

Elizabeth L. Clare

List of Publications by Year in descending order

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Version: 2024-02-01

50
papers

3,684
citations

196777

29
h-index

232693

48
g-index

58
all docs

58
docs citations

58
times ranked

4143
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Measuring biodiversity from DNA in the air. <i>Current Biology</i> , 2022, 32, 693-700.e5. | 1.8 | 58 |
| 2 | The structure of tropical bat–plant interaction networks during an extreme El Niño–Southern Oscillation event. <i>Molecular Ecology</i> , 2022, 31, 1892-1906. | 2.0 | 7 |
| 3 | Shunning the scoop: Sidestepping the race to publish. <i>iScience</i> , 2022, 25, 104080. | 1.9 | 0 |
| 4 | Differential effects of fertilisers on pollination and parasitoid interaction networks. <i>Journal of Animal Ecology</i> , 2021, 90, 404-414. | 1.3 | 4 |
| 5 | Assessing the impact of taxon resolution on network structure. <i>Ecology</i> , 2021, 102, e03256. | 1.5 | 19 |
| 6 | Leech blood–meal invertebrate–derived DNA reveals differences in Bornean mammal diversity across habitats. <i>Molecular Ecology</i> , 2021, 30, 3299-3312. | 2.0 | 24 |
| 7 | eDNAir: proof of concept that animal DNA can be collected from air sampling. <i>PeerJ</i> , 2021, 9, e11030. | 0.9 | 58 |
| 8 | Biodiversity assessment across a dynamic riverine system: A comparison of eDNA metabarcoding versus traditional fish surveying methods. <i>Environmental DNA</i> , 2021, 3, 1247-1266. | 3.1 | 29 |
| 9 | Dung beetles as samplers of mammals in Malaysian Borneo—a test of high throughput metabarcoding of iDNA. <i>PeerJ</i> , 2021, 9, e11897. | 0.9 | 21 |
| 10 | Molecular diet analysis of the marine fish-eating bat (<i>Myotis vivax</i>) and potential mercury exposure. <i>Canadian Journal of Zoology</i> , 2021, 99, 752-759. | 0.4 | 3 |
| 11 | Altered structure of bat–prey interaction networks in logged tropical forests revealed by metabarcoding. <i>Molecular Ecology</i> , 2021, 30, 5844-5857. | 2.0 | 10 |
| 12 | Selective Logging Shows No Impact on the Dietary Breadth of a Generalist Bat Species: The Fawn Leaf-Nosed Bat (<i>Hipposideros cervinus</i>). <i>Frontiers in Ecology and Evolution</i> , 2021, 9, . | 1.1 | 0 |
| 13 | Occurrence of blood–feeding terrestrial leeches (Haemadipsidae) in a degraded forest ecosystem and their potential as ecological indicators. <i>Biotropica</i> , 2020, 52, 302-312. | 0.8 | 9 |
| 14 | Trophic resource partitioning drives fine-scale coexistence in cryptic bat species. <i>Ecology and Evolution</i> , 2020, 10, 14122-14136. | 0.8 | 14 |
| 15 | Wing morphology predicts individual niche specialization in <i>Pteronotus mesoamericanus</i> (Mammalia: Tj ETQq1 1 0,784314 rgBT /Overl | 1.1 | 14 |
| 16 | Counting with <i>scp>DNA</scp></i> in metabarcoding studies: How should we convert sequence reads to dietary data?. <i>Molecular Ecology</i> , 2019, 28, 391-406. | 2.0 | 455 |
| 17 | <i>scp>DNA</scp></i> metabarcoding reveals changes in the contents of carnivorous plants along an elevation gradient. <i>Molecular Ecology</i> , 2019, 28, 281-292. | 2.0 | 6 |
| 18 | Molecular diet analysis finds an insectivorous desert bat community dominated by resource sharing despite diverse echolocation and foraging strategies. <i>Ecology and Evolution</i> , 2019, 9, 3117-3129. | 0.8 | 38 |

