

William C Black

List of Publications by Year in descending order

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75
papers

4,530
citations

136740

32
h-index

118652

62
g-index

78
all docs

78
docs citations

78
times ranked

4628
citing authors

#	ARTICLE	IF	CITATIONS
1	Improved reference genome of <i>Aedes aegypti</i> informs arbovirus vector control. <i>Nature</i> , 2018, 563, 501-507.	13.7	426
2	Flavivirus Susceptibility in <i>Aedes aegypti</i> . <i>Archives of Medical Research</i> , 2002, 33, 379-388.	1.5	303
3	Genomic analysis of detoxification genes in the mosquito <i>Aedes aegypti</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2008, 38, 113-123.	1.2	289
4	Population Genetics of <i>Ixodes scapularis</i> (Acari: Ixodidae) Based on Mitochondrial 16S and 12S Genes. <i>Journal of Medical Entomology</i> , 1996, 33, 78-89.	0.9	231
5	Vector Competence of American Mosquitoes for Three Strains of Zika Virus. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0005101.	1.3	172
6	Quantitative Trait Loci That Control Vector Competence for Dengue-2 Virus in the Mosquito <i>Aedes aegypti</i> . <i>Genetics</i> , 2000, 156, 687-698.	1.2	166
7	Population genetics with RAPD-PCR markers: the breeding structure of <i>Aedes aegypti</i> in Puerto Rico. <i>Heredity</i> , 1996, 76, 325-334.	1.2	158
8	Does <i>kdr</i> genotype predict insecticide-resistance phenotype in mosquitoes?. <i>Trends in Parasitology</i> , 2009, 25, 213-219.	1.5	138
9	Recent Rapid Rise of a Permethrin Knock Down Resistance Allele in <i>Aedes aegypti</i> in México. <i>PLoS Neglected Tropical Diseases</i> , 2009, 3, e531.	1.3	130
10	Genetic Drift during Systemic Arbovirus Infection of Mosquito Vectors Leads to Decreased Relative Fitness during Host Switching. <i>Cell Host and Microbe</i> , 2016, 19, 481-492.	5.1	125
11	Spatial population genetic structure and limited dispersal in a Rocky Mountain alpine stream insect. <i>Molecular Ecology</i> , 2006, 15, 3553-3566.	2.0	124
12	RAPD-PCR and SSCP analysis for insect population genetic studies. , 1997, , 361-373.		116
13	Coevolution of the Ile1,016 and Cys1,534 Mutations in the Voltage Gated Sodium Channel Gene of <i>Aedes aegypti</i> in Mexico. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0004263.	1.3	116
14	Breeding structure of <i>Aedes aegypti</i> populations in Mexico varies by region.. <i>American Journal of Tropical Medicine and Hygiene</i> , 2002, 66, 213-222.	0.6	105
15	Population genomics reveals that an anthropophilic population of <i>Aedes aegypti</i> mosquitoes in West Africa recently gave rise to American and Asian populations of this major disease vector. <i>BMC Biology</i> , 2017, 15, 16.	1.7	96
16	Systematics and Biogeography of Hard Ticks, a Total Evidence Approach. <i>Cladistics</i> , 2000, 16, 79-102.	1.5	92
17	Parallel evolution of <i>vgsc</i> mutations at domains IS6, IIS6 and IIIS6 in pyrethroid resistant <i>Aedes aegypti</i> from Mexico. <i>Scientific Reports</i> , 2018, 8, 6747.	1.6	89
18	Intensive Linkage Mapping in a Wasp (<i>Bracon hebetor</i>) and a Mosquito (<i>Aedes aegypti</i>) With Single-Strand Conformation Polymorphism Analysis of Random Amplified Polymorphic DNA Markers. <i>Genetics</i> , 1996, 143, 1727-1738.	1.2	89

