

Omar Akbari

List of Publications by Citations

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119
papers

3,247
citations

31
h-index

54
g-index

131
ext. papers

4,383
ext. citations

8.8
avg, IF

5.85
L-index

| # | Paper | IF | Citations |
|-----|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 119 | Cheating evolution: engineering gene drives to manipulate the fate of wild populations. <i>Nature Reviews Genetics</i> , 2016 , 17, 146-59 | 30.1 | 283 |
| 118 | Improved reference genome of <i>Aedes aegypti</i> informs arbovirus vector control. <i>Nature</i> , 2018 , 563, 501-507 | 50.4 | 235 |
| 117 | BIOSAFETY. Safeguarding gene drive experiments in the laboratory. <i>Science</i> , 2015 , 349, 927-9 | 33.3 | 215 |
| 116 | Malaria eradication within a generation: ambitious, achievable, and necessary. <i>Lancet, The</i> , 2019 , 394, 1056-1112 | 40 | 130 |
| 115 | A synthetic gene drive system for local, reversible modification and suppression of insect populations. <i>Current Biology</i> , 2013 , 23, 671-7 | 6.3 | 127 |
| 114 | The developmental transcriptome of the mosquito <i>Aedes aegypti</i> , an invasive species and major arbovirus vector. <i>G3: Genes, Genomes, Genetics</i> , 2013 , 3, 1493-509 | 3.2 | 122 |
| 113 | Overcoming evolved resistance to population-suppressing homing-based gene drives. <i>Scientific Reports</i> , 2017 , 7, 3776 | 4.9 | 113 |
| 112 | Germline Cas9 expression yields highly efficient genome engineering in a major worldwide disease vector,. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017 , 114, E10540-E10549 | 11.5 | 105 |
| 111 | Mapping a multiplexed zoo of mRNA expression. <i>Development (Cambridge)</i> , 2016 , 143, 3632-3637 | 6.6 | 95 |
| 110 | Transforming insect population control with precision guided sterile males with demonstration in flies. <i>Nature Communications</i> , 2019 , 10, 84 | 17.4 | 85 |
| 109 | Development of a confinable gene drive system in the human disease vector. <i>ELife</i> , 2020 , 9, | 8.9 | 82 |
| 108 | Synthetically engineered gene drive system in the worldwide crop pest. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018 , 115, 4725-4730 | 11.5 | 73 |
| 107 | Novel synthetic <i>Medea</i> selfish genetic elements drive population replacement in <i>Drosophila</i> ; a theoretical exploration of <i>Medea</i> -dependent population suppression. <i>ACS Synthetic Biology</i> , 2014 , 3, 915-28 | 5.7 | 71 |
| 106 | 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 | 11.5 | 63 |
| 105 | Rules of the road for insect gene drive research and testing. <i>Nature Biotechnology</i> , 2017 , 35, 716-718 | 44.5 | 62 |
| 104 | Can CRISPR-Based Gene Drive Be Confined in the Wild? A Question for Molecular and Population Biology. <i>ACS Chemical Biology</i> , 2018 , 13, 424-430 | 4.9 | 58 |
| 103 | Engineered resistance to Zika virus in transgenic expressing a polycistronic cluster of synthetic small RNAs. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 3656-3661 | 11.5 | 53 |

