

# Diego Ayala

## List of Publications by Year in descending order

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Version: 2024-02-01

53  
papers

2,688  
citations

218677

26  
h-index

223800

46  
g-index

61  
all docs

61  
docs citations

61  
times ranked

2992  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic diversity of the African malaria vector <i>Anopheles gambiae</i> . <i>Nature</i> , 2017, 552, 96-100.	27.8	288
2	Ecological niche partitioning between <i>Anopheles gambiae</i> molecular forms in Cameroon: the ecological side of speciation. <i>BMC Ecology</i> , 2009, 9, 17.	3.0	211
3	Global genetic diversity of <i>Aedes aegypti</i> . <i>Molecular Ecology</i> , 2016, 25, 5377-5395.	3.9	195
4	Living at the edge: biogeographic patterns of habitat segregation conform to speciation by niche expansion in <i>Anopheles gambiae</i> . <i>BMC Ecology</i> , 2009, 9, 16.	3.0	174
5	Carryover effects of larval exposure to different environmental bacteria drive adult trait variation in a mosquito vector. <i>Science Advances</i> , 2017, 3, e1700585.	10.3	172
6	Climate and Urbanization Drive Mosquito Preference for Humans. <i>Current Biology</i> , 2020, 30, 3570-3579.e6.	3.9	153
7	Habitat suitability and ecological niche profile of major malaria vectors in Cameroon. <i>Malaria Journal</i> , 2009, 8, 307.	2.3	84
8	REPRODUCTIVE ISOLATION AND LOCAL ADAPTATION QUANTIFIED FOR A CHROMOSOME INVERSION IN A MALARIA MOSQUITO. <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, 946-958.	2.3	84
9	Whole-genome sequencing reveals high complexity of copy number variation at insecticide resistance loci in malaria mosquitoes. <i>Genome Research</i> , 2019, 29, 1250-1261.	5.5	79
10	Adaptation through chromosomal inversions in <i>Anopheles</i> . <i>Frontiers in Genetics</i> , 2014, 5, 129.	2.3	75
11	Habitat segregation and ecological character displacement in cryptic African malaria mosquitoes. <i>Evolutionary Applications</i> , 2015, 8, 326-345.	3.1	75
12	Chromosomal Inversions, Natural Selection and Adaptation in the Malaria Vector <i>Anopheles funestus</i> . <i>Molecular Biology and Evolution</i> , 2011, 28, 745-758.	8.9	62
13	Ape malaria transmission and potential for ape-to-human transfers in Africa. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 5329-5334.	7.1	59
14	Population structure of a vector of human diseases: <i>Aedes aegypti</i> in its ancestral range, Africa. <i>Ecology and Evolution</i> , 2018, 8, 7835-7848.	1.9	57
15	A new species in the major malaria vector complex sheds light on reticulated species evolution. <i>Scientific Reports</i> , 2019, 9, 14753.	3.3	56
16	An <i>ace-1</i> gene duplication resorbs the fitness cost associated with resistance in <i>Anopheles gambiae</i> , the main malaria mosquito. <i>Scientific Reports</i> , 2015, 5, 14529.	3.3	52
17	Chromosomal and environmental determinants of morphometric variation in natural populations of the malaria vector <i>Anopheles funestus</i> in Cameroon. <i>Infection, Genetics and Evolution</i> , 2011, 11, 940-947.	2.3	51
18	Chromosome inversions and ecological plasticity in the main African malaria mosquitoes. <i>Evolution; International Journal of Organic Evolution</i> , 2017, 71, 686-701.	2.3	51

