Gabriele Buchmann

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Loss of mitochondrial diversity in invasive populations of Asian honey bees, <scp><i>Apis cerana</i></scp> (Hymenoptera: Apidae), in the Australâ€Pacific. Austral Entomology, 2022, 61, 97-103.	0.8	3
2	Split or combine? Effects of repeated sampling and data pooling on the estimation of colony numbers obtained from drone genotyping. Apidologie, 2021, 52, 620-631.	0.9	2
3	Adaptation to vectorâ€based transmission in a honeybee virus. Journal of Animal Ecology, 2021, 90, 2254-2267.	1.3	20
4	Reply to Soley: DNA methylation marks are stably transferred across generations in honey bees. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	2
5	A Single Gene Causes Thelytokous Parthenogenesis, the Defining Feature of the Cape Honeybee Apis mellifera capensis. Current Biology, 2020, 30, 2248-2259.e6.	1.8	23
6	Cross-Kingdom RNAi of Pathogen Effectors Leads to Quantitative Adult Plant Resistance in Wheat. Frontiers in Plant Science, 2020, 11, 253.	1.7	24
7	What mechanistic factors affect thelytokous parthenogenesis in Apis mellifera caponises queens?. Apidologie, 2020, 51, 329-341.	0.9	1
8	Accumulation and Competition Amongst Deformed Wing Virus Genotypes in NaÃ ⁻ ve Australian Honeybees Provides Insight Into the Increasing Global Prevalence of Genotype B. Frontiers in Microbiology, 2020, 11, 620.	1.5	32
9	Paternallyâ€biased gene expression follows kinâ€selected predictions in female honey bee embryos. Molecular Ecology, 2020, 29, 1523-1533.	2.0	16
10	Intergenerational transfer of DNA methylation marks in the honey bee. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 32519-32527.	3.3	45
11	Unique DNA Methylation Profiles Are Associated with cis-Variation in Honey Bees. Genome Biology and Evolution, 2019, 11, 2517-2530.	1.1	31
12	Workers' sons rescue genetic diversity at the sex locus in an invasive honey bee population. Molecular Ecology, 2019, 28, 1585-1592.	2.0	15
13	Genetic Diversity in the Progeny of Commercial Australian Queen Honey Bees (Hymenoptera: Apidae) Produced in Autumn and Early Spring. Journal of Economic Entomology, 2019, 112, 33-39.	0.8	5
14	Viable Triploid Honey Bees (Apis mellifera capensis) Are Reliably Produced in the Progeny of CO2 Narcotised Queens. G3: Genes, Genomes, Genetics, 2018, 8, 3357-3366.	0.8	5
15	Chromatin Modifiers SET-25 and SET-32 Are Required for Establishment but Not Long-Term Maintenance of Transgenerational Epigenetic Inheritance. Cell Reports, 2018, 25, 2259-2272.e5.	2.9	50
16	Evolutionary divergence of the rye Pm17 and Pm8 resistance genes reveals ancient diversity. Plant Molecular Biology, 2018, 98, 249-260.	2.0	75
17	A Diverse Range of Novel RNA Viruses in Geographically Distinct Honey Bee Populations. Journal of Virology, 2017, 91,	1.5	138
18	An invasive social insect overcomes genetic load at the sex locus. Nature Ecology and Evolution, 2017, 1, 11.	3.4	45

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19	Parent-of-origin effects on genome-wide DNA methylation in the Cape honey bee (Apis mellifera) Tj ETQq1 1 0.784	4314 rgBT 1.2	/Overlock 1
20	The wheat resistance gene <i>Lr34</i> results in the constitutive induction of multiple defense pathways in transgenic barley. Plant Journal, 2015, 84, 202-215.	2.8	45
21	Genetic and molecular characterization of a locus involved in avirulence of Blumeria graminis f. sp. tritici on wheat Pm3 resistance alleles. Fungal Genetics and Biology, 2015, 82, 181-192.	0.9	50
22	Rye <i><scp>P</scp>m8</i> and wheat <i><scp>P</scp>m3</i> are orthologous genes and show evolutionary conservation of resistance function against powdery mildew. Plant Journal, 2013, 76, 957-969.	2.8	178