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| 102 | Engineered Reciprocal Chromosome Translocations Drive High Threshold, Reversible Population Replacement in <i>Drosophila</i> . <i>ACS Synthetic Biology</i> , 2018 , 7, 1359-1370 | 5.7 | 53 |
| 101 | The olfactory basis of orchid pollination by mosquitoes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 708-716 | 11.5 | 49 |
| 100 | Generation of heritable germline mutations in the jewel wasp <i>Nasonia vitripennis</i> using CRISPR/Cas9. <i>Scientific Reports</i> , 2017 , 7, 901 | 4.9 | 47 |
| 99 | Modulation of Host Learning in <i>Aedes aegypti</i> Mosquitoes. <i>Current Biology</i> , 2018 , 28, 333-344.e8 | 6.3 | 47 |
| 98 | A novel promoter-tethering element regulates enhancer-driven gene expression at the bithorax complex in the <i>Drosophila</i> embryo. <i>Development (Cambridge)</i> , 2008 , 135, 123-31 | 6.6 | 44 |
| 97 | Opinion: Standardizing the definition of gene drive. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 30864-30867 | 11.5 | 38 |
| 96 | Assessment of a Split Homing Based Gene Drive for Efficient Knockout of Multiple Genes. <i>G3: Genes, Genomes, Genetics</i> , 2020 , 10, 827-837 | 3.2 | 38 |
| 95 | Gene editing technologies and applications for insects. <i>Current Opinion in Insect Science</i> , 2018 , 28, 66-72 | 5.1 | 38 |
| 94 | Visual-Olfactory Integration in the Human Disease Vector Mosquito <i>Aedes aegypti</i> . <i>Current Biology</i> , 2019 , 29, 2509-2516.e5 | 6.3 | 37 |
| 93 | Broad dengue neutralization in mosquitoes expressing an engineered antibody. <i>PLoS Pathogens</i> , 2020 , 16, e1008103 | 7.6 | 36 |
| 92 | Unraveling cis-regulatory mechanisms at the abdominal-A and Abdominal-B genes in the <i>Drosophila</i> bithorax complex. <i>Developmental Biology</i> , 2006 , 293, 294-304 | 3.1 | 36 |
| 91 | Transcriptome profiling of <i>Nasonia vitripennis</i> testis reveals novel transcripts expressed from the selfish B chromosome, paternal sex ratio. <i>G3: Genes, Genomes, Genetics</i> , 2013 , 3, 1597-605 | 3.2 | 35 |
| 90 | Core commitments for field trials of gene drive organisms. <i>Science</i> , 2020 , 370, 1417-1419 | 33.3 | 35 |
| 89 | Gene Drive Strategies for Population Replacement 2016 , 169-200 | | 33 |
| 88 | Progress towards engineering gene drives for population control. <i>Journal of Experimental Biology</i> , 2020 , 223, | 3 | 29 |
| 87 | Programmable RNA Targeting Using CasRx in Flies. <i>CRISPR Journal</i> , 2020 , 3, 164-176 | 2.5 | 26 |
| 86 | An Entry/Gateway cloning system for general expression of genes with molecular tags in <i>Drosophila melanogaster</i> . <i>BMC Cell Biology</i> , 2009 , 10, 8 | | 26 |
| 85 | Improved reference genome of the arboviral vector <i>Aedes albopictus</i> . <i>Genome Biology</i> , 2020 , 21, 215 | 18.3 | 26 |

