

Sulaiman S S Ibrahim

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28

papers

1,084

citations

16

h-index

32

g-index

35

ext. papers

1,434

ext. citations

5.3

avg, IF

4.15

L-index

#	Paper	IF	Citations
28	A single mutation in the GSTe2 gene allows tracking of metabolically based insecticide resistance in a major malaria vector. <i>Genome Biology</i> , 2014 , 15, R27	18.3	180
27	Directionally selected cytochrome P450 alleles are driving the spread of pyrethroid resistance in the major malaria vector <i>Anopheles funestus</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 252-7	11.5	147
26	Rise of multiple insecticide resistance in <i>Anopheles funestus</i> in Malawi: a major concern for malaria vector control. <i>Malaria Journal</i> , 2015 , 14, 344	3.6	71
25	Widespread pyrethroid and DDT resistance in the major malaria vector <i>Anopheles funestus</i> in East Africa is driven by metabolic resistance mechanisms. <i>PLoS ONE</i> , 2014 , 9, e110058	3.7	70
24	The highly polymorphic CYP6M7 cytochrome P450 gene partners with the directionally selected CYP6P9a and CYP6P9b genes to expand the pyrethroid resistance front in the malaria vector <i>Anopheles funestus</i> in Africa. <i>BMC Genomics</i> , 2014 , 15, 817	4.5	68
23	The cytochrome P450 CYP6P4 is responsible for the high pyrethroid resistance in knockdown resistance-free <i>Anopheles arabiensis</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2016 , 68, 23-32	4.5	61
22	Multiple Insecticide Resistance in the Malaria Vector <i>Anopheles funestus</i> from Northern Cameroon Is Mediated by Metabolic Resistance Alongside Potential Target Site Insensitivity Mutations. <i>PLoS ONE</i> , 2016 , 11, e0163261	3.7	59
21	A cytochrome P450 allele confers pyrethroid resistance on a major African malaria vector, reducing insecticide-treated bednet efficacy. <i>Science Translational Medicine</i> , 2019 , 11,	17.5	58
20	Allelic Variation of Cytochrome P450s Drives Resistance to Bednet Insecticides in a Major Malaria Vector. <i>PLoS Genetics</i> , 2015 , 11, e1005618	6	50
19	Multiple insecticide resistance in the major malaria vector <i>Anopheles funestus</i> in southern Ghana: implications for malaria control. <i>Parasites and Vectors</i> , 2016 , 9, 504	4	45
18	The P450 CYP6Z1 confers carbamate/pyrethroid cross-resistance in a major African malaria vector beside a novel carbamate-insensitive N485I acetylcholinesterase-1 mutation. <i>Molecular Ecology</i> , 2016 , 25, 3436-52	5.7	41
17	The Cytochrome P450 gene CYP6P12 confers pyrethroid resistance in kdr-free Malaysian populations of the dengue vector <i>Aedes albopictus</i> . <i>Scientific Reports</i> , 2016 , 6, 24707	4.9	40
16	Positional cloning of rp2 QTL associates the P450 genes CYP6Z1, CYP6Z3 and CYP6M7 with pyrethroid resistance in the malaria vector <i>Anopheles funestus</i> . <i>Heredity</i> , 2012 , 109, 383-92	3.6	38
15	High frequency of kdr L1014F is associated with pyrethroid resistance in <i>Anopheles coluzzii</i> in Sudan savannah of northern Nigeria. <i>BMC Infectious Diseases</i> , 2014 , 14, 441	4	36
14	Pyrethroid Resistance in Malaysian Populations of Dengue Vector <i>Aedes aegypti</i> Is Mediated by CYP9 Family of Cytochrome P450 Genes. <i>PLoS Neglected Tropical Diseases</i> , 2017 , 11, e0005302	4.8	26
13	Genome-Wide Transcription and Functional Analyses Reveal Heterogeneous Molecular Mechanisms Driving Pyrethroids Resistance in the Major Malaria Vector Across Africa. <i>G3: Genes, Genomes, Genetics</i> , 2017 , 7, 1819-1832	3.2	25
12	Temporal escalation of Pyrethroid Resistance in the major malaria vector <i>Anopheles coluzzii</i> from Sahelo-Sudanian Region of northern Nigeria. <i>Scientific Reports</i> , 2019 , 9, 7395	4.9	14

11	Pyrethroid Resistance in the Major Malaria Vector <i>Anopheles funestus</i> is Exacerbated by Overexpression and Overactivity of the P450 CYP6AA1 Across Africa. <i>Genes</i> , 2018 , 9,	4.2	12
10	A combination of metabolic resistance and high frequency of the 1014F kdr mutation is driving pyrethroid resistance in <i>Anopheles coluzzii</i> population from Guinea savanna of Cameroon. <i>Parasites and Vectors</i> , 2019 , 12, 263	4	11
9	Identification of Mutations in Antimalarial Resistance Gene from Isolates in Kano, Nigeria. <i>Tropical Medicine and Infectious Disease</i> , 2020 , 5,	3.5	6
8	Determination of Insecticide Susceptibility of Field Populations of Tomato Leaf Miner (<i>Tuta absoluta</i>) in Northern Nigeria. <i>Agriculture (Switzerland)</i> , 2019 , 9, 7	3	5
7	High Plasmodium infection and multiple insecticide resistance in a major malaria vector <i>Anopheles coluzzii</i> from Sahel of Niger Republic. <i>Malaria Journal</i> , 2019 , 18, 181	3.6	5
6	High insecticide resistance in the major malaria vector <i>Anopheles coluzzii</i> in Chad Republic. <i>Infectious Diseases of Poverty</i> , 2019 , 8, 100	10.4	5
5	Genome-Wide Transcriptional Analysis and Functional Validation Linked a Cluster of Epsilon Glutathione S-Transferases with Insecticide Resistance in the Major Malaria Vector across Africa. <i>Genes</i> , 2021 , 12,	4.2	4
4	Exploring the Mechanisms of Multiple Insecticide Resistance in a Highly -Infected Malaria Vector <i>Sensu Stricto</i> from Sahel of Northern Nigeria. <i>Genes</i> , 2020 , 11,	4.2	3
3	The cytochrome P450 CYP325A is a major driver of pyrethroid resistance in the major malaria vector <i>Anopheles funestus</i> in Central Africa. <i>Insect Biochemistry and Molecular Biology</i> , 2021 , 138, 103647-5	4.5	2
2	High pyrethroid/DDT resistance in major malaria vector <i>Anopheles coluzzii</i> from Niger-Delta of Nigeria is probably driven by metabolic resistance mechanisms. <i>PLoS ONE</i> , 2021 , 16, e0247944	3.7	1
1	Pyrethroid resistance in the New World malaria vector <i>Anopheles albimanus</i> is mediated by cytochrome P450 CYP6P5.. <i>Pesticide Biochemistry and Physiology</i> , 2022 , 183, 105061	4.9	