

Elsbeth A Mclellan

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5684899/publications.pdf>

Version: 2024-02-01

13
papers

284
citations

1307594

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h-index

1125743

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13
all docs

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docs citations

13
times ranked

432
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | DNA metabarcoding reveals a broad dietary range for Tasmanian devils introduced to a naive ecosystem. <i>Ecology and Evolution</i> , 2022, 12, . | 1.9 | 4 |
| 2 | Restoring faith in conservation action: Maintaining wild genetic diversity through the Tasmanian devil insurance program. <i>IScience</i> , 2022, 25, 104474. | 4.1 | 8 |
| 3 | Assessing evolutionary processes over time in a conservation breeding program: a combined approach using molecular data, simulations and pedigree analysis. <i>Biodiversity and Conservation</i> , 2021, 30, 1011-1029. | 2.6 | 12 |
| 4 | How much is enough? Sampling intensity influences estimates of reproductive variance in an introduced population. <i>Ecological Applications</i> , 2021, , e02462. | 3.8 | 2 |
| 5 | Metapopulation management of a critically endangered marsupial in the age of genomics. <i>Global Ecology and Conservation</i> , 2021, 31, e01869. | 2.1 | 11 |
| 6 | Investigating inbreeding in a free-ranging, captive population of an Australian marsupial. <i>Conservation Genetics</i> , 2020, 21, 665-675. | 1.5 | 5 |
| 7 | A demonstration of conservation genomics for threatened species management. <i>Molecular Ecology Resources</i> , 2020, 20, 1526-1541. | 4.8 | 54 |
| 8 | Mixing genetically differentiated populations successfully boosts diversity of an endangered carnivore. <i>Animal Conservation</i> , 2020, 23, 700-712. | 2.9 | 23 |
| 9 | Too much of a good thing? Finding the most informative genetic data set to answer conservation questions. <i>Molecular Ecology Resources</i> , 2019, 19, 659-671. | 4.8 | 25 |
| 10 | From reference genomes to population genomics: comparing three reference-aligned reduced-representation sequencing pipelines in two wildlife species. <i>BMC Genomics</i> , 2019, 20, 453. | 2.8 | 48 |
| 11 | Complex problems need detailed solutions: Harnessing multiple data types to inform genetic management in the wild. <i>Evolutionary Applications</i> , 2019, 12, 280-291. | 3.1 | 28 |
| 12 | Pedigree reconstruction using molecular data reveals an early warning sign of gene diversity loss in an island population of Tasmanian devils (<i>Sarcophilus harrisii</i>). <i>Conservation Genetics</i> , 2018, 19, 439-450. | 1.5 | 27 |
| 13 | Immunization Strategies Producing a Humoral IgG Immune Response against Devil Facial Tumor Disease in the Majority of Tasmanian Devils Destined for Wild Release. <i>Frontiers in Immunology</i> , 2018, 9, 259. | 4.8 | 37 |