

# Conrad A Matthee

## List of Publications by Year in descending order

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133  
papers

4,876  
citations

76294

40  
h-index

110317

64  
g-index

134  
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134  
docs citations

134  
times ranked

4936  
citing authors

#	ARTICLE	IF	CITATIONS
1	Pattern and timing of diversification of Cetartiodactyla (Mammalia, Laurasiatheria), as revealed by a comprehensive analysis of mitochondrial genomes. <i>Comptes Rendus - Biologies</i> , 2012, 335, 32-50.	0.1	448
2	A Nuclear DNA Phylogenetic Perspective on the Evolution of Echolocation and Historical Biogeography of Extant Bats (Chiroptera). <i>Molecular Biology and Evolution</i> , 2005, 22, 1869-1886.	3.5	211
3	A Molecular Supermatrix of the Rabbits and Hares (Leporidae) Allows for the Identification of Five Intercontinental Exchanges During the Miocene. <i>Systematic Biology</i> , 2004, 53, 433-447.	2.7	198
4	Resolution of a Supertree/Supermatrix Paradox. <i>Systematic Biology</i> , 2002, 51, 652-664.	2.7	190
5	Mining the Mammalian Genome for Artiodactyl Systematics. <i>Systematic Biology</i> , 2001, 50, 367-390.	2.7	136
6	Molecular Insights into the Evolution of the Family Bovidae: A Nuclear DNA Perspective. <i>Molecular Biology and Evolution</i> , 2001, 18, 1220-1230.	3.5	125
7	Biogeographic patterns and phylogeography of dwarf chameleons ( <i>Bradypodion</i> ) in an African biodiversity hotspot. <i>Molecular Ecology</i> , 2006, 15, 781-793.	2.0	107
8	Ancient forest fragmentation or recent radiation? Testing refugial speciation models in chameleons within an African biodiversity hotspot. <i>Journal of Biogeography</i> , 2011, 38, 1748-1760.	1.4	87
9	Suprafamilial relationships among Rodentia and the phylogenetic effect of removing fast-evolving nucleotides in mitochondrial, exon and intron fragments. <i>BMC Evolutionary Biology</i> , 2008, 8, 321.	3.2	84
10	The evolutionary history of seahorses (Syngnathidae: Hippocampus): molecular data suggest a West Pacific origin and two invasions of the Atlantic Ocean. <i>Molecular Phylogenetics and Evolution</i> , 2004, 30, 273-286.	1.2	82
11	Molecular evidence for long-distance colonization in an Indo-Pacific seahorse lineage. <i>Marine Ecology - Progress Series</i> , 2005, 286, 249-260.	0.9	78
12	Cytochrome b Phylogeny of the Family Bovidae: Resolution within the Alcelaphini, Antilopini, Neotragini, and Tragelaphini. <i>Molecular Phylogenetics and Evolution</i> , 1999, 12, 31-46.	1.2	77
13	Phylogenetic relationships of the southern African freshwater crab fauna (Decapoda: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 20 Molecular Phylogenetics and Evolution, 2002, 25, 511-523.	1.2	77
14	Mining the Mammalian Genome for Artiodactyl Systematics. <i>Systematic Biology</i> , 2001, 50, 367-390.	2.7	74
15	Biome specificity of distinct genetic lineages within the four-striped mouse <i>Rhabdomys pumilio</i> (Rodentia: Muridae) from southern Africa with implications for taxonomy. <i>Molecular Phylogenetics and Evolution</i> , 2012, 65, 75-86.	1.2	74
16	Biotic diversity in the Southern African winter-rainfall region. <i>Current Opinion in Environmental Sustainability</i> , 2010, 2, 109-116.	3.1	73
17	Population fragmentation in the southern rock agama, <i>Agama atra</i> : more evidence for vicariance in Southern Africa. <i>Molecular Ecology</i> , 2002, 11, 465-471.	2.0	71
18	Mitochondrial DNA analyses of the Cape hakes reveal an expanding, panmictic population for <i>Merluccius capensis</i> and population structuring for mature fish in <i>Merluccius paradoxus</i> . <i>Molecular Phylogenetics and Evolution</i> , 2007, 42, 517-527.	1.2	69

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19	Genetic isolation by distance reveals restricted dispersal across a range of life histories: implications for biodiversity conservation planning across highly variable marine environments. <i>Diversity and Distributions</i> , 2015, 21, 698-710.	1.9	67
20	Phylogenetics of the southern African dwarf chameleons, <i>Bradypodion</i> (Squamata: Chamaeleonidae). <i>Molecular Phylogenetics and Evolution</i> , 2004, 30, 354-365.	1.2	65
21	Mitochondrial DNA variation in spiny lobster <i>Palinurus delagoae</i> suggests genetically structured populations in the southwestern Indian Ocean. <i>Marine Ecology - Progress Series</i> , 2006, 319, 191-198.	0.9	65
22	Molecular phylogenetics and historical biogeography of <i>Rhinolophus</i> bats. <i>Molecular Phylogenetics and Evolution</i> , 2010, 54, 1-9.	1.2	64
23	Phylogeny and evolutionary origins of the Leporidae: a review of cytogenetics, molecular analyses and a supermatrix analysis. <i>Mammal Review</i> , 2005, 35, 231-247.	2.2	62
24	Mitochondrial DNA population structure of roan and sable antelope: implications for the translocation and conservation of the species. <i>Molecular Ecology</i> , 1999, 8, 227-238.	2.0	61
25	A phylogenetic review of the African leaf chameleons: genus <i>Rhampholeon</i> (Chamaeleonidae): the role of vicariance and climate change in speciation. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2004, 271, 1967-1975.	1.2	60
26	Range expansions across ecoregions: interactions of climate change, physiology and genetic diversity. <i>Global Ecology and Biogeography</i> , 2014, 23, 76-88.	2.7	59
27	Misleading the masses: detection of mislabelled and substituted frozen fish products in South Africa. <i>ICES Journal of Marine Science</i> , 2010, 67, 176-185.	1.2	58
28	Climate change drives speciation in the southern rock agama ( <i>Agama atra</i> ) in the Cape Floristic Region, South Africa. <i>Journal of Biogeography</i> , 2009, 36, 78-87.	1.4	57
29	Mitochondrial DNA panmixia in spiny lobster <i>Palinurus gilchristi</i> suggests a population expansion. <i>Marine Ecology - Progress Series</i> , 2005, 297, 225-231.	0.9	57
30	Utility of nuclear DNA intron markers at lower taxonomic levels: Phylogenetic resolution among nine <i>Tragelaphus</i> spp.. <i>Molecular Phylogenetics and Evolution</i> , 2005, 35, 624-636.	1.2	56
31	Population genetics of the endangered Knysna seahorse, <i>Hippocampus capensis</i> . <i>Molecular Ecology</i> , 2003, 12, 1703-1715.	2.0	55
32	Indel evolution of mammalian introns and the utility of non-coding nuclear markers in eutherian phylogenetics. <i>Molecular Phylogenetics and Evolution</i> , 2007, 42, 827-837.	1.2	55
33	Evidence for panmixia despite barriers to gene flow in the southern African endemic, <i>Caffrogobius caffer</i> (Teleostei: Gobiidae). <i>BMC Evolutionary Biology</i> , 2008, 8, 325.	3.2	54
34	Title is missing!. <i>Journal of Mammalian Evolution</i> , 1997, 4, 53-73.	1.0	53
35	Molecular systematics of dormice (Rodentia: Gliridae) and the radiation of <i>Graphiurus</i> in Africa. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2003, 270, 1947-1955.	1.2	52
36	Biogeography and host-related factors trump parasite life history: limited congruence among the genetic structures of specific ectoparasitic lice and their rodent hosts. <i>Molecular Ecology</i> , 2013, 22, 5185-5204.	2.0	50

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37	Signatures of seaway closures and founder dispersal in the phylogeny of a circumglobally distributed seahorse lineage. <i>BMC Evolutionary Biology</i> , 2007, 7, 138.	3.2	46
38	Effects of tectonics and large scale climatic changes on the evolutionary history of <i>Hyalomma</i> ticks. <i>Molecular Phylogenetics and Evolution</i> , 2017, 114, 153-165.	1.2	45
39	CRYPTIC SPECIES IN AN INSECTIVOROUS BAT, <i>SCOTOPHILUS DINGANII</i> . <i>Journal of Mammalogy</i> , 2006, 87, 161-170.	0.6	44
40	The potential for predicted climate shifts to impact genetic landscapes of lizards in the South African Cape Floristic Region. <i>Molecular Phylogenetics and Evolution</i> , 2009, 51, 120-130.	1.2	44
41	Mitochondrial DNA differentiation among geographical populations of <i>Pronolagus rupestris</i> , Smith's red rock rabbit (Mammalia: Lagomorpha). <i>Heredity</i> , 1996, 76, 514-523.	1.2	42
42	Mitochondrial DNA sequence data of the Cape fur seal ( <i>Arctocephalus pusillus pusillus</i> ) suggest that population numbers may be affected by climatic shifts. <i>Marine Biology</i> , 2006, 148, 899-905.	0.7	41
43	Remarkably low mtDNA control region diversity in an abundant demersal fish. <i>Molecular Phylogenetics and Evolution</i> , 2010, 55, 1183-1188.	1.2	41
44	Molecular phylogeny of the springhare, <i>Pedetes capensis</i> , based on mitochondrial DNA sequences. <i>Molecular Biology and Evolution</i> , 1997, 14, 20-29.	3.5	36
45	Integrative taxonomy and species delimitation of <i>Rhipicephalus turanicus</i> (Acari: Ixodida: Ixodidae). <i>International Journal for Parasitology</i> , 2020, 50, 577-594.	1.3	34
46	Spatio-temporal genetic structure and the effects of long-term fishing in two partially sympatric offshore demersal fishes. <i>Molecular Ecology</i> , 2016, 25, 5843-5861.	2.0	33
47	An integrated mark-recapture and genetic approach to estimate the population size of white sharks in South Africa. <i>Marine Ecology - Progress Series</i> , 2016, 552, 241-253.	0.9	33
48	Mining the mammalian genome for artiodactyl systematics. <i>Systematic Biology</i> , 2001, 50, 367-90.	2.7	33
49	Linking lineage diversification to climate and habitat heterogeneity: phylogeography of the southern African shrew <i>Myosorex varius</i> . <i>Journal of Biogeography</i> , 2011, 38, 1976-1991.	1.4	32
50	Nuclear and mtDNA-based phylogeny of southern African sand lizards, <i>Pedioplanis</i> (Sauria: Lacertidae). <i>Molecular Phylogenetics and Evolution</i> , 2007, 44, 622-633.	1.2	30
51	Molecular phylogeny of the spiny lobster genus <i>Palinurus</i> (Decapoda: Palinuridae) with hypotheses on speciation in the NE Atlantic/Mediterranean and SW Indian Ocean. <i>Molecular Phylogenetics and Evolution</i> , 2007, 45, 102-110.	1.2	29
52	Phylogeographic analysis of nuclear and mtDNA supports subspecies designations in the ostrich ( <i>Struthio camelus</i> ). <i>Conservation Genetics</i> , 2011, 12, 423-431.	0.8	29
53	Evolutionary history of the Karoo bush rat, <i>Myotomys unisulcatus</i> (Rodentia: Muridae): discordance between morphology and genetics. <i>Biological Journal of the Linnean Society</i> , 2011, 102, 510-526.	0.7	28
54	The Divergence of Echolocation Frequency in Horseshoe Bats: Moth Hearing, Body Size or Habitat?. <i>Journal of Mammalian Evolution</i> , 2011, 18, 117-129.	1.0	28

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55	Mitochondrial DNA variation of the west-coast rock lobster, <i>Jasus lalandii</i> : marked genetic diversity differences among sampling sites. <i>Marine and Freshwater Research</i> , 2007, 58, 1130.	0.7	27
56	The evolution of the southern African members of the shrew genus <i>Myosorex</i> : Understanding the origin and diversification of a morphologically cryptic group. <i>Molecular Phylogenetics and Evolution</i> , 2009, 51, 394-398.	1.2	27
57	A New Species of <i>Ixodes</i> (Acari: Ixodidae) From South African Mammals. <i>Journal of Parasitology</i> , 2011, 97, 389-398.	0.3	27
58	Oceanic circulation, local upwelling and palaeoclimatic changes linked to the phylogeography of the Cape sea urchin <i>Parechinus angulosus</i> . <i>Marine Ecology - Progress Series</i> , 2012, 468, 203-215.	0.9	27
59	Species-specific genetic markers for identification of early life-history stages of Cape hakes, <i>Merluccius capensis</i> and <i>Merluccius paradoxus</i> in the southern Benguela Current. <i>Journal of Fish Biology</i> , 2007, 70, 262-268.	0.7	26
60	Four p67 alleles identified in South African <i>Theileria parva</i> field samples. <i>Veterinary Parasitology</i> , 2010, 167, 244-254.	0.7	26
61	Correlated Genetic and Ecological Diversification in a Widespread Southern African Horseshoe Bat. <i>PLoS ONE</i> , 2012, 7, e31946.	1.1	25
62	Comparative phylogeography between two generalist flea species reveal a complex interaction between parasite life history and host vicariance: parasite-host association matters. <i>BMC Evolutionary Biology</i> , 2015, 15, 105.	3.2	24
63	A shared unusual genetic change at the chemokine receptor type 5 between <i>Oryctolagus</i> , <i>Bunolagus</i> and <i>Pentalagus</i> . <i>Conservation Genetics</i> , 2011, 12, 325-330.	0.8	23
64	Poecilogony in <i>Polydora hoplura</i> (Polychaeta: Spionidae) from commercially important molluscs in South Africa. <i>Marine Biology</i> , 2014, 161, 887-898.	0.7	23
65	The Influence of Interspecific Competition and Host Preference on the Phylogeography of Two African Ixodid Tick Species. <i>PLoS ONE</i> , 2013, 8, e76930.	1.1	23
66	Molecular genetic relationships of the extinct ostrich, <i>Struthio camelus syriacus</i> : consequences for ostrich introductions into Saudi Arabia. <i>Animal Conservation</i> , 1999, 2, 165-171.	1.5	22
67	Long current to nowhere? Genetic connectivity of <i>Jasus tristani</i> populations in the southern Atlantic Ocean. <i>African Journal of Marine Science</i> , 2007, 29, 491-497.	0.4	22
68	Limited dispersal in an ectoparasitic mite, <i>Laelaps giganteus</i> , contributes to significant phylogeographic congruence with the rodent host, <i>Rhabdomys</i> . <i>Molecular Ecology</i> , 2016, 25, 1006-1021.	2.0	22
69	High connectivity and lack of mtDNA differentiation among two previously recognized spiny lobster species in the southern Atlantic and Indian Oceans. <i>Marine Biology Research</i> , 2012, 8, 764-770.	0.3	21
70	Dispersal and genetic structure of <i>Boccardia polybranchia</i> and <i>Polydora hoplura</i> (Annelida: Tj ETQq0 0 0 rgBT /Overlock 10 Tf, 50 142 Td	1.7	21
71	Implications of spatial genetic patterns for conserving African leopards. <i>Comptes Rendus - Biologies</i> , 2015, 338, 728-737.	0.1	17
72	New insights into the evolutionary history of white sharks, <i>Carcharodon carcharias</i> . <i>Journal of Biogeography</i> , 2016, 43, 328-339.	1.4	17

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73	The effect of host vicariance and parasite life history on the dispersal of the multi-host ectoparasite, <i>Hyalomma truncatum</i> . <i>Journal of Biogeography</i> , 2017, 44, 1124-1136.	1.4	17
74	Species delineation in the speciation grey zone—The case of <i>Diopatra</i> (Annelida, Onuphidae). <i>Zoologica Scripta</i> , 2020, 49, 516-534.	0.7	17
75	Effects of the Sharksafe barrier on white shark ( <i>Carcharodon carcharias</i> ) behavior and its implications for future conservation technologies. <i>Journal of Experimental Marine Biology and Ecology</i> , 2014, 460, 37-46.	0.7	16
76	Evidence of cryptic speciation in mesostigmatid mites from South Africa. <i>Parasitology</i> , 2014, 141, 1322-1332.	0.7	16
77	New taxonomic and evolutionary insights relevant to the cat flea, <i>Ctenocephalides felis</i> : A geographic perspective. <i>Molecular Phylogenetics and Evolution</i> , 2021, 155, 106990.	1.2	16
78	When homoplasy mimics hybridization: a case study of Cape hakes ( <i>Merluccius capensis</i> and <i>M. Tj ETQq0,00 rgBT /Overlock 1</i> ).	0.9	16
79	Western Zambian sable: Are they a Geographic Extension of the Giant sable Antelope?. <i>South African Journal of Wildlife Research</i> , 2010, 40, 35-42.	1.4	15
80	The Influence of Pleistocene Climatic Changes and Ocean Currents on the Phylogeography of the Southern African Barnacle, <i>Tetraclita serrata</i> (Thoracica; Cirripedia). <i>PLoS ONE</i> , 2014, 9, e102115.	1.1	15
81	Predicting the Dispersal Potential of an Invasive Polychaete Pest along a Complex Coastal Biome. <i>Integrative and Comparative Biology</i> , 2016, 56, 600-610.	0.9	15
82	Testing the exclusion capabilities and durability of the Sharksafe Barrier to determine its viability as an eco-friendly alternative to current shark culling methodologies. <i>Aquatic Conservation: Marine and Freshwater Ecosystems</i> , 2018, 28, 252-258.	0.9	15
83	Comparative phylogeography of parasitic Laelaps mites contribute new insights into the specialist-generalist variation hypothesis (SGVH). <i>BMC Evolutionary Biology</i> , 2018, 18, 131.	3.2	15
84	Tempo of genetic diversification in southern African rodents: The role of Plio-Pleistocene climatic oscillations as drivers for speciation. <i>Acta Oecologica</i> , 2012, 42, 50-57.	0.5	14
85	Nanger, Eudorcas, Gazella, and Antelope form a well-supported chromosomal clade within Antilopini (Bovidae, Cetartiodactyla). <i>Chromosoma</i> , 2015, 124, 235-247.	1.0	14
86	Towards resolving familial relationships within the Gadiformes, and the resurrection of the Lyconidae. <i>Molecular Phylogenetics and Evolution</i> , 2008, 48, 764-769.	1.2	13
87	First record of the pantropical blue tick <i>Rhipicephalus microplus</i> in Namibia. <i>Experimental and Applied Acarology</i> , 2013, 61, 503-507.	0.7	13
88	Adaptive radiation and speciation in <i>Rhipicephalus</i> ticks: A medley of novel hosts, nested predator-prey food webs, off-host periods and dispersal along temperature variation gradients. <i>Molecular Phylogenetics and Evolution</i> , 2021, 162, 107178.	1.2	13
89	The influence of life history characteristics on flea (Siphonaptera) species distribution models. <i>Parasites and Vectors</i> , 2016, 9, 178.	1.0	12
90	<i>Hippocampus queenslandicus</i> Horne, 2001 - a new seahorse species or yet another synonym?. <i>Australian Journal of Zoology</i> , 2007, 55, 139.	0.6	11

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91	A multiple data set phylogeny for the endemic South African freshwater phreatoicidan isopod genus <i>Mesamphisopus</i> : Taxonomic and biogeographic implications. <i>Molecular Phylogenetics and Evolution</i> , 2010, 55, 541-551.	1.2	11
92	Natural hosts of the larvae of <i>Nuttalliella</i> sp. ( <i>N. namaqua</i> ?) (Acari: Tj ETQq0 0 0 rgBT /Overlock 10 Tf	0.6	11
93	The sympatric occurrence of two genetically divergent lineages of sucking louse, <i>Polyplax arvicantis</i> (Phthiraptera: Anoplura), on the four-striped mouse genus, <i>Rhabdomys</i> (Rodentia: Muridae). <