

# Nora J Besansky

## List of Publications by Year in descending order

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92  
papers

7,285  
citations

57758

44  
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64796

79  
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97  
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97  
docs citations

97  
times ranked

6099  
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	A PCR-RFLP method for genotyping of inversion 2Rc in <i>Anopheles coluzzii</i> . <i>Parasites and Vectors</i> , 2021, 14, 174.	2.5	3
3	Highly specific PCR-RFLP assays for karyotyping the widespread 2Rb inversion in malaria vectors of the <i>Anopheles gambiae</i> complex. <i>Parasites and Vectors</i> , 2020, 13, 16.	2.5	9
4	Evolutionary superscaffolding and chromosome anchoring to improve <i>Anopheles</i> genome assemblies. <i>BMC Biology</i> , 2020, 18, 1.	3.8	177
5	High-Throughput Genotyping of Common Chromosomal Inversions in the Afrotropical Malaria Mosquito <i>Anopheles Funestus</i> . <i>Insects</i> , 2020, 11, 693.	2.2	7
6	Assessing connectivity despite high diversity in island populations of a malaria mosquito. <i>Evolutionary Applications</i> , 2020, 13, 417-431.	3.1	11
7	Radiation with reticulation marks the origin of a major malaria vector. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 31583-31590.	7.1	29
8	Inversion Genotyping in the <i>Anopheles gambiae</i> Complex Using High-Throughput Array and Sequencing Platforms. <i>G3: Genes, Genomes, Genetics</i> , 2020, 10, 3299-3307.	1.8	8
9	Malaria mosquitoes go with the flow. <i>Nature</i> , 2019, 574, 340-341.	27.8	0
10	A chromosome-scale assembly of the major African malaria vector <i>Anopheles funestus</i> . <i>GigaScience</i> , 2019, 8, .	6.4	56
11	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
12	Fine-Mapping Complex Inversion Breakpoints and Investigating Somatic Pairing in the <i>Anopheles gambiae</i> Species Complex Using Proximity-Ligation Sequencing. <i>Genetics</i> , 2019, 213, 1495-1511.	2.9	27
13	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
14	Spatio-temporal genetic structure of <i>Anopheles gambiae</i> in the Northwestern Lake Victoria Basin, Uganda: implications for genetic control trials in malaria endemic regions. <i>Parasites and Vectors</i> , 2018, 11, 246.	2.5	11
15	Systems genetic analysis of inversion polymorphisms in the malaria mosquito <i>Anopheles gambiae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E7005-E7014.	7.1	47
16	Reduced-representation sequencing identifies small effective population sizes of <i>Anopheles gambiae</i> in the north-western Lake Victoria basin, Uganda. <i>Malaria Journal</i> , 2018, 17, 285.	2.3	7
17	Powerful methods for detecting introgressed regions from population genomic data. <i>Molecular Ecology</i> , 2016, 25, 2387-2397.	3.9	78
18	Radical remodeling of the Y chromosome in a recent radiation of malaria mosquitoes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E2114-23.	7.1	92

