

# James G Baldwin-Brown

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3589276/publications.pdf>

Version: 2024-02-01

8  
papers

527  
citations

1478505

6  
h-index

1588992

8  
g-index

11  
all docs

11  
docs citations

11  
times ranked

1153  
citing authors

#	ARTICLE	IF	CITATIONS
1	The assembled and annotated genome of the pigeon louse <i>Columbicola columbae</i> , a model ectoparasite. <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	1.8	18
2	Genomic Signatures of Local Adaptation in Clam Shrimp ( <i>Eulimnadia texana</i> ) from Natural Vernal Pools. <i>Genome Biology and Evolution</i> , 2020, 12, 1194-1206.	2.5	1
3	The genome of <i>Peromyscus leucopus</i> , natural host for Lyme disease and other emerging infections. <i>Science Advances</i> , 2019, 5, eaaw6441.	10.3	41
4	The gene structure and hypervariability of the complete <i>Penaeus monodon</i> Dscam gene. <i>Scientific Reports</i> , 2019, 9, 16595.	3.3	8
5	Genomes, expression profiles, and diversity of mitochondria of the White-footed Deermouse <i>Peromyscus leucopus</i> , reservoir of Lyme disease and other zoonoses. <i>Scientific Reports</i> , 2019, 9, 17618.	3.3	9
6	A New Standard for Crustacean Genomes: The Highly Contiguous, Annotated Genome Assembly of the Clam Shrimp <i>Eulimnadia texana</i> Reveals HOX Gene Order and Identifies the Sex Chromosome. <i>Genome Biology and Evolution</i> , 2018, 10, 143-156.	2.5	33
7	Contiguous and accurate <i>de novo</i> assembly of metazoan genomes with modest long read coverage. <i>Nucleic Acids Research</i> , 2016, 44, gkw654.	14.5	329
8	The Power to Detect Quantitative Trait Loci Using Resequenced, Experimentally Evolved Populations of Diploid, Sexual Organisms. <i>Molecular Biology and Evolution</i> , 2014, 31, 1040-1055.	8.9	88