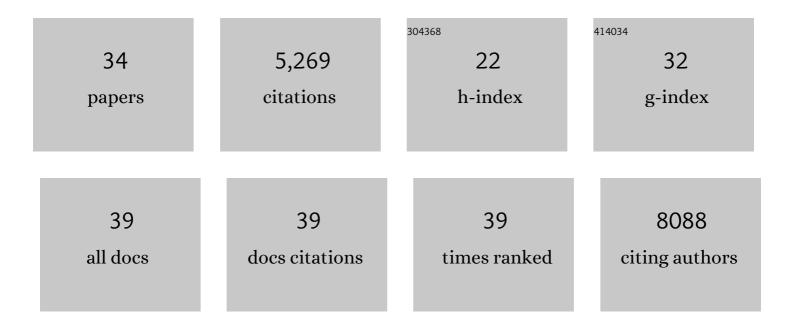
Floyd A Reed

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9433018/publications.pdf Version: 2024-02-01



FLOVD A REED

#	Article	IF	CITATIONS
1	Wild gut microbiomes reveal individuals, species, and location as drivers of variation in two critically endangered Hawaiian honeycreepers. PeerJ, 2021, 9, e12291.	0.9	7
2	Linking genotype to phenotype in a changing ocean: inferring the genomic architecture of a blue mussel stress response with genomeâ€wide association. Journal of Evolutionary Biology, 2018, 31, 346-361.	0.8	6
3	Gene expression across tissues, sex, and life stages in the sea urchin Tripneustes gratilla [Toxopneustidae, Odontophora, Camarodonta]. Marine Genomics, 2018, 41, 12-18.	0.4	4
4	CRISPR/Cas9 Gene Drive: Growing Pains for a New Technology. Genetics, 2017, 205, 1037-1039.	1.2	14
5	Transcriptional effects of a positive feedback circuit in Drosophila melanogaster. BMC Genomics, 2017, 18, 990.	1.2	5
6	Stability of underdominant genetic polymorphisms in population networks. Journal of Theoretical Biology, 2016, 390, 156-163.	0.8	18
7	Reply to Garner et al Trends in Ecology and Evolution, 2016, 31, 83-84.	4.2	24
8	Genomics and the challenging translation into conservation practice. Trends in Ecology and Evolution, 2015, 30, 78-87.	4.2	469
9	First Steps towards Underdominant Genetic Transformation of Insect Populations. PLoS ONE, 2014, 9, e97557.	1.1	81
10	Dynamics of a combined medea-underdominant population transformation system. BMC Evolutionary Biology, 2014, 14, 98.	3.2	24
11	Scientific Standards and the Regulation of Genetically Modified Insects. PLoS Neglected Tropical Diseases, 2012, 6, e1502.	1.3	60
12	Modern Human Migrations: The First 200,000 Years. , 2012, , 315-326.		0
13	From genes to games: Cooperation and cyclic dominance in meiotic drive. Journal of Theoretical Biology, 2012, 299, 120-125.	0.8	42
14	Rapid formation of distinct hybrid lineages after secondary contact of two fish species (<i>Cottus</i>) Tj ETQq0	0 0 rgBT / 2.0	Overlock 101
15	Alignment-free estimation of nucleotide diversity. Bioinformatics, 2011, 27, 449-455.	1.8	20
16	Stability Properties of Underdominance in Finite Subdivided Populations. PLoS Computational Biology, 2011, 7, e1002260.	1.5	61
17	Using underdominance to bi-stably transform local populations. Journal of Theoretical Biology, 2010, 267, 62-75.	0.8	64
18	The Genetic Structure and History of Africans and African Americans. Science, 2009, 324, 1035-1044.	6.0	1,267

2

FLOYD A REED

#	Article	IF	CITATIONS
19	The collective-risk social dilemma and the prevention of simulated dangerous climate change. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 2291-2294.	3.3	429
20	The Genetic Structure of Pacific Islanders. PLoS Genetics, 2008, 4, e19.	1.5	251
21	Two-Locus Epistasis With Sexually Antagonistic Selection: A Genetic Parrondo's Paradox. Genetics, 2007, 176, 1923-1929.	1.2	53
22	History of Click-Speaking Populations of Africa Inferred from mtDNA and Y Chromosome Genetic Variation. Molecular Biology and Evolution, 2007, 24, 2180-2195.	3.5	202
23	Whole-mtDNA Genome Sequence Analysis of Ancient African Lineages. Molecular Biology and Evolution, 2007, 24, 757-768.	3.5	234
24	Convergent adaptation of human lactase persistence in Africa and Europe. Nature Genetics, 2007, 39, 31-40.	9.4	1,375
25	African human diversity, origins and migrations. Current Opinion in Genetics and Development, 2006, 16, 597-605.	1.5	98
26	Evaluation of Real-Time PCR Amplification Efficiencies to Detect PCR Inhibitors. Journal of Forensic Sciences, 2006, 51, 795-804.	0.9	136
27	Mutation, selection and the future of human evolution. Trends in Genetics, 2006, 22, 479-484.	2.9	29
28	Positive Selection Can Create False Hotspots of Recombination. Genetics, 2006, 172, 2011-2014.	1.2	48
29	EVIDENCE OF SUSCEPTIBILITY AND RESISTANCE TO CRYPTIC X-LINKED MEIOTIC DRIVE IN NATURAL POPULATIONS OF DROSOPHILA MELANOGASTER. Evolution; International Journal of Organic Evolution, 2005, 59, 1280-1291.	1.1	13
30	EVIDENCE OF SUSCEPTIBILITY AND RESISTANCE TO CRYPTIC X-LINKED MEIOTIC DRIVE IN NATURAL POPULATIONS OF DROSOPHILA MELANOGASTER. Evolution; International Journal of Organic Evolution, 2005, 59, 1280.	1.1	0
31	Fitting background-selection predictions to levels of nucleotide variation and divergence along the human autosomes. Genome Research, 2005, 15, 1211-1221.	2.4	44
32	Evidence of susceptibility and resistance to cryptic X-linked meiotic drive in natural populations of Drosophila melanogaster. Evolution; International Journal of Organic Evolution, 2005, 59, 1280-91.	1.1	11
33	Brief communication: Ancient DNA prospects from Sri Lankan highland dry caves support an emerging global pattern. American Journal of Physical Anthropology, 2003, 121, 112-116.	2.1	27
34	Genome-wide variation in the human and fruitfly: a comparison. Current Opinion in Genetics and Development, 2001, 11, 627-634.	1.5	91