

Mariah H Meek

List of Publications by Year in descending order

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Version: 2024-02-01

32
papers

1,179
citations

623734

14
h-index

454955

30
g-index

34
all docs

34
docs citations

34
times ranked

1933
citing authors

#	ARTICLE	IF	CITATIONS
1	A nonfunctional copy of the salmonid sex-determining gene (<i>sdY</i>) is responsible for the apparent XY females in Chinook salmon, <i>Oncorhynchus tshawytscha</i> . <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	1.8	3
2	The Coalition for Conservation Genetics: Working across organizations to build capacity and achieve change in policy and practice. <i>Conservation Science and Practice</i> , 2022, 4, .	2.0	17
3	Trends in ecology and conservation over eight decades. <i>Frontiers in Ecology and the Environment</i> , 2021, 19, 274-282.	4.0	48
4	Global Commitments to Conserving and Monitoring Genetic Diversity Are Now Necessary and Feasible. <i>BioScience</i> , 2021, 71, 964-976.	4.9	96
5	Identifying hidden biocomplexity and genomic diversity in Chinook salmon, an imperiled species with a history of anthropogenic influence. <i>Canadian Journal of Fisheries and Aquatic Sciences</i> , 2020, 77, 534-547.	1.4	4
6	Attack of the PCR clones: Rates of clonality have little effect on RADseq genotype calls. <i>Molecular Ecology Resources</i> , 2020, 20, 66-78.	4.8	16
7	Evaluating Bioinformatic Pipeline Performance for Forensic Microbiome Analysis ^{*,†,‡} . <i>Journal of Forensic Sciences</i> , 2020, 65, 513-525.	1.6	10
8	Research management partnerships: An opportunity to integrate genetics in conservation actions. <i>Conservation Science and Practice</i> , 2020, 2, e218.	2.0	31
9	Dysbiosis in the Dead: Human Postmortem Microbiome Beta-Dispersion as an Indicator of Manner and Cause of Death. <i>Frontiers in Microbiology</i> , 2020, 11, 555347.	3.5	25
10	Linking gene expression patterns with survival studies elucidates adaptive potential in changing environments. <i>Molecular Ecology</i> , 2020, 29, 1031-1034.	3.9	3
11	The future is now: Amplicon sequencing and sequence capture usher in the conservation genomics era. <i>Molecular Ecology Resources</i> , 2019, 19, 795-803.	4.8	94
12	Physical defenses and herbivory vary more within plants than among plants in the tropical understory shrub <i>Piper polytrichum</i> . <i>Botany</i> , 2019, 97, 113-121.	1.0	9
13	Genomic Analysis Reveals Genetic Distinctiveness of the Paiute Cutthroat Trout <i>Oncorhynchus clarkii seleniris</i> . <i>Transactions of the American Fisheries Society</i> , 2017, 146, 1291-1302.	1.4	8
14	Sequencing improves our ability to study threatened migratory species: Genetic population assignment in California's Central Valley Chinook salmon. <i>Ecology and Evolution</i> , 2016, 6, 7706-7716.	1.9	24
15	Migration-related phenotypic divergence is associated with epigenetic modifications in rainbow trout. <i>Molecular Ecology</i> , 2016, 25, 1785-1800.	3.9	121
16	RAD Capture (Rapture): Flexible and Efficient Sequence-Based Genotyping. <i>Genetics</i> , 2016, 202, 389-400.	2.9	366
17	We should not be afraid to talk about fear of failure in conservation. <i>Biological Conservation</i> , 2016, 194, 218-219.	4.1	1
18	Transcriptional Response to Acute Thermal Exposure in Juvenile Chinook Salmon Determined by RNAseq. <i>G3: Genes, Genomes, Genetics</i> , 2015, 5, 1335-1349.	1.8	61

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19	Fear of failure in conservation: The problem and potential solutions to aid conservation of extremely small populations. <i>Biological Conservation</i> , 2015, 184, 209-217.	4.1	60
20	Using Next-Generation Sequencing to Assist a Conservation Hatchery: a Single-Nucleotide Polymorphism Panel for the Genetic Management of Endangered Delta Smelt. <i>Transactions of the American Fisheries Society</i> , 2015, 144, 767-779.	1.4	17
21	Genetic Considerations for Sourcing Steelhead Reintroductions: Investigating Possibilities for the San Joaquin River. <i>San Francisco Estuary and Watershed Science</i> , 2014, 12, .	0.4	4
22	Examining the Causes and Consequences of Hybridization During Chinook Salmon Reintroductions: Using the San Joaquin River as a Restoration Case Study of Management Options. <i>San Francisco Estuary and Watershed Science</i> , 2014, 12, .	0.4	0
23	Genetic characterization of California's Central Valley chinook salmon. <i>Ecology</i> , 2014, 95, 1431-1431.	3.2	1
24	Genetic diversity and reproductive mode in two non-native hydromedusae, <i>Maeotias marginata</i> and <i>Moerisia</i> sp., in the upper San Francisco Estuary, California. <i>Biological Invasions</i> , 2013, 15, 199-212.	2.4	7
25	Graduate Student's Guide to Necessary Skills for Nonacademic Conservation Careers. <i>Conservation Biology</i> , 2013, 27, 24-34.	4.7	77
26	Abundance, size, and diel feeding ecology of <i>Blackfordia virginica</i> (Mayer, 1910), a non-native hydrozoan in the lower Napa and Petaluma Rivers, California (USA). <i>Aquatic Invasions</i> , 2013, 8, 147-156.	1.6	16
27	Climate Change Likely to Facilitate the Invasion of the Non-Native Hydroid, <i>Cordylophora caspia</i> , in the San Francisco Estuary. <i>PLoS ONE</i> , 2012, 7, e46373.	2.5	11
28	Trophic ecology of two non-native hydrozoan medusae in the upper San Francisco Estuary. <i>Marine and Freshwater Research</i> , 2011, 62, 952.	1.3	15
29	Life history and population dynamics of <i>Moerisia</i> sp., a non-native hydrozoan, in the upper San Francisco Estuary (U.S.A.). <i>Estuarine, Coastal and Shelf Science</i> , 2011, 94, 48-55.	2.1	7
30	Ecological insights into the polyp stage of non-native hydrozoans in the San Francisco Estuary. <i>Aquatic Ecology</i> , 2011, 45, 151-161.	1.5	20
31	Isolation and characterization of microsatellite loci in two non-native hydromedusae in the San Francisco Estuary: <i>Maeotias marginata</i> and <i>Moerisia</i> sp.. <i>Conservation Genetics Resources</i> , 2009, 1, 205-208.	0.8	1
32	Black Sea Jellyfish: Shocking Newcomers to Suisun Marsh. <i>Frontiers for Young Minds</i> , 0, 9, .	0.8	0