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19	Rapid and specific detection of Asian- and African-lineage Zika viruses. <i>Science Translational Medicine</i> , 2017, 9, .	5.8	86
20	Gene Flow, Subspecies Composition, and Dengue Virus-2 Susceptibility among <i>Aedes aegypti</i> Collections in Senegal. <i>PLoS Neglected Tropical Diseases</i> , 2009, 3, e408.	1.3	82
21	Quantitative Trait Loci Mapping of Genome Regions Controlling Permethrin Resistance in the Mosquito <i>Aedes aegypti</i> . <i>Genetics</i> , 2008, 180, 1137-1152.	1.2	75
22	Transcription of detoxification genes after permethrin selection in the mosquito <i>Aedes aegypti</i> . <i>Insect Molecular Biology</i> , 2012, 21, 61-77.	1.0	75
23	Why RIDL is not SIT. <i>Trends in Parasitology</i> , 2011, 27, 362-370.	1.5	71
24	Fitness Impact and Stability of a Transgene Conferring Resistance to Dengue-2 Virus following Introgression into a Genetically Diverse <i>Aedes aegypti</i> Strain. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e2833.	1.3	70
25	Characterization of the target of ivermectin, the glutamate-gated chloride channel, from <i>Anopheles gambiae</i> . <i>Journal of Experimental Biology</i> , 2015, 218, 1478-1486.	0.8	65
26	Taxonomic Status of <i>Ixodes neotomae</i> and <i>I. spinipalpis</i> (Acari: Ixodidae) Based on Mitochondrial DNA Evidence. <i>Journal of Medical Entomology</i> , 1997, 34, 696-703.	0.9	58
27	Vector Competence in West African <i>Aedes aegypti</i> Is Flavivirus Species and Genotype Dependent. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e3153.	1.3	56
28	Wide Spread Cross Resistance to Pyrethroids in <i>Aedes aegypti</i> (Diptera: Culicidae) From Veracruz State Mexico. <i>Journal of Economic Entomology</i> , 2013, 106, 959-969.	0.8	54
29	Experimental Evolution of an RNA Virus in Wild Birds: Evidence for Host-Dependent Impacts on Population Structure and Competitive Fitness. <i>PLoS Pathogens</i> , 2015, 11, e1004874.	2.1	51
30	Local Evolution of Pyrethroid Resistance Offsets Gene Flow Among <i>Aedes aegypti</i> Collections in Yucatan State, Mexico. <i>American Journal of Tropical Medicine and Hygiene</i> , 2015, 92, 201-209.	0.6	42
31	Age and prior blood feeding of <i>Anopheles gambiae</i> influences their susceptibility and gene expression patterns to ivermectin-containing blood meals. <i>BMC Genomics</i> , 2015, 16, 797.	1.2	39
32	Demonstration of efficient vertical and venereal transmission of dengue virus type-2 in a genetically diverse laboratory strain of <i>Aedes aegypti</i> . <i>PLoS Neglected Tropical Diseases</i> , 2018, 12, e0006754.	1.3	38
33	Insecticide resistance to permethrin and malathion and associated mechanisms in <i>Aedes aegypti</i> mosquitoes from St. Andrew Jamaica. <i>PLoS ONE</i> , 2017, 12, e0179673.	1.1	36
34	Variation in competence for ZIKV transmission by <i>Aedes aegypti</i> and <i>Aedes albopictus</i> in Mexico. <i>PLoS Neglected Tropical Diseases</i> , 2018, 12, e0006599.	1.3	36
35	Sequential Infection of <i>Aedes aegypti</i> Mosquitoes with Chikungunya Virus and Zika Virus Enhances Early Zika Virus Transmission. <i>Insects</i> , 2018, 9, 177.	1.0	34
36	The Neovolcanic Axis Is a Barrier to Gene Flow among <i>Aedes aegypti</i> Populations in Mexico That Differ in Vector Competence for Dengue 2 Virus. <i>PLoS Neglected Tropical Diseases</i> , 2009, 3, e468.	1.3	34

