

Emily J Remnant

List of Publications by Year in descending order

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Version: 2024-02-01

31
papers

791
citations

623188

14
h-index

552369

26
g-index

33
all docs

33
docs citations

33
times ranked

904
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Cold case: The disappearance of Egypt bee virus, a fourth distinct master strain of deformed wing virus linked to honeybee mortality in 1970s Egypt. <i>Virology Journal</i> , 2022, 19, 12. | 1.4 | 17 |
| 2 | Abundant small RNAs in the reproductive tissues and eggs of the honey bee, <i>Apis mellifera</i> . <i>BMC Genomics</i> , 2022, 23, 257. | 1.2 | 6 |
| 3 | Viral communities in the parasite <i>Varroa destructor</i> and in colonies of their honey bee host (<i>Apis mellifera</i>). <i>Journal of Animal Ecology</i> , 2021, 90, 2254-2267. | 1.6 | 12 |
| 4 | Adaptation to vector-based transmission in a honeybee virus. <i>Journal of Animal Ecology</i> , 2021, 90, 2254-2267. | 1.3 | 20 |
| 5 | A Diverse Viral Community from Predatory Wasps in Their Native and Invaded Range, with a New Virus Infectious to Honey Bees. <i>Viruses</i> , 2021, 13, 1431. | 1.5 | 10 |
| 6 | Reply to Soley: DNA methylation marks are stably transferred across generations in honey bees. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, . | 3.3 | 2 |
| 7 | DNA methylation is not a driver of gene expression reprogramming in young honey bee workers. <i>Molecular Ecology</i> , 2021, 30, 4804-4818. | 2.0 | 21 |
| 8 | High-Quality Assemblies for Three Invasive Social Wasps from the <i>Vespula</i> Genus. <i>G3: Genes, Genomes, Genetics</i> , 2020, 10, 3479-3488. | 0.8 | 19 |
| 9 | A Single Gene Causes Thelytokous Parthenogenesis, the Defining Feature of the Cape Honeybee <i>Apis mellifera capensis</i> . <i>Current Biology</i> , 2020, 30, 2248-2259.e6. | 1.8 | 23 |
| 10 | Accumulation and Competition Amongst Deformed Wing Virus Genotypes in Naïve Australian Honeybees Provides Insight Into the Increasing Global Prevalence of Genotype B. <i>Frontiers in Microbiology</i> , 2020, 11, 620. | 1.5 | 32 |
| 11 | Paternally-biased gene expression follows kin-selected predictions in female honey bee embryos. <i>Molecular Ecology</i> , 2020, 29, 1523-1533. | 2.0 | 16 |
| 12 | Intergenerational transfer of DNA methylation marks in the honey bee. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 32519-32527. | 3.3 | 45 |
| 13 | Unique DNA Methylation Profiles Are Associated with cis-Variation in Honey Bees. <i>Genome Biology and Evolution</i> , 2019, 11, 2517-2530. | 1.1 | 31 |
| 14 | The frequency of thelytokous parthenogenesis in European-derived <i>Apis mellifera</i> virgin queens. <i>Apidologie</i> , 2019, 50, 295-303. | 0.9 | 5 |
| 15 | Direct transmission by injection affects competition among RNA viruses in honeybees. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2019, 286, 20182452. | 1.2 | 37 |
| 16 | Genetic origins of honey bees (<i>Apis mellifera</i>) on Kangaroo Island and Norfolk Island (Australia) and the Kingdom of Tonga. <i>Apidologie</i> , 2019, 50, 28-39. | 0.9 | 5 |
| 17 | Viable Triploid Honey Bees (<i>Apis mellifera capensis</i>) Are Reliably Produced in the Progeny of CO ₂ Narcotised Queens. <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 3357-3366. | 0.8 | 5 |
| 18 | Honey Bees, Royal Jelly, Epigenetics. , 2018, , 722-727. | | 0 |

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|----|--|-----|-----------|
| 19 | A Diverse Range of Novel RNA Viruses in Geographically Distinct Honey Bee Populations. <i>Journal of Virology</i> , 2017, 91, . | 1.5 | 138 |
| 20 | Paternal effects on <i>Apis mellifera capensis</i> worker ovary size. <i>Apidologie</i> , 2017, 48, 660-665. | 0.9 | 7 |
| 21 | DNA methylation of Kr-h1 is involved in regulating ovary activation in worker honeybees (<i>Apis</i>) <i>Tj ETQq1 1 0.784314 rgBT /Overlock 1</i> | 0.7 | 3 |
| 22 | No evidence that DNA methylation is associated with the regulation of fertility in the adult honey bee <i>Apis mellifera</i> (Hymenoptera: Apidae) worker ovary. <i>Austral Entomology</i> , 2017, 56, 115-121. | 0.8 | 1 |
| 23 | Nutrition and Epigenetic Change in Insects: Evidence and Implications. <i>Advances in Insect Physiology</i> , 2017, 53, 31-54. | 1.1 | 4 |
| 24 | Evolution, Expression, and Function of Nonneuronal Ligand-Gated Chloride Channels in <i>Drosophila melanogaster</i> . <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 2003-2012. | 0.8 | 13 |
| 25 | Parent-of-origin effects on genome-wide DNA methylation in the Cape honey bee (<i>Apis mellifera</i>) <i>Tj ETQq1 1 0.784314 rgBT /Overlock 1</i> | 1.2 | 54 |
| 26 | Reproductive interference between honeybee species in artificial sympatry. <i>Molecular Ecology</i> , 2014, 23, 1096-1107. | 2.0 | 20 |
| 27 | Whole-Genome DNA Methylation Profile of the Jewel Wasp (<i>Nasonia vitripennis</i>). <i>G3: Genes, Genomes, Genetics</i> , 2014, 4, 383-388. | 0.8 | 59 |
| 28 | A parent-of-origin effect on honeybee worker ovary size. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2014, 281, 20132388. | 1.2 | 34 |
| 29 | The role of Rdl in resistance to phenylpyrazoles in <i>Drosophila melanogaster</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2014, 54, 11-21. | 1.2 | 30 |
| 30 | The dynamic DNA methylation cycle from egg to sperm in the honey bee <i>Apis mellifera</i> . <i>Development (Cambridge)</i> , 2014, 141, 2702-2711. | 1.2 | 58 |
| 31 | Gene duplication in the major insecticide target site, <i>Rdl</i> , in <i>Drosophila melanogaster</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 14705-14710. | 3.3 | 63 |