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19	How reticulated are species?. <i>BioEssays</i> , 2016, 38, 140-149.	2.5	449
20	Chromosomal inversions and ecotypic differentiation in <i>Anopheles gambiae</i> : the perspective from whole-genome sequencing. <i>Molecular Ecology</i> , 2016, 25, 5889-5906.	3.9	35
21	Transcriptomic differences between euryhaline and stenohaline malaria vector sibling species in response to salinity stress. <i>Molecular Ecology</i> , 2016, 25, 2210-2225.	3.9	17
22	Habitat segregation and ecological character displacement in cryptic African malaria mosquitoes. <i>Evolutionary Applications</i> , 2015, 8, 326-345.	3.1	75
23	Extensive introgression in a malaria vector species complex revealed by phylogenomics. <i>Science</i> , 2015, 347, 1258524.	12.6	527
24	Highly evolvable malaria vectors: The genomes of 16 <i>Anopheles</i> mosquitoes. <i>Science</i> , 2015, 347, 1258522.	12.6	492
25	How vector mosquitoes beat the heat. <i>Nature</i> , 2014, 516, 334-335.	27.8	5
26	Gene expression divergence between malaria vector sibling species <i>Anopheles gambiae</i> and <i>An. coluzzii</i> from rural and urban Yaoundé Cameroon. <i>Molecular Ecology</i> , 2014, 23, 2242-2259.	3.9	28
27	Cuticular differences associated with aridity acclimation in African malaria vectors carrying alternative arrangements of inversion 2La. <i>Parasites and Vectors</i> , 2014, 7, 176.	2.5	34
28	Mosquitoes. <i>Current Biology</i> , 2014, 24, R14-R15.	3.9	12
29	Physiological correlates of ecological divergence along an urbanization gradient: differential tolerance to ammonia among molecular forms of the malaria mosquito <i>Anopheles gambiae</i> . <i>BMC Ecology</i> , 2013, 13, 1.	3.0	67
30	Dose and developmental responses of <i>Anopheles merus</i> larvae to salinity. <i>Journal of Experimental Biology</i> , 2013, 216, 3433-3441.	1.7	32
31	The Evolution of the <i>Anopheles</i> 16 Genomes Project. <i>G3: Genes, Genomes, Genetics</i> , 2013, 3, 1191-1194.	1.8	49
32	<i>Anopheles coluzzii</i> and <i>Anopheles amharicus</i> , new members of the <i>Anopheles gambiae</i> complex. <i>Zootaxa</i> , 2013, 3619, .	0.5	411
33	<i>Anopheles coluzzii</i> and <i>Anopheles amharicus</i> , new members of the <i>Anopheles gambiae</i> complex. <i>Zootaxa</i> , 2013, 3619, 246-74.	0.5	272
34	Ecological Genomics of <i>Anopheles gambiae</i> Along a Latitudinal Cline: A Population-Resequencing Approach. <i>Genetics</i> , 2012, 190, 1417-1432.	2.9	157
35	Patterns of Genomic Differentiation between Ecologically Differentiated M and S Forms of <i>Anopheles gambiae</i> in West and Central Africa. <i>Genome Biology and Evolution</i> , 2012, 4, 1202-1212.	2.5	57
36	Spatially Explicit Analyses of Anopheline Mosquitoes Indoor Resting Density: Implications for Malaria Control. <i>PLoS ONE</i> , 2012, 7, e31843.	2.5	16

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37	Adaptation to Aridity in the Malaria Mosquito <i>Anopheles gambiae</i> : Chromosomal Inversion Polymorphism and Body Size Influence Resistance to Desiccation. <i>PLoS ONE</i> , 2012, 7, e34841.	2.5	80
38	Anthropogenic Habitat Disturbance and Ecological Divergence between Incipient Species of the Malaria Mosquito <i>Anopheles gambiae</i> . <i>PLoS ONE</i> , 2012, 7, e39453.	2.5	123
39	Evolution of <i>Anopheles gambiae</i> in Relation to Humans and Malaria. <i>Annual Review of Ecology, Evolution, and Systematics</i> , 2011, 42, 111-132.	8.3	87
40	Divergent transcriptional response to thermal stress by <i>Anopheles gambiae</i> larvae carrying alternative arrangements of inversion 2La. <i>Molecular Ecology</i> , 2011, 20, 2567-2580.	3.9	37
41	Adaptive divergence between incipient species of <i>Anopheles gambiae</i> increases resistance to <i>Plasmodium</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 244-249.	7.1	97
42	Evolutionary Dynamics of the Ty3/Gypsy LTR Retrotransposons in the Genome of <i>Anopheles gambiae</i> . <i>PLoS ONE</i> , 2011, 6, e16328.	2.5	15
43	The "Far-West" of <i>Anopheles gambiae</i> Molecular Forms. <i>PLoS ONE</i> , 2011, 6, e16415.	2.5	62
44	Genetic association of physically unlinked islands of genomic divergence in incipient species of <i>Anopheles gambiae</i> . <i>Molecular Ecology</i> , 2010, 19, 925-939.	3.9	123
45	A behavioral mechanism underlying ecological divergence in the malaria mosquito <i>Anopheles gambiae</i> . <i>Behavioral Ecology</i> , 2010, 21, 1087-1092.	2.2	76
46	Breakpoint structure of the <i>Anopheles gambiae</i> 2Rb chromosomal inversion. <i>Malaria Journal</i> , 2010, 9, 293.	2.3	40
47	Comparative Analysis of the Global Transcriptome of <i>Anopheles funestus</i> from Mali, West Africa. <i>PLoS ONE</i> , 2009, 4, e7976.	2.5	13
48	The Population Genomics of Trans-Specific Inversion Polymorphisms in <i>Anopheles gambiae</i> . <i>Genetics</i> , 2009, 183, 275-288.	2.9	47
49	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
50	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
51	Seasonal distribution of <i>Anopheles funestus</i> chromosomal forms from Burkina Faso. <i>Malaria Journal</i> , 2009, 8, 239.	2.3	24
52	Inversion 2La is associated with enhanced desiccation resistance in <i>Anopheles gambiae</i> . <i>Malaria Journal</i> , 2009, 8, 215.	2.3	77
53	2La chromosomal inversion enhances thermal tolerance of <i>Anopheles gambiae</i> larvae. <i>Malaria Journal</i> , 2009, 8, 147.	2.3	54
54	Differential gene expression in incipient species of <i>Anopheles gambiae</i> . <i>Molecular Ecology</i> , 2008, 17, 2491-2504.	3.9	46