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| 84 | Identification of germline transcriptional regulatory elements in <i>Aedes aegypti</i> . <i>Scientific Reports</i> , 2014 , 4, 3954 | 4.9 | 25 |
| 83 | Active Genetic Neutralizing Elements for Halting or Deleting Gene Drives. <i>Molecular Cell</i> , 2020 , 80, 246-262 | 6.4 | 25 |
| 82 | Inherently confinable split-drive systems in <i>Drosophila</i> . <i>Nature Communications</i> , 2021 , 12, 1480 | 17.4 | 24 |
| 81 | Winning the Tug-of-War Between Effector Gene Design and Pathogen Evolution in Vector Population Replacement Strategies. <i>Frontiers in Genetics</i> , 2019 , 10, 1072 | 4.5 | 24 |
| 80 | Genome elimination mediated by gene expression from a selfish chromosome. <i>Science Advances</i> , 2020 , 6, eaaz9808 | 14.3 | 24 |
| 79 | Highly Efficient Site-Specific Mutagenesis in Malaria Mosquitoes Using CRISPR. <i>G3: Genes, Genomes, Genetics</i> , 2018 , 8, 653-658 | 3.2 | 24 |
| 78 | Male-Killing Spiroplasma Alters Behavior of the Dosage Compensation Complex during <i>Drosophila melanogaster</i> Embryogenesis. <i>Current Biology</i> , 2016 , 26, 1339-45 | 6.3 | 18 |
| 77 | Live calcium imaging of <i>Aedes aegypti</i> neuronal tissues reveals differential importance of chemosensory systems for life-history-specific foraging strategies. <i>BMC Neuroscience</i> , 2019 , 20, 27 | 3.2 | 16 |
| 76 | Unique sequence organization and small RNA expression of a "selfish" B chromosome. <i>Chromosoma</i> , 2017 , 126, 753-768 | 2.8 | 16 |
| 75 | Germline mutagenesis of <i>Nasonia vitripennis</i> through ovarian delivery of CRISPR-Cas9 ribonucleoprotein. <i>Insect Molecular Biology</i> , 2020 , 29, 569-577 | 3.4 | 16 |
| 74 | The Developmental Transcriptome of , a Major Worldwide Human Disease Vector. <i>G3: Genes, Genomes, Genetics</i> , 2020 , 10, 1051-1062 | 3.2 | 16 |
| 73 | Modeling confinement and reversibility of threshold-dependent gene drive systems in spatially-explicit <i>Aedes aegypti</i> populations. <i>BMC Biology</i> , 2020 , 18, 50 | 7.3 | 15 |
| 72 | Methods for the generation of heritable germline mutations in the disease vector <i>Culex quinquefasciatus</i> using clustered regularly interspaced short palindrome repeats-associated protein 9. <i>Insect Molecular Biology</i> , 2020 , 29, 214-220 | 3.4 | 15 |
| 71 | A confinable home-and-rescue gene drive for population modification. <i>ELife</i> , 2021 , 10, | 8.9 | 15 |
| 70 | Sequence Expression of Supernumerary B Chromosomes: Function or Fluff?. <i>Genes</i> , 2019 , 10, | 4.2 | 14 |
| 69 | A typology of community and stakeholder engagement based on documented examples in the field of novel vector control. <i>PLoS Neglected Tropical Diseases</i> , 2019 , 13, e0007863 | 4.8 | 14 |
| 68 | Suppressing mosquito populations with precision guided sterile males. <i>Nature Communications</i> , 2021 , 12, 5374 | 17.4 | 14 |
| 67 | Identification of Genes Uniquely Expressed in the Germ-Line Tissues of the Jewel Wasp <i>Nasonia vitripennis</i> . <i>G3: Genes, Genomes, Genetics</i> , 2015 , 5, 2647-53 | 3.2 | 13 |

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| 66 | Vectored antibody gene delivery mediates long-term contraception. <i>Current Biology</i> , 2015 , 25, R820-2 | 6.3 | 12 |
| 65 | Site-specific transgenesis of the <i>Drosophila melanogaster</i> Y-chromosome using CRISPR/Cas9. <i>Insect Molecular Biology</i> , 2019 , 28, 65-73 | 3.4 | 12 |
| 64 | A drug-inducible sex-separation technique for insects. <i>Nature Communications</i> , 2020 , 11, 2106 | 17.4 | 11 |
| 63 | The abdominal-B promoter tethering element mediates promoter-enhancer specificity at the <i>Drosophila bithorax</i> complex. <i>Fly</i> , 2007 , 1, 337-9 | 1.3 | 11 |
| 62 | Sex ratio manipulation for insect population control. 2014 , 83-100 | | 11 |
| 61 | Combating mosquito-borne diseases using genetic control technologies. <i>Nature Communications</i> , 2021 , 12, 4388 | 17.4 | 11 |
| 60 | A mosquito small RNA genomics resource reveals dynamic evolution and host responses to viruses and transposons. <i>Genome Research</i> , 2021 , 31, 512-528 | 9.7 | 11 |
| 59 | Gene drives may be the next step towards sustainable control of malaria. <i>Pathogens and Global Health</i> , 2017 , 111, 399-400 | 3.1 | 10 |
| 58 | Germline Cas9 Expression Yields Highly Efficient Genome Engineering in a Major Worldwide Disease Vector, <i>Aedes aegypti</i> | | 10 |
| 57 | Improved <i>Aedes aegypti</i> mosquito reference genome assembly enables biological discovery and vector control | | 10 |
| 56 | Development of a Confinable Gene-Drive System in the Human Disease Vector, <i>Aedes aegypti</i> | | 9 |
| 55 | Engineered reciprocal chromosome translocations drive high threshold, reversible population replacement in <i>Drosophila</i> | | 8 |
| 54 | A home and rescue gene drive efficiently spreads and persists in populations | | 8 |
| 53 | Suppression of female fertility in with a CRISPR-targeted male-sterile mutation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118, | 11.5 | 8 |
| 52 | Broad Dengue Neutralization in Mosquitoes Expressing an Engineered Antibody. <i>SSRN Electronic Journal</i> , | 1 | 7 |
| 51 | A Sensitive, Rapid, and Portable CasRx-based Diagnostic Assay for SARS-CoV-2 2020 , | | 7 |
| 50 | Interdisciplinary development of a standardized introduction to gene drives for lay audiences. <i>BMC Medical Research Methodology</i> , 2020 , 20, 273 | 4.7 | 6 |
| 49 | Translating gene drive science to promote linguistic diversity in community and stakeholder engagement. <i>Global Public Health</i> , 2020 , 15, 1551-1565 | 3.5 | 6 |