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37	First report of V1016G and S989P knockdown resistant (kdr) mutations in pyrethroid-resistant Sri Lankan <i>Aedes aegypti</i> mosquitoes. <i>Parasites and Vectors</i> , 2018, 11, 526.	1.0	33
38	Mosquitoes and West Nile virus along a river corridor from prairie to montane habitats in eastern Colorado. <i>Journal of Vector Ecology</i> , 2009, 34, 276-293.	0.5	30
39	Challenges in Estimating Insecticide Selection Pressures from Mosquito Field Data. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e1387.	1.3	27
40	QCal: A Software Application for the Calculation of Dose-Response Curves In Insecticide Resistance Bioassays. <i>Journal of the American Mosquito Control Association</i> , 2012, 28, 59-61.	0.2	26
41	Selection of D2S3, an <i>Aedes aegypti</i> (Diptera: Culicidae) Strain with High Oral Susceptibility to Dengue 2 Virus and D2MEB, a Strain with a Midgut Barrier to Dengue 2 Escape. <i>Journal of Medical Entomology</i> , 2005, 42, 110-119.	0.9	24
42	Linkage Analysis of Sex Determination in <i>Bracon</i> sp. <i>Near hebetor</i> (Hymenoptera: Braconidae). <i>Genetics</i> , 2000, 154, 205-212.	1.2	23
43	Assessing Insecticide Susceptibility of Laboratory <i>Lutzomyia longipalpis</i> and <i>Phlebotomus papatasi</i> Sand Flies (Diptera: Psychodidae: Phlebotominae). <i>Journal of Medical Entomology</i> , 2015, 52, 1003-1012.	0.9	22
44	Exon-Enriched Libraries Reveal Large Genic Differences Between <i>Aedes aegypti</i> from Senegal, West Africa, and Populations Outside Africa. <i>G3: Genes, Genomes, Genetics</i> , 2017, 7, 571-582.	0.8	22
45	The Genetic Basis for Salivary Gland Barriers to Arboviral Transmission. <i>Insects</i> , 2021, 12, 73.	1.0	22
46	A reverse-transcription/RNase H based protocol for depletion of mosquito ribosomal RNA facilitates viral intrahost evolution analysis, transcriptomics and pathogen discovery. <i>Virology</i> , 2019, 528, 181-197.	1.1	21
47	Reproductive Incompatibility Involving Senegalese <i>Aedes aegypti</i> (L) Is Associated with Chromosome Rearrangements. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004626.	1.3	21
48	Temporal and geographic evidence for evolution of Sin Nombre virus using molecular analyses of viral RNA from Colorado, New Mexico and Montana. <i>Virology Journal</i> , 2009, 6, 102.	1.4	19
49	Induction of RNA interference to block Zika virus replication and transmission in the mosquito <i>Aedes aegypti</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2019, 111, 103169.	1.2	19
50	Breeding structure of three snow pool <i>Aedes</i> mosquito species in northern Colorado. <i>Heredity</i> , 1998, 81, 371-380.	1.2	18
51	Differential Lymphocyte and Antibody Responses in Deer Mice Infected with Sin Nombre Hantavirus or Andes Hantavirus. <i>Journal of Virology</i> , 2014, 88, 8319-8331.	1.5	18
52	Susceptibility to Chlorpyrifos in Pyrethroid-Resistant Populations of <i>Aedes aegypti</i> (Diptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.9	16
53	Vgsc-interacting proteins are genetically associated with pyrethroid resistance in <i>Aedes aegypti</i> . <i>PLoS ONE</i> , 2019, 14, e0211497.	1.1	16
54	Resistance to commonly used insecticides and underlying mechanisms of resistance in <i>Aedes aegypti</i> (L.) from Sri Lanka. <i>Parasites and Vectors</i> , 2020, 13, 407.	1.0	15

