , 2015, 24, 4901-4911.	2.0	65
59	Altered Acetylcholinesterase Confers Organophosphate Resistance in the Olive Fruit Fly <i>Bactrocera oleae</i> . <i>Pesticide Biochemistry and Physiology</i> , 2001, 71, 124-132.	1.6	64
60	A Simple Colorimetric Assay for Specific Detection of Glutathione-S Transferase Activity Associated with DDT Resistance in Mosquitoes. <i>PLoS Neglected Tropical Diseases</i> , 2010, 4, e808.	1.3	64
61	The <i>Anopheles gambiae</i> ATP-binding cassette transporter family: phylogenetic analysis and tissue localization provide clues on function and role in insecticide resistance. <i>Insect Molecular Biology</i> , 2018, 27, 110-122.	1.0	64
62	Cytochrome P450-based metabolic insecticide resistance in <i>Anopheles</i> and <i>Aedes</i> mosquito vectors: Muddying the waters. <i>Pesticide Biochemistry and Physiology</i> , 2020, 170, 104666.	1.6	64
63	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
64	Dissecting the organ specificity of insecticide resistance candidate genes in <i>Anopheles gambiae</i> : known and novel candidate genes. <i>BMC Genomics</i> , 2014, 15, 1018.	1.2	63
65	Assessment of the <i>Bemisia tabaci</i> CYP6CM1vQ transcript and protein levels in laboratory and field-derived imidacloprid-resistant insects and cross-metabolism potential of the recombinant enzyme. <i>Insect Science</i> , 2011, 18, 23-29.	1.5	62
66	Large-scale field trial of attractive toxic sugar baits (ATSB) for the control of malaria vector mosquitoes in Mali, West Africa. <i>Malaria Journal</i> , 2020, 19, 72.	0.8	61
67	Development of high-throughput real-time PCR assays for the identification of insensitive acetylcholinesterase (ace-1R) in <i>Anopheles gambiae</i> . <i>Pesticide Biochemistry and Physiology</i> , 2010, 96, 80-85.	1.6	60
68	Transcription analysis of neonicotinoid resistance in Mediterranean (MED) populations of <i>B. tabaci</i> reveal novel cytochrome P450s, but no nAChR mutations associated with the phenotype. <i>BMC Genomics</i> , 2015, 16, 939.	1.2	59
69	Transcriptomic responses of the olive fruit fly <i>Bactrocera oleae</i> and its symbiont <i>Candidatus Erwinia dacicola</i> to olive feeding. <i>Scientific Reports</i> , 2017, 7, 42633.	1.6	58
70	Genetic elimination of field-cage populations of Mediterranean fruit flies. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2014, 281, 20141372.	1.2	57
71	Mosquitoes cloak their legs to resist insecticides. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2019, 286, 20191091.	1.2	56
72	Global patterns in genomic diversity underpinning the evolution of insecticide resistance in the aphid crop pest <i>Myzus persicae</i> . <i>Communications Biology</i> , 2021, 4, 847.	2.0	55

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73	Analysis of the Olive Fruit Fly <i>Bactrocera oleae</i> Transcriptome and Phylogenetic Classification of the Major Detoxification Gene Families. <i>PLoS ONE</i> , 2013, 8, e66533.	1.1	55
74	How do oral insecticidal compounds cross the insect midgut epithelium?. <i>Insect Biochemistry and Molecular Biology</i> , 2018, 103, 22-35.	1.2	54
75	Detection of resistance-associated point mutations of organophosphate-insensitive acetylcholinesterase in the olive fruit fly, <i>Bactrocera oleae</i> (Gmelin). <i>Pesticide Biochemistry and Physiology</i> , 2005, 81, 154-163.	1.6	53
76	Identification of mutations in the para sodium channel of <i>Bemisia tabaci</i> from Crete, associated with resistance to pyrethroids. <i>Pesticide Biochemistry and Physiology</i> , 2006, 85, 161-166.	1.6	53
77	Insecticide resistance status of the codling moth <i>Cydia pomonella</i> (Lepidoptera: Tortricidae) from Greece. <i>Pesticide Biochemistry and Physiology</i> , 2011, 100, 229-238.	1.6	53
78	Biological and molecular characterization of laboratory mutants of <i>Cercospora beticola</i> resistant to Qo inhibitors. <i>European Journal of Plant Pathology</i> , 2006, 116, 155-166.	0.8	52