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| 48 | Targeting female flight for genetic control of mosquitoes. <i>PLoS Neglected Tropical Diseases</i> , 2020 , 14, e0008876 | 4.8 | 6 |
| 47 | A CRISPR endonuclease gene drive reveals two distinct mechanisms of inheritance bias | | 6 |
| 46 | Embryo Microinjection and Transplantation Technique for <i>Nasonia vitripennis</i> Genome Manipulation. <i>Journal of Visualized Experiments</i> , 2017 , | 1.6 | 5 |
| 45 | Broad Dengue Neutralization in Mosquitoes Expressing an Engineered Antibody | | 5 |
| 44 | Reply to the Concerns about the feasibility of using "precision guided sterile males" to control insectsS <i>Nature Communications</i> , 2019 , 10, 3955 | 17.4 | 4 |
| 43 | Human attractive cues and mosquito host-seeking behavior. <i>Trends in Parasitology</i> , 2021 , | 6.4 | 4 |
| 42 | Genetically Encoded CRISPR Components Yield Efficient Gene Editing in the Invasive Pest. <i>CRISPR Journal</i> , 2021 , 4, 739-751 | 2.5 | 4 |
| 41 | Overcoming evolved resistance to population-suppressing homing-based gene drives | | 4 |
| 40 | Synthetically Engineered Medea Gene Drive System in the Worldwide Crop Pest, <i>D. suzukii</i> | | 4 |
| 39 | Can CRISPR-based gene drive be confined in the wild? A question for molecular and population biology | | 4 |
| 38 | Inherently confinable split-drive systems in <i>Drosophila</i> | | 4 |
| 37 | Engineered reproductively isolated species drive reversible population replacement. <i>Nature Communications</i> , 2021 , 12, 3281 | 17.4 | 4 |
| 36 | Embryo Microinjection Techniques for Efficient Site-Specific Mutagenesis in <i>Culex quinquefasciatus</i> . <i>Journal of Visualized Experiments</i> , 2020 , | 1.6 | 3 |
| 35 | Diverse Defenses: A Perspective Comparing Dipteran Piwi-piRNA Pathways. <i>Cells</i> , 2020 , 9, | 7.9 | 3 |
| 34 | Live calcium imaging of <i>Aedes aegypti</i> neuronal tissues reveals differential importance of chemosensory systems for life-history-specific foraging strategies | | 3 |
| 33 | Assessment of a split homing based gene drive for efficient knockout of multiple genes | | 3 |
| 32 | Synergistic Coding of Carbon Dioxide and a Human Sweat Odorant in the Mosquito Brain | | 2 |
| 31 | Modeling confinement and reversibility of threshold-dependent gene drive systems in spatially-explicit <i>Aedes aegypti</i> populations | | 2 |