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55	Permethrin resistance in <i>Aedes aegypti</i> : Genomic variants that confer knockdown resistance, recovery, and death. <i>PLoS Genetics</i> , 2021, 17, e1009606.	1.5	14
56	Loss of pyrethroid resistance in newly established laboratory colonies of <i>Aedes aegypti</i> . <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0007753.	1.3	13
57	<i>Aedes</i> species in treeholes and fruit husks between dry and wet seasons in southeastern Senegal. <i>Journal of Vector Ecology</i> , 2013, 38, 237-244.	0.5	12
58	From Global to Local—New Insights into Features of Pyrethroid Detoxification in Vector Mosquitoes. <i>Insects</i> , 2021, 12, 276.	1.0	12
59	Nootkatone Is an Effective Repellent against <i>Aedes aegypti</i> and <i>Aedes albopictus</i> . <i>Insects</i> , 2021, 12, 386.	1.0	12
60	The Yin and Yang of Linkage Disequilibrium: Mapping of Genes and Nucleotides Conferring Insecticide Resistance in Insect Disease Vectors. <i>Advances in Experimental Medicine and Biology</i> , 2008, 627, 71-83.	0.8	11
61	Insecticide resistance in <i>Aedes aegypti</i> from Tapachula, Mexico: Spatial variation and response to historical insecticide use. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009746.	1.3	10
62	Alternative patterns of sex chromosome differentiation in <i>Aedes aegypti</i> (L). <i>BMC Genomics</i> , 2017, 18, 943.	1.2	9
63	Learning to use <i>Ochlerotatus</i> is just the beginning. <i>Journal of the American Mosquito Control Association</i> , 2004, 20, 215-6.	0.2	8
64	<i>Aedes</i> (<i>Stegomyia</i>) <i>aegypti</i> and <i>Aedes</i> (<i>Howardina</i>) <i>cozumelensis</i> in Yucatán State, México, with a summary of published collection records for <i>Ae. cozumelensis</i> . <i>Journal of Vector Ecology</i> , 2012, 37, 365-372.	0.5	7
65	Profiles of Amino Acids and Acylcarnitines Related with Insecticide Exposure in <i>Culex quinquefasciatus</i> (Say). <i>PLoS ONE</i> , 2017, 12, e0169514.	1.1	7
66	A Linkage Map of the Asian Tiger Mosquito (<i>Aedes albopictus</i>) Based on cDNA Markers. <i>Journal of Heredity</i> , 2011, 102, 102-112.	1.0	6
67	QTL Mapping of Genome Regions Controlling Temephos Resistance in Larvae of the Mosquito <i>Aedes aegypti</i> . <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e3177.	1.3	6
68	Mitochondrial metabolic genes provide phylogeographic relationships of global collections of <i>Aedes aegypti</i> (Diptera: Culicidae). <i>PLoS ONE</i> , 2020, 15, e0235430.	1.1	5
69	Gene Flow Patterns among <i>Aedes aegypti</i> (Diptera: Culicidae) Populations in Sri Lanka. <i>Insects</i> , 2020, 11, 169.	1.0	4
70	Mosquitoes and West Nile Virus Along a River Corridor from Prairie to Montane Habitats in Eastern Colorado. <i>Journal of Vector Ecology</i> , 2009, 34, 276-293.	0.5	4
71	The Use of Insecticide-Treated Curtains for Control of <i>Aedes aegypti</i> and Dengue Virus Transmission in Fraccionamiento-Style Houses in México. <i>Journal of Tropical Medicine</i> , 2018, 2018, 1-22.	0.6	3
72	Permethrin Resistance Status and Associated Mechanisms in <i>Aedes albopictus</i> (Diptera: Culicidae) From Chiapas, Mexico. <i>Journal of Medical Entomology</i> , 2021, 58, 739-748.	0.9	3

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73	<i>Aedes aegypti</i> miRNA-33 modulates permethrin induced toxicity by regulating VGSC transcripts. <i>Scientific Reports</i> , 2021, 11, 7301.	1.6	3
74	Patterns of Variation in the Inhibitor of Apoptosis 1 Gene of <i>Aedes triseriatus</i> , a Transovarial Vector of La Crosse Virus. <i>Journal of Molecular Evolution</i> , 2009, 68, 403-413.	0.8	2
75	Effect of Selection for Pyrethroid Resistance on Abiotic Stress Tolerance in <i>Aedes aegypti</i> from Merida, Yucatan, Mexico. <i>Insects</i> , 2021, 12, 124.	1.0	2