79	Molecular characterization and detection of overexpressed C-14 alpha-demethylase-based DMI resistance in <i>Cercospora beticola</i> field isolates. <i>Pesticide Biochemistry and Physiology</i> , 2009, 95, 18-27.	1.6	52
80	Striking diflubenzuron resistance in <i>Culex pipiens</i> , the prime vector of West Nile Virus. <i>Scientific Reports</i> , 2017, 7, 11699.	1.6	52
81	Resurgence of the cotton bollworm <i>Helicoverpa armigera</i> in northern Greece associated with insecticide resistance. <i>Insect Science</i> , 2013, 20, 505-512.	1.5	51
82	Insecticide resistance is mediated by multiple mechanisms in recently introduced <i>Aedes aegypti</i> from Madeira Island (Portugal). <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005799.	1.3	51
83	Insecticide resistance in <i>Bemisia tabaci</i> from Cyprus. <i>Insect Science</i> , 2011, 18, 30-39.	1.5	50
84	Multiple recombination events between two cytochrome P450 loci contribute to global pyrethroid resistance in <i>Helicoverpa armigera</i> . <i>PLoS ONE</i> , 2018, 13, e0197760.	1.1	50
85	Rapid multiplex gene expression assays for monitoring metabolic resistance in the major malaria vector <i>Anopheles gambiae</i> . <i>Parasites and Vectors</i> , 2019, 12, 9.	1.0	50
86	Framework for rapid assessment and adoption of new vector control tools. <i>Trends in Parasitology</i> , 2014, 30, 191-204.	1.5	49
87	Transcriptome Profiling and Genetic Study Reveal Amplified Carboxylesterase Genes Implicated in Temephos Resistance, in the Asian Tiger Mosquito <i>Aedes albopictus</i> . <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0003771.	1.3	49
88	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
89	Efficacy of the pyrethroid alpha-cypermethrin against <i>Bactrocera oleae</i> populations from Greece, and improved diagnostic for an iAChE mutation. <i>Pest Management Science</i> , 2008, 64, 900-908.	1.7	45
90	Engineering sensitive glutathione transferase for the detection of xenobiotics. <i>Biosensors and Bioelectronics</i> , 2008, 24, 498-503.	5.3	45

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91	Distribution and hybridization of <i>Culex pipiens</i> forms in Greece during the West Nile virus outbreak of 2010. <i>Infection, Genetics and Evolution</i> , 2013, 16, 218-225.	1.0	45
92	Development of a lateral flow test to detect metabolic resistance in <i>Bemisia tabaci</i> mediated by CYP6CM1, a cytochrome P450 with broad spectrum catalytic efficiency. <i>Pesticide Biochemistry and Physiology</i> , 2015, 121, 3-11.	1.6	44
93	Two functionally distinct CYP4G genes of <i>Anopheles gambiae</i> contribute to cuticular hydrocarbon biosynthesis. <i>Insect Biochemistry and Molecular Biology</i> , 2019, 110, 52-59.	1.2	44
94	Tracking Insecticide Resistance in Mosquito Vectors of Arboviruses: The Worldwide Insecticide resistance Network (WIN). <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0005054.	1.3	43
95	A Simple Biochemical Assay for Glutathione S-Transferase Activity and Its Possible Field Application for Screening Glutathione S-Transferase-Based Insecticide Resistance. <i>Pesticide Biochemistry and Physiology</i> , 2000, 68, 184-192.	1.6	42
96	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
97	“What I cannot create, I do not understand”: functionally validated synergism of metabolic and target site insecticide resistance. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2020, 287, 20200838.	1.2	42
98	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
99	Identification of a novel point mutation in the β -tubulin gene of <i>Botrytis cinerea</i> and detection of benzimidazole resistance by a diagnostic PCR-RFLP assay. <i>European Journal of Plant Pathology</i> , 2009, 125, 97-107.	0.8	41
100	The Vector Population Monitoring Tool (VPMT): High-Throughput DNA-Based Diagnostics for the Monitoring of Mosquito Vector Populations. <i>Malaria Research and Treatment</i> , 2010, 2010, 1-8.	2.0	41
