## Nicole E White

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2427249/publications.pdf

Version: 2024-02-01

840776 839539 22 445 11 18 citations h-index g-index papers 22 22 22 622 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Testing multiple substrates for terrestrial biodiversity monitoring using environmental DNA metabarcoding. Molecular Ecology Resources, 2020, 20, 732-745.	4.8	60
2	Development and evaluation of fish eDNA metabarcoding assays facilitate the detection of cryptic seahorse taxa (family: Syngnathidae). Environmental DNA, 2020, 2, 614-626.	5.8	48
3	New light in the dark - a proposed multidisciplinary framework for studying functional ecology of groundwater fauna. Science of the Total Environment, 2019, 662, 963-977.	8.0	47
4	Salt to conserve: a review on the ecology and preservation of hypersaline ecosystems. Biological Reviews, 2021, 96, 2828-2850.	10.4	47
5	Development of a multi-assay approach for monitoring coral diversity using eDNA metabarcoding. Coral Reefs, 2020, 39, 159-171.	2.2	42
6	High-throughput sequencing of ancient plant and mammal DNA preserved in herbivore middens. Quaternary Science Reviews, 2012, 58, 135-145.	3.0	40
7	eDNA in subterranean ecosystems: Applications, technical aspects, and future prospects. Science of the Total Environment, 2022, 820, 153223.	8.0	38
8	Rainfall as a trigger of ecological cascade effects in an Australian groundwater ecosystem. Scientific Reports, 2021, 11, 3694.	3.3	20
9	Detection of the rare Australian endemic blind cave eel (Ophisternon candidum) with environmental DNA: implications for threatened species management in subterranean environments. Hydrobiologia, 2020, 847, 3201-3211.	2.0	16
10	Identifying conservation units after large-scale land clearing: a spatio-temporal molecular survey of endangered white-tailed black cockatoos (Calyptorhynchusspp.). Diversity and Distributions, 2014, 20, 1208-1220.	4.1	15
11	The applicability of eDNA metabarcoding approaches for sessile benthic surveying in the Kimberley region, northâ€western Australia. Environmental DNA, 2022, 4, 34-49.	5.8	15
12	Tracking down carbon inputs underground from an arid zone Australian calcrete. PLoS ONE, 2020, 15, e0237730.	2.5	14
13	Considerations for future environmental DNA accreditation and proficiency testing schemes. Environmental DNA, 2021, 3, 1049-1058.	5.8	12
14	Determination of the diet of the ghost bat (Macroderma gigas) in the Pilbara region of Western Australia from dried prey remains and DNA metabarcoding. Australian Journal of Zoology, 2018, 66, 195.	1.0	10
15	Metabarcoding under Brine: Microbial Ecology of Five Hypersaline Lakes at Rottnest Island (WA,) Tj ETQq1 1 0.784	1314 rgBT	/Overlock 1
16	Phylogeography of recent Plesiastrea (Scleractinia: Plesiastreidae) based on an integrated taxonomic approach. Molecular Phylogenetics and Evolution, 2022, 172, 107469.	2.7	6
17	Evaluating restoration trajectories using DNA metabarcoding of groundâ€dwelling and airborne invertebrates and associated plant communities. Molecular Ecology, 2022, 31, 2172-2188.	3.9	5
18	Crop-Zone Weed Mycobiomes of the South-Western Australian Grain Belt. Frontiers in Microbiology, 2020, 11, 581592.	3.5	4

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19	Tracking down carbon inputs underground from an arid zone Australian calcrete. , 2020, 15, e0237730.		O
20	Tracking down carbon inputs underground from an arid zone Australian calcrete., 2020, 15, e0237730.		0
21	Tracking down carbon inputs underground from an arid zone Australian calcrete. , 2020, 15, e0237730.		O
22	Tracking down carbon inputs underground from an arid zone Australian calcrete., 2020, 15, e0237730.		0