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55	Chromosomal plasticity and evolutionary potential in the malaria vector <i>Anopheles gambiae</i> sensu stricto: insights from three decades of rare paracentric inversions. <i>BMC Evolutionary Biology</i> , 2008, 8, 309.	3.2	60
56	A test of the chromosomal theory of ecotypic speciation in <i>Anopheles gambiae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 2940-2945.	7.1	74
57	Localization of Candidate Regions Maintaining a Common Polymorphic Inversion (2La) in <i>Anopheles gambiae</i> . <i>PLoS Genetics</i> , 2007, 3, e217.	3.5	75
58	PCR-based karyotyping of <i>Anopheles gambiae</i> inversion 2Rj identifies the BAMAKO chromosomal form. <i>Malaria Journal</i> , 2007, 6, 133.	2.3	17
59	Segmental Duplication Implicated in the Genesis of Inversion 2Rj of <i>Anopheles gambiae</i> . <i>PLoS ONE</i> , 2007, 2, e849.	2.5	28
60	Polymorphism at the defensin gene in the <i>Anopheles gambiae</i> complex: Testing different selection hypotheses. <i>Infection, Genetics and Evolution</i> , 2007, 7, 285-292.	2.3	30
61	MOLECULAR KARYOTYPING OF THE 2LA INVERSION IN ANOPHELES GAMBIAE. <i>American Journal of Tropical Medicine and Hygiene</i> , 2007, 76, 334-339.	1.4	67
62	Molecular karyotyping of the 2La inversion in <i>Anopheles gambiae</i> . <i>American Journal of Tropical Medicine and Hygiene</i> , 2007, 76, 334-9.	1.4	48
63	Effective population size of <i>Anopheles funestus</i> chromosomal forms in Burkina Faso. <i>Malaria Journal</i> , 2006, 5, 115.	2.3	16
64	Analysis of the complete mitochondrial DNA from <i>Anopheles funestus</i> : An improved dipteran mitochondrial genome annotation and a temporal dimension of mosquito evolution. <i>Molecular Phylogenetics and Evolution</i> , 2006, 39, 417-423.	2.7	184
65	Gene Finding on the Y: Fruitful Strategy in <i>Drosophila</i> does not Deliver in <i>Anopheles</i> . <i>Genetica</i> , 2006, 126, 369-375.	1.1	9
66	Breakpoint structure reveals the unique origin of an interspecific chromosomal inversion (2La) in the <i>Anopheles gambiae</i> complex. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 6258-6262.	7.1	102
67	Divergence With Gene Flow in <i>Anopheles funestus</i> From the Sudan Savanna of Burkina Faso, West Africa. <i>Genetics</i> , 2006, 173, 1389-1395.	2.9	32
68	Variation in recombination rate across the X chromosome of <i>Anopheles gambiae</i> . <i>American Journal of Tropical Medicine and Hygiene</i> , 2006, 75, 901-3.	1.4	33
69	Sex-Linked Differentiation Between Incipient Species of <i>Anopheles gambiae</i> . <i>Genetics</i> , 2005, 169, 1509-1519.	2.9	50
70	An Integrated Genetic and Physical Map for the Malaria Vector <i>Anopheles funestus</i> . <i>Genetics</i> , 2005, 171, 1779-1787.	2.9	20
71	Satellite DNA From the Y Chromosome of the Malaria Vector <i>Anopheles gambiae</i> Sequence data from this article have been deposited with the EMBL/GenBank Data Libraries under accession nos. AY754141, AY754312. <i>Genetics</i> , 2005, 169, 185-196.	2.9	33
72	Centromere-proximal differentiation and speciation in <i>Anopheles gambiae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 15930-15935.	7.1	96