101	Molecular diagnostics for detecting pyrethroid and organophosphate resistance mutations in the Q biotype of the whitefly <i>Bemisia tabaci</i> (Hemiptera: Aleyrodidae). <i>Pesticide Biochemistry and Physiology</i> , 2009, 94, 49-54.	1.6	40
102	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. <i>Insect Biochemistry and Molecular Biology</i> , 2012, 42, 881-889.	1.2	40
103	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
104	Heterologous expression of insect P450 enzymes that metabolize xenobiotics. <i>Current Opinion in Insect Science</i> , 2021, 43, 78-84.	2.2	40
105	Insecticide resistance status in the major West Nile virus vector <i>Culex pipiens</i> from Greece. <i>Pest Management Science</i> , 2014, 70, 623-627.	1.7	38
106	Identification and characterization of abamectin resistance in <i>Tetranychus urticae</i> Koch populations from greenhouses in Turkey. <i>Crop Protection</i> , 2018, 112, 112-117.	1.0	38
107	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
108	Analysis of population structure and insecticide resistance in mosquitoes of the genus <i>Culex</i> , <i>Anopheles</i> and <i>Aedes</i> from different environments of Greece with a history of mosquito borne disease transmission. <i>Acta Tropica</i> , 2017, 174, 29-37.	0.9	37

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109	Molecular diagnostics for detecting pyrethroid and abamectin resistance mutations in <i>Tetranychus urticae</i> . <i>Pesticide Biochemistry and Physiology</i> , 2017, 135, 9-14.	1.6	37
110	The genetic architecture of a host shift: An adaptive walk protected an aphid and its endosymbiont from plant chemical defenses. <i>Science Advances</i> , 2020, 6, eaba1070.	4.7	37
111	Carboxylesterase gene amplifications associated with insecticide resistance in <i>Aedes albopictus</i> : Geographical distribution and evolutionary origin. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005533.	1.3	36
112	First evidence of resistance to pyrethroid insecticides in Italian <i>Aedes albopictus</i> populations 26 years after invasion. <i>Pest Management Science</i> , 2018, 74, 1319-1327.	1.7	36
113	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
114	Activity of flonicamid on the sweet potato whitefly <i>Bemisia tabaci</i> (Homoptera: Aleyrodidae) and its natural enemies. <i>Pest Management Science</i> , 2014, 70, 1460-1467.	1.7	35
115	Status of Insecticide Resistance and Its Mechanisms in <i>Anopheles gambiae</i> and <i>Anopheles coluzzii</i> Populations from Forest Settings in South Cameroon. <i>Genes</i> , 2019, 10, 741.	1.0	35
116	Molecular characterization of the amplified aldehyde oxidase from insecticide resistant <i>Culex quinquefasciatus</i> . <i>FEBS Journal</i> , 2002, 269, 768-779.	0.2	34
117	Affordable assays for genotyping single nucleotide polymorphisms in insects. <i>Insect Molecular Biology</i> , 2007, 16, 377-387.	1.0	33
118	Functional and immunohistochemical characterization of CCEae3a, a carboxylesterase associated with temephos resistance in the major arbovirus vectors <i>Aedes aegypti</i> and <i>Ae. albopictus</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2016, 74, 61-67.	1.2	33
119	Identification and detection of indoxacarb resistance mutations in the sodium channel of the tomato leafminer, <i>Tuta absoluta</i> . <i>Pest Management Science</i> , 2017, 73, 1679-1688.	1.7	33
120	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
121	Vertically transmitted rhabdoviruses are found across three insect families and have dynamic interactions with their hosts. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20162381.	1.2	32
122	The evolution of multiple insecticide resistance in UK populations of tomato leafminer, <i>Tuta absoluta</i> . <i>Pest Management Science</i> , 2019, 75, 2079-2085.	1.7	32
123	Detection and Monitoring of Insecticide Resistance Mutations in <i>Anopheles gambiae</i> : Individual vs Pooled Specimens. <i>Genes</i> , 2018, 9, 479.	1.0	31