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|----|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|---|
| 30 | Improved reference genome of the arboviral vector <i>Aedes albopictus</i> | | 2 |
| 29 | Development of a Rapid and Sensitive CasRx-Based Diagnostic Assay for SARS-CoV-2. <i>ACS Sensors</i> , 2021 , 6, 3957-3966 | 9.2 | 2 |
| 28 | Highly efficient site-specific mutagenesis in Malaria mosquitoes using CRISPR | | 2 |
| 27 | An integrated mosquito small RNA genomics resource reveals dynamic evolution and host responses to viruses and transposons | | 2 |
| 26 | Engineered Reproductively Isolated Species Drive Reversible Population Replacement | | 2 |
| 25 | Engineered resistance to Zika virus in transgenic <i>Ae. aegypti</i> expressing a polycistronic cluster of synthetic miRNAs | | 2 |
| 24 | The Developmental Transcriptome of <i>Ae. albopictus</i> , a Major Worldwide Human Disease Vector | | 2 |
| 23 | Eliminating Mosquitoes with Precision Guided Sterile Males | | 2 |
| 22 | Eliminating Mosquitoes with Precision Guided Sterile Males | | 2 |
| 21 | Parasitic nematode fatty acid- and retinol-binding proteins compromise host immunity by interfering with host lipid signaling pathways | | 2 |
| 20 | Oxitec and MosquitoMate in the United States: lessons for the future of gene drive mosquito control. <i>Pathogens and Global Health</i> , 2021 , 115, 365-376 | 3.1 | 2 |
| 19 | Transgenic refractory <i>Aedes aegypti</i> lines are resistant to multiple serotypes of dengue virus.. <i>Scientific Reports</i> , 2021 , 11, 23865 | 4.9 | 1 |
| 18 | Programmable RNA Targeting using CasRx in Flies | | 1 |
| 17 | Spatial control of gene expression in flies using bacterially derived binary transactivation systems | | 1 |
| 16 | Parasitic nematode fatty acid- and retinol-binding proteins compromise host immunity by interfering with host lipid signaling pathways. <i>PLoS Pathogens</i> , 2021 , 17, e1010027 | 7.6 | 1 |
| 15 | Germline mutagenesis of <i>Nasonia vitripennis</i> through ovarian delivery of CRISPR-Cas9 ribonucleoprotein | | 1 |
| 14 | Genome Elimination Mediated by Gene Expression from a Selfish Chromosome | | 1 |
| 13 | Mechanistically comparing reproductive manipulations caused by selfish chromosomes and bacterial symbionts. <i>Heredity</i> , 2021 , 126, 707-716 | 3.6 | 1 |

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| 12 | Spatial control of gene expression in flies using bacterially derived binary transactivation systems. <i>Insect Molecular Biology</i> , 2021 , 30, 461-471 | 3.4 | 1 |
| 11 | Temperature-Inducible Precision Guided Sterile Insect Technique | | 1 |
| 10 | Ubiquitous and Tissue-specific RNA Targeting in <i>Drosophila Melanogaster</i> using CRISPR/CasRx. <i>Journal of Visualized Experiments</i> , 2021 , | 1.6 | 1 |
| 9 | CRISPR Diagnostics: Advances toward the Point of Care.. <i>Biochemistry</i> , 2022 , | 3.2 | 1 |
| 8 | California Residents' Perceptions of Gene Drive Systems to Control Mosquito-Borne Disease.. <i>Frontiers in Bioengineering and Biotechnology</i> , 2022 , 10, 848707 | 5.8 | 1 |
| 7 | A day in the life of a mosquito insectary team: pushing for solutions to mosquito-borne diseases. <i>Lab Animal</i> , 2020 , 49, 241-243 | 0.4 | 0 |
| 6 | Exploiting a Y chromosome-linked Cas9 for sex selection and gene drive. <i>Nature Communications</i> , 2021 , 12, 7202 | 17.4 | 0 |
| 5 | CRISPR-Mediated Genome Engineering in <i>Aedes aegypti</i> . <i>Methods in Molecular Biology</i> , 2022 , 23-51 | 1.4 | 0 |
| 4 | Broad dengue neutralization in mosquitoes expressing an engineered antibody 2020 , 16, e1008103 | | |
| 3 | Broad dengue neutralization in mosquitoes expressing an engineered antibody 2020 , 16, e1008103 | | |
| 2 | Broad dengue neutralization in mosquitoes expressing an engineered antibody 2020 , 16, e1008103 | | |
| 1 | Broad dengue neutralization in mosquitoes expressing an engineered antibody 2020 , 16, e1008103 | | |