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73	Genetic structure of <i>Anopheles gambiae</i> populations on islands in northwestern Lake Victoria, Uganda. <i>Malaria Journal</i> , 2005, 4, 59.	2.3	21
74	Isolation and Characterization of Y Chromosome Sequences From the African Malaria Mosquito <i>Anopheles gambiae</i> . <i>Genetics</i> , 2004, 166, 1291-1302.	2.9	47
75	No accounting for taste: host preference in malaria vectors. <i>Trends in Parasitology</i> , 2004, 20, 249-251.	3.3	73
76	Dynamics of the pyrethroid knockdown resistance allele in western Kenyan populations of <i>Anopheles gambiae</i> in response to insecticide-treated bed net trials. <i>American Journal of Tropical Medicine and Hygiene</i> , 2004, 70, 591-6.	1.4	93
77	DNA barcoding of parasites and invertebrate disease vectors: what you don't know can hurt you. <i>Trends in Parasitology</i> , 2003, 19, 545-546.	3.3	118
78	Molecular Systematics of <i>Anopheles</i> : From Subgenera to Subpopulations. <i>Annual Review of Entomology</i> , 2003, 48, 111-139.	11.8	150
79	Frequent Intron Loss in the White Gene: A Cautionary Tale for Phylogeneticists. <i>Molecular Biology and Evolution</i> , 2002, 19, 362-366.	8.9	38
80	Inversions and Gene Order Shuffling in <i>Anopheles gambiae</i> and <i>A. funestus</i> . <i>Science</i> , 2002, 298, 182-185.	12.6	110
81	Structure and Evolution of mtanga, a Retrotransposon Actively Expressed on the Y Chromosome of the African Malaria Vector <i>Anopheles gambiae</i> . <i>Molecular Biology and Evolution</i> , 2002, 19, 149-162.	8.9	22
82	Bloodthirsty Hitchhikers?. <i>Science</i> , 2002, 295, 973-973.	12.6	0
83	Evolution of Mitochondrial and Ribosomal Gene Sequences in Anophelinae (Diptera: Culicidae): Implications for Phylogeny Reconstruction. <i>Molecular Phylogenetics and Evolution</i> , 2001, 18, 479-487.	2.7	80
84	The <i>Anopheles gambiae</i> tryptophan oxygenase gene expressed from a baculovirus promoter complements <i>Drosophila melanogaster</i> vermilion. <i>Insect Biochemistry and Molecular Biology</i> , 1997, 27, 803-805.	2.7	13
85	The <i>Anopheles albimanus</i> white gene: molecular characterization of the gene and a spontaneous white gene mutation. <i>Genetica</i> , 1997, 101, 87-96.	1.1	7
86	Patterns of Mitochondrial Variation Within and Between African Malaria Vectors, <i>Anopheles gambiae</i> and <i>An. arabiensis</i> , Suggest Extensive Gene Flow. <i>Genetics</i> , 1997, 147, 1817-1828.	2.9	119
87	An <i>Anopheles gambiae</i> cDNA predicts a protein similar to a yeast Suil translation factor. <i>Gene</i> , 1994, 141, 299-300.	2.2	5
88	Unintegrated Two-Long Terminal Repeat Circular Human T Lymphotropic Virus DNA Accumulation During Chronic HTLV Infection. <i>AIDS Research and Human Retroviruses</i> , 1993, 9, 1167-1172.	1.1	6
89	Molecular Perspectives on the Genetics of Mosquitoes. <i>Advances in Genetics</i> , 1992, 30, 123-184.	1.8	36
90	Reassociation Kinetics of <i>Anopheles gambiae</i> (Diptera: Culicidae) DNA. <i>Journal of Medical Entomology</i> , 1992, 29, 125-128.	1.8	63

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91	Extrachromosomal human immunodeficiency virus type-1 DNA can initiate a spreading infection of HL-60 cells. <i>Journal of Cellular Biochemistry</i> , 1991, 45, 366-373.	2.6	12
92	A Ribosomal RNA Gene Probe Differentiates Member Species of the <i>Anopheles gambiae</i> Complex. <i>American Journal of Tropical Medicine and Hygiene</i> , 1987, 37, 37-41.	1.4	508