124	Insecticide resistance status and mechanisms in <i>Aedes aegypti</i> populations from Senegal. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009393.	1.3	31
125	Draft Genome Sequence of the <i>Bactrocera oleae</i> Symbiont <i>Candidatus</i> <i>Erwinia dacicola</i> . <i>Genome Announcements</i> , 2016, 4, .	0.8	30
126	Insecticide resistance in <i>Trialeurodes vaporariorum</i> populations and novel diagnostics for <i>ksr</i> mutations. <i>Pest Management Science</i> , 2018, 74, 59-69.	1.7	30

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128	Mapping insecticide resistance and characterization of resistance mechanisms in <i>Anopheles arabiensis</i> (Diptera: Culicidae) in Ethiopia. <i>Parasites and Vectors</i> , 2017, 10, 407.	1.0	29
129	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 ketoenol acaricide spiroticlofen. <i>Pest Management Science</i> , 2020, 76, 1142-1153.	1.7	29
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133	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
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135	Geographical distribution and evolutionary history of organophosphate-resistant Ace alleles in the olive fly (<i>Bactrocera oleae</i>). <i>Insect Biochemistry and Molecular Biology</i> , 2006, 36, 593-602.	1.2	27
136	Focal distribution of diflubenzuron resistance mutations in <i>Culex pipiens</i> mosquitoes from Northern Italy. <i>Acta Tropica</i> , 2019, 193, 106-112.	0.9	27
137	Only a minority of broad-range detoxification genes respond to a variety of phytotoxins in generalist <i>Bemisia tabaci</i> species. <i>Scientific Reports</i> , 2015, 5, 17975.	1.6	26
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139	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
140	Transgenic expression of the <i>Aedes aegypti</i> CYP9J28 confers pyrethroid resistance in <i>Drosophila melanogaster</i> . <i>Pesticide Biochemistry and Physiology</i> , 2012, 104, 132-135.	1.6	25
141	Identification of Climatic Factors Affecting the Epidemiology of Human West Nile Virus Infections in Northern Greece. <i>PLoS ONE</i> , 2016, 11, e0161510.	1.1	24
142	Detection of West Nile Virus "Lineage 2 in <i>Culex pipiens</i> mosquitoes, associated with disease outbreak in Greece, 2017. <i>Acta Tropica</i> , 2018, 182, 64-68.	0.9	24
143	International workshop on insecticide resistance in vectors of arboviruses, December 2016, Rio de Janeiro, Brazil. <i>Parasites and Vectors</i> , 2017, 10, 278.	1.0	23
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153	Efficacy of ketoenols on insecticide resistant field populations of two-spotted spider mite <i>Tetranychus urticae</i> and sweet potato whitefly <i>Bemisia tabaci</i> from Greece. <i>Crop Protection</i> , 2012, 42, 305-311.	1.0	20
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159	A new dibenzoylhydrazine with insecticidal activity against <i>Anopheles</i> mosquito larvae. <i>Pest Management Science</i> , 2013, 69, 827-833.	1.7	18
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165	Development of efficient RNAi in <i>Nezara viridula</i> for use in insecticide target discovery. <i>Archives of Insect Biochemistry and Physiology</i> , 2020, 103, e21650.	0.6	17
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167	Susceptibility Profiles of <i>Helicoverpa armigera</i> (Lepidoptera: Noctuidae) to Deltamethrin Reveal a Contrast between the Northern and the Southern Benin. <i>International Journal of Environmental Research and Public Health</i> , 2019, 16, 1882.	1.2	16
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171	Pyrethroid and Etofenprox Resistance in <i>Anopheles gambiae</i> and <i>Anopheles coluzzii</i> from Vegetable Farms in Yaoundé, Cameroon: Dynamics, Intensity and Molecular Basis. <i>Molecules</i> , 2021, 26, 5543.	1.7	16
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