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|----|--|-----|-----------|
| 19 | Using metabarcoding to compare the suitability of two blood-feeding leech species for sampling mammalian diversity in North Borneo. <i>Molecular Ecology Resources</i> , 2019, 19, 105-117. | 2.2 | 31 |
| 20 | Approaches to integrating genetic data into ecological networks. <i>Molecular Ecology</i> , 2019, 28, 503-519. | 2.0 | 37 |
| 21 | Assessing niche partitioning of co-occurring sibling bat species by <i>scp>DNA</scp></i> metabarcoding. <i>Molecular Ecology</i> , 2018, 27, 1273-1283. | 2.0 | 52 |
| 22 | Impact of urbanisation and agriculture on the diet of fruit bats. <i>Urban Ecosystems</i> , 2018, 21, 61-70. | 1.1 | 30 |
| 23 | The effects of pastoral intensification on the feeding interactions of generalist predators in streams. <i>Molecular Ecology</i> , 2018, 27, 590-602. | 2.0 | 9 |
| 24 | Diet tracing in ecology: Method comparison and selection. <i>Methods in Ecology and Evolution</i> , 2018, 9, 278-291. | 2.2 | 320 |
| 25 | Spatiotemporal and demographic variation in the diet of New Zealand lesser short-tailed bats (<i>Mystacina tuberculata</i>). <i>Ecology and Evolution</i> , 2018, 8, 7599-7610. | 0.8 | 17 |
| 26 | Flower preferences and pollen transport networks for cavity-nesting solitary bees: Implications for the design of agricultural environment schemes. <i>Ecology and Evolution</i> , 2018, 8, 7574-7587. | 0.8 | 44 |
| 27 | The effects of parameter choice on defining molecular operational taxonomic units and resulting ecological analyses of metabarcoding data. <i>Genome</i> , 2016, 59, 981-990. | 0.9 | 73 |
| 28 | Barcoding the food chain: from Sanger to high-throughput sequencing. <i>Genome</i> , 2016, 59, 946-958. | 0.9 | 27 |
| 29 | Dietary overlap and seasonality in three species of mormoopid bats from a tropical dry forest. <i>Molecular Ecology</i> , 2015, 24, 5296-5307. | 2.0 | 52 |
| 30 | Acoustic shadows help gleaning bats find prey, but may be defeated by prey acoustic camouflage on rough surfaces. <i>ELife</i> , 2015, 4, . | 2.8 | 16 |
| 31 | Molecular detection of trophic interactions: emerging trends, distinct advantages, significant considerations and conservation applications. <i>Evolutionary Applications</i> , 2014, 7, 1144-1157. | 1.5 | 163 |
| 32 | An inordinate fondness for beetles? Variation in seasonal dietary preferences of night-roosting big brown bats (<i>Myotisotis fuscus</i>). <i>Molecular Ecology</i> , 2014, 23, 3633-3647. | 2.0 | 105 |
| 33 | Diet of the insectivorous bat <i>Myotisotis nathusii</i> during autumn migration and summer residence. <i>Molecular Ecology</i> , 2014, 23, 3672-3683. | 2.0 | 57 |
| 34 | Dietary competition between the alien <i>Myotisotis sianuskshrew</i> (<i>Myotisotis murinus</i>) and a re-introduced population of <i>Myotisotis elfair's</i> <i>Myotisotis kink</i> (<i>Myotisotis telairii</i>). <i>Molecular Ecology</i> , 2014, 23, 3695-3705. | 2.0 | 65 |
| 35 | An integrative approach to detect subtle trophic niche differentiation in the sympatric trawling bat species <i>Myotisotis dasynceme</i> and <i>Myotisotis daubentonii</i> . <i>Molecular Ecology</i> , 2014, 23, 3657-3671. | 2.0 | 50 |
| 36 | A pragmatic approach to the analysis of diets of generalist predators: the use of next-generation sequencing with no blocking probes. <i>Molecular Ecology Resources</i> , 2014, 14, 18-26. | 2.2 | 147 |

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|----|--|-----|-----------|
| 37 | Resource partitioning by insectivorous bats in Jamaica. <i>Molecular Ecology</i> , 2014, 23, 3648-3656. | 2.0 | 68 |
| 38 | Trophic niche flexibility in <i>Geophaga soricina</i> : how a nectar seeker sneaks an insect snack. <i>Functional Ecology</i> , 2014, 28, 632-641. | 1.7 | 51 |
| 39 | The diet of <i>Myotis lucifugus</i> across Canada: assessing foraging quality and diet variability. <i>Molecular Ecology</i> , 2014, 23, 3618-3632. | 2.0 | 111 |
| 40 | Island bat diets: does it matter more who you are or where you live?. <i>Molecular Ecology</i> , 2014, 23, 3684-3694. | 2.0 | 19 |
| 41 | Diversification and reproductive isolation: cryptic species in the only New World high-duty cycle bat, <i>Pteronotus parnellii</i> . <i>BMC Evolutionary Biology</i> , 2013, 13, 26. | 3.2 | 54 |
| 42 | DNA Barcoding in Mammals. <i>Methods in Molecular Biology</i> , 2012, 858, 153-182. | 0.4 | 63 |
| 43 | Neotropical Bats: Estimating Species Diversity with DNA Barcodes. <i>PLoS ONE</i> , 2011, 6, e22648. | 1.1 | 138 |
| 44 | Eating local: influences of habitat on the diet of little brown bats (<i>Myotis lucifugus</i>). <i>Molecular Ecology</i> , 2011, 20, 1772-1780. | 2.0 | 170 |
| 45 | High-throughput sequencing offers insight into mechanisms of resource partitioning in cryptic bat species. <i>Ecology and Evolution</i> , 2011, 1, 556-570. | 0.8 | 163 |
| 46 | Molecular Diet Analysis of Two African Free-Tailed Bats (Molossidae) Using High Throughput Sequencing. <i>PLoS ONE</i> , 2011, 6, e21441. | 1.1 | 175 |
| 47 | Cryptic Species? Patterns of Maternal and Paternal Gene Flow in Eight Neotropical Bats. <i>PLoS ONE</i> , 2011, 6, e21460. | 1.1 | 55 |
| 48 | Species on the menu of a generalist predator, the eastern red bat (<i>Lasiurus borealis</i>): using a molecular approach to detect arthropod prey. <i>Molecular Ecology</i> , 2009, 18, 2532-2542. | 2.0 | 225 |
| 49 | Diagnosing Mitochondrial DNA Diversity: Applications of a Sentinel Gene Approach. <i>Journal of Molecular Evolution</i> , 2008, 66, 362-367. | 0.8 | 39 |
| 50 | DNA barcoding of Neotropical bats: species identification and discovery within Guyana. <i>Molecular Ecology Notes</i> , 2007, 7, 184-190. | 1.7 | 261 |