

# 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	A Diverse Range of Novel RNA Viruses in Geographically Distinct Honey Bee Populations. <i>Journal of Virology</i> , 2017, 91, .	1.5	138
2	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
3	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
4	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
5	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 11</i>	1.2	54
6	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
7	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
8	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
9	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
10	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
11	The role of <i>Rdl</i> in resistance to phenylpyrazoles in <i>Drosophila melanogaster</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2014, 54, 11-21.	1.2	30
12	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
13	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
14	Reproductive interference between honeybee species in artificial sympatry. <i>Molecular Ecology</i> , 2014, 23, 1096-1107.	2.0	20
15	Adaptation to vector-based transmission in a honeybee virus. <i>Journal of Animal Ecology</i> , 2021, 90, 2254-2267.	1.3	20
16	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
17	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
18	Paternally-biased gene expression follows kin-selected predictions in female honey bee embryos. <i>Molecular Ecology</i> , 2020, 29, 1523-1533.	2.0	16

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19	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
20	Viral communities in the parasite <i>Varroa destructor</i> and in colonies of their honey bee host ( <i>Apis mellifera</i> ). <i>Overlook</i> , 2010, 10, 50-71.	1.6	12
21	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
22	Paternal effects on <i>Apis mellifera capensis</i> worker ovary size. <i>Apidologie</i> , 2017, 48, 660-665.	0.9	7
23	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
24	Viable Triploid Honey Bees ( <i>Apis mellifera capensis</i> ) Are Reliably Produced in the Progeny of CO <sub>2</sub> Narcotised Queens. <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 3357-3366.	0.8	5
25	The frequency of thelytokous parthenogenesis in European-derived <i>Apis mellifera</i> virgin queens. <i>Apidologie</i> , 2019, 50, 295-303.	0.9	5
26	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
27	Nutrition and Epigenetic Change in Insects: Evidence and Implications. <i>Advances in Insect Physiology</i> , 2017, 53, 31-54.	1.1	4
28	DNA methylation of Kr-h1 is involved in regulating ovary activation in worker honeybees ( <i>Apis mellifera</i> ). <i>Overlook</i> , 2010, 10, 382-391.	0.7	3
29	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
30	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
31	Honey Bees, Royal Jelly, Epigenetics. , 2018, , 722-727.		0