#	ARTICLE	IF	CITATIONS
19	Association mapping desiccation resistance within chromosomal inversions in the African malaria vector <i>Anopheles gambiae</i> . <i>Molecular Ecology</i> , 2019, 28, 1333-1342.	3.9	51
20	Microbial community structure reveals instability of nutritional symbiosis during the evolutionary radiation of <i>Amblyomma</i> ticks. <i>Molecular Ecology</i> , 2020, 29, 1016-1029.	3.9	48
21	Diurnal biting of malaria mosquitoes in the Central African Republic indicates residual transmission may be 'out of control'. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2104282119.	7.1	44
22	Dissecting functional components of reproductive isolation among closely related sympatric species of the <i>Anopheles gambiae</i> complex. <i>Evolutionary Applications</i> , 2017, 10, 1102-1120.	3.1	39
23	Population genomics in the arboviral vector <i>Aedes aegypti</i> reveals the genomic architecture and evolution of endogenous viral elements. <i>Molecular Ecology</i> , 2021, 30, 1594-1611.	3.9	37
24	Bat flies (Diptera: Nycteribiidae and Streblidae) infesting cave-dwelling bats in Gabon: diversity, dynamics and potential role in <i>Polychromophilus melanipherus</i> transmission. <i>Parasites and Vectors</i> , 2016, 9, 333.	2.5	36
25	Natural <i>Wolbachia</i> infections are common in the major malaria vectors in Central Africa. <i>Evolutionary Applications</i> , 2019, 12, 1583-1594.	3.1	36
26	Inversion breakpoints and the evolution of supergenes. <i>Molecular Ecology</i> , 2021, 30, 2738-2755.	3.9	36
27	Tracking zoonotic pathogens using blood-sucking flies as 'flying syringes'. <i>ELife</i> , 2017, 6, .	6.0	35
28	Population genetic structure of the malaria vector <i>Anopheles nili</i> in sub-Saharan Africa. <i>Malaria Journal</i> , 2010, 9, 161.	2.3	34
29	Advances and Perspectives in the Study of the Malaria Mosquito <i>Anopheles funestus</i> . , 0, , .		34
30	Population structure of the malaria vector <i>Anopheles funestus</i> (Diptera: Culicidae) in Madagascar and Comoros. <i>Acta Tropica</i> , 2006, 97, 292-300.	2.0	25
31	Discovery of Ongoing Selective Sweeps within <i>Anopheles</i> Mosquito Populations Using Deep Learning. <i>Molecular Biology and Evolution</i> , 2021, 38, 1168-1183.	8.9	25
32	In Silico Karyotyping of Chromosomally Polymorphic Malaria Mosquitoes in the <i>Anopheles gambiae</i> Complex. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 3249-3262.	1.8	24
33	Tolerance of disease-vector mosquitoes to brackish water and their osmoregulatory ability. <i>Ecosphere</i> , 2019, 10, e02783.	2.2	20
34	A targeted amplicon sequencing panel to simultaneously identify mosquito species and <i>Plasmodium</i> presence across the entire <i>Anopheles</i> genus. <i>Molecular Ecology Resources</i> , 2022, 22, 28-44.	4.8	18
35	Larval habitat determines the bacterial and fungal microbiota of the mosquito vector <i>Aedes aegypti</i> . <i>FEMS Microbiology Ecology</i> , 2022, 98, .	2.7	17
36	Population genetic structure of the malaria vector <i>Anopheles funestus</i> , in a recently re-colonized area of the Senegal River basin and human-induced environmental changes. <i>Parasites and Vectors</i> , 2012, 5, 188.	2.5	16

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37	Genetic structure of the mosquito <i>Aedes aegypti</i> in local forest and domestic habitats in Gabon and Kenya. <i>Parasites and Vectors</i> , 2020, 13, 417.	2.5	16
38	Larval sites of the mosquito <i>Aedes aegypti formosus</i> in forest and domestic habitats in Africa and the potential association with oviposition evolution. <i>Ecology and Evolution</i> , 2021, 11, 16327-16343.	1.9	16
39	Exome-wide association study reveals largely distinct gene sets underlying specific resistance to dengue virus types 1 and 3 in <i>Aedes aegypti</i> . <i>PLoS Genetics</i> , 2020, 16, e1008794.	3.5	13
40	Exploring the diversity of blood-sucking Diptera in caves of Central Africa. <i>Scientific Reports</i> , 2017, 7, 250.	3.3	12
41	“Show me which parasites you carry and I will tell you what you eat” or how to infer the trophic behavior of hematophagous arthropods feeding on wildlife. <i>Ecology and Evolution</i> , 2017, 7, 7578-7584.	1.9	12
42	Description of <i>Anopheles gabonensis</i> , a new species potentially involved in rodent malaria transmission in Gabon, Central Africa. <i>Infection, Genetics and Evolution</i> , 2014, 28, 628-634.	2.3	11
43	The origin of island populations of the African malaria mosquito, <i>Anopheles coluzzii</i> . <i>Communications Biology</i> , 2021, 4, 630.	4.4	7
44	Ecological plasticity to ions concentration determines genetic response and dominance of <i>Anopheles coluzzii</i> larvae in urban coastal habitats of Central Africa. <i>Scientific Reports</i> , 2021, 11, 15781.	3.3	7
45	Transposable element variants and their potential adaptive impact in urban populations of the malaria vector <i>Anopheles coluzzii</i> . <i>Genome Research</i> , 2022, 32, 189-202.	5.5	5
46	A molecular study of the genus <i>Spelaomyia</i> (Diptera: Phlebotominae) with description of the male of <i>Spelaomyia moucheti</i> . <i>Parasites and Vectors</i> , 2016, 9, 367.	2.5	3
47	Intraspecific Transcriptome Variation and Sex-Biased Expression in <i>Anopheles arabiensis</i> . <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	3
48	Experimental infections with Zika virus strains reveal high vector competence of <i>Aedes albopictus</i> and <i>Aedes aegypti</i> populations from Gabon (Central Africa) for the African virus lineage. <i>Emerging Microbes and Infections</i> , 2021, 10, 1244-1253.	6.5	1
49	Chapitre 10. Les anophèles (Diptera: Culicidae: Anophelinae)., 2017, , 181-241.		1
50	Title is missing!. , 2020, 16, e1008794.		0
51	Title is missing!. , 2020, 16, e1008794.		0
52	Title is missing!. , 2020, 16, e1008794.		0
53	Title is missing!. , 2020, 16, e1008794.		0