

Tony Nolan

List of Publications by Year in descending order

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55
papers

5,034
citations

172457

29
h-index

161849

54
g-index

67
all docs

67
docs citations

67
times ranked

4163
citing authors

#	ARTICLE	IF	CITATIONS
1	A CRISPR-Cas9 gene drive system targeting female reproduction in the malaria mosquito vector <i>Anopheles gambiae</i> . <i>Nature Biotechnology</i> , 2016, 34, 78-83.	17.5	985
2	A CRISPR-Cas9 gene drive targeting doublesex causes complete population suppression in caged <i>Anopheles gambiae</i> mosquitoes. <i>Nature Biotechnology</i> , 2018, 36, 1062-1066.	17.5	648
3	Highly evolvable malaria vectors: The genomes of 16 <i>Anopheles</i> mosquitoes. <i>Science</i> , 2015, 347, 1258522.	12.6	492
4	Stable germline transformation of the malaria mosquito <i>Anopheles stephensi</i> . <i>Nature</i> , 2000, 405, 959-962.	27.8	344
5	The creation and selection of mutations resistant to a gene drive over multiple generations in the malaria mosquito. <i>PLoS Genetics</i> , 2017, 13, e1007039.	3.5	243
6	A comprehensive gene expression atlas of sex- and tissue-specificity in the malaria vector, <i>Anopheles gambiae</i> . <i>BMC Genomics</i> , 2011, 12, 296.	2.8	169
7	Bee Venom Phospholipase Inhibits Malaria Parasite Development in Transgenic Mosquitoes. <i>Journal of Biological Chemistry</i> , 2002, 277, 40839-40843.	3.4	168
8	A CRISPR-Cas9 sex-ratio distortion system for genetic control. <i>Scientific Reports</i> , 2016, 6, 31139.	3.3	160
9	A male-biased sex-distorter gene drive for the human malaria vector <i>Anopheles gambiae</i> . <i>Nature Biotechnology</i> , 2020, 38, 1054-1060.	17.5	153
10	The post-transcriptional gene silencing machinery functions independently of DNA methylation to repress a LINE1-like retrotransposon in <i>Neurospora crassa</i> . <i>Nucleic Acids Research</i> , 2005, 33, 1564-1573.	14.5	97
11	Tools for <i>Anopheles gambiae</i> Transgenesis. <i>G3: Genes, Genomes, Genetics</i> , 2015, 5, 1151-1163.	1.8	95
12	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
13	piggyBac-mediated Germline Transformation of the Malaria Mosquito <i>Anopheles stephensi</i> Using the Red Fluorescent Protein dsRED as a Selectable Marker. <i>Journal of Biological Chemistry</i> , 2002, 277, 8759-8762.	3.4	87
14	Transcription Regulation of Sex-Biased Genes during Ontogeny in the Malaria Vector <i>Anopheles gambiae</i> . <i>PLoS ONE</i> , 2011, 6, e21572.	2.5	82
15	Regulating the expression of gene drives is key to increasing their invasive potential and the mitigation of resistance. <i>PLoS Genetics</i> , 2021, 17, e1009321.	3.5	72
16	Phenylalanine Metabolism Regulates Reproduction and Parasite Melanization in the Malaria Mosquito. <i>PLoS ONE</i> , 2014, 9, e84865.	2.5	65
17	Requirements for Driving Antipathogen Effector Genes into Populations of Disease Vectors by Homing. <i>Genetics</i> , 2017, 205, 1587-1596.	2.9	62
18	Toward <i>Anopheles</i> transformation: Minos element activity in anopheline cells and embryos. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 2157-2162.	7.1	61

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19	Rapid evolution of female-biased genes among four species of <i>Anopheles</i> malaria mosquitoes. <i>Genome Research</i> , 2017, 27, 1536-1548.	5.5	60
20	Toward the Definition of Efficacy and Safety Criteria for Advancing Gene Drive-Modified Mosquitoes to Field Testing. <i>Vector-Borne and Zoonotic Diseases</i> , 2020, 20, 237-251.	1.5	60
21	Gene-drive suppression of mosquito populations in large cages as a bridge between lab and field. <i>Nature Communications</i> , 2021, 12, 4589.	12.8	59
22	Disruption of aminergic signalling reveals novel compounds with distinct inhibitory effects on mosquito reproduction, locomotor function and survival. <i>Scientific Reports</i> , 2014, 4, 5526.	3.3	49
23	Developing transgenic <i>Anopheles</i> mosquitoes for the sterile insect technique. <i>Genetica</i> , 2011, 139, 33-39.	1.1	44
24	The germline of the malaria mosquito produces abundant miRNAs, endo-siRNAs, piRNAs and 29-nt small RNAs. <i>BMC Genomics</i> , 2015, 16, 100.	2.8	44
25	Analysis of Two Novel Midgut-Specific Promoters Driving Transgene Expression in <i>Anopheles stephensi</i> Mosquitoes. <i>PLoS ONE</i> , 2011, 6, e16471.	2.5	40
26	Gene drive for population genetic control: non-functional resistance and parental effects. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2019, 286, 20191586.	2.6	39
27	Mosquito Transgenic Technologies to Reduce <i>Plasmodium</i> Transmission. <i>Methods in Molecular Biology</i> , 2012, 923, 601-622.	0.9	35
28	Control of malaria-transmitting mosquitoes using gene drives. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2021, 376, 20190803.	4.0	35
29	An <i>Anopheles gambiae</i> salivary gland promoter analysis in <i>Drosophila melanogaster</i> and <i>Anopheles stephensi</i> . <i>Insect Molecular Biology</i> , 2005, 14, 207-216.	2.0	34
30	Homology effects in <i>Neurospora crassa</i> . <i>FEMS Microbiology Letters</i> , 2006, 254, 182-189.	1.8	34
31	The RNA-dependent RNA polymerase essential for post-transcriptional gene silencing in <i>Neurospora crassa</i> interacts with replication protein A. <i>Nucleic Acids Research</i> , 2008, 36, 532-538.	14.5	32
32	Reprogramming homing endonuclease specificity through computational design and directed evolution. <i>Nucleic Acids Research</i> , 2014, 42, 2564-2576.	14.5	31
33	Deciphering the olfactory repertoire of the tiger mosquito <i>Aedes albopictus</i> . <i>BMC Genomics</i> , 2017, 18, 770.	2.8	30
34	A mosquito small RNA genomics resource reveals dynamic evolution and host responses to viruses and transposons. <i>Genome Research</i> , 2021, 31, 512-528.	5.5	29
35	The <i>Anopheles</i> FBN9 immune factor mediates <i>Plasmodium</i> species-specific defense through transgenic fat body expression. <i>Developmental and Comparative Immunology</i> , 2017, 67, 257-265.	2.3	28
36	CRISPR/Cas9 modified <i>An. gambiae</i> carrying <i>kdr</i> mutation L1014F functionally validate its contribution in insecticide resistance and combined effect with metabolic enzymes. <i>PLoS Genetics</i> , 2021, 17, e1009556.	3.5	27

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37	High-resolution transcriptional profiling of <i>Anopheles gambiae</i> spermatogenesis reveals mechanisms of sex chromosome regulation. <i>Scientific Reports</i> , 2019, 9, 14841.	3.3	26
38	Resistance to a CRISPR-based gene drive at an evolutionarily conserved site is revealed by mimicking genotype fixation. <i>PLoS Genetics</i> , 2021, 17, e1009740.	3.5	21
39	Cross-Species Y Chromosome Function Between Malaria Vectors of the <i>Anopheles gambiae</i> Species Complex. <i>Genetics</i> , 2017, 207, 729-740.	2.9	18
40	Post-integration behavior of a Minos transposon in the malaria mosquito <i>Anopheles stephensi</i> . <i>Molecular Genetics and Genomics</i> , 2007, 278, 575-584.	2.1	17
41	Nuclease-based gene drives, an innovative tool for insect vector control: advantages and challenges of the technology. <i>Current Opinion in Insect Science</i> , 2020, 39, 77-83.	4.4	17
42	Detecting the population dynamics of an autosomal sex ratio distorter transgene in malaria vector mosquitoes. <i>Journal of Applied Ecology</i> , 2020, 57, 2086-2096.	4.0	14
43	Transcript profiles of long- and short-lived adults implicate protein synthesis in evolved differences in ageing in the nematode <i>Strongyloides ratti</i> . <i>Mechanisms of Ageing and Development</i> , 2009, 130, 167-172.	4.6	12
44	Crystallographic analyses illustrate significant plasticity and efficient recoding of meganuclease target specificity. <i>Nucleic Acids Research</i> , 2017, 45, 8621-8634.	14.5	12
45	Introgression of a synthetic sex ratio distortion system from <i>Anopheles gambiae</i> into <i>Anopheles arabiensis</i> . <i>Scientific Reports</i> , 2019, 9, 5158.	3.3	11
46	Molecular tools and genetic markers for the generation of transgenic sexing strains in Anopheline mosquitoes. <i>Parasites and Vectors</i> , 2018, 11, 660.	2.5	10
47	CRISPR-mediated knock-in of transgenes into the malaria vector <i>Anopheles funestus</i> . <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	1.8	8
48	<i>Anopheles gambiae</i> Genome Conservation as a Resource for Rational Gene Drive Target Site Selection. <i>Insects</i> , 2021, 12, 97.	2.2	8
49	Transcriptional variation of sensory-related genes in natural populations of <i>Aedes albopictus</i> . <i>BMC Genomics</i> , 2020, 21, 547.	2.8	6
50	The long hand of the small RNAs reaches into several levels of gene regulation. <i>Biochemistry and Cell Biology</i> , 2004, 82, 472-481.	2.0	4
51	Identifying an essential interaction between malaria parasites and erythrocytes unlocks the door to promising vaccine targets. <i>Pathogens and Global Health</i> , 2012, 106, 64-64.	2.3	4
52	Ultra-conserved sequences in the genomes of highly diverse <i>Anopheles</i> mosquitoes, with implications for malaria vector control. <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	1.8	3
53	How to handle gene drives in arthropods?. <i>Pathogens and Global Health</i> , 2017, 111, 403-403.	2.3	0
54	Safe Driving: CRISPR and the Gene Drive Landscape. <i>CRISPR Journal</i> , 2018, 1, 16-18.	2.9	0

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55	Sex-, tissue- and stage-specific transgene expression.. , 2014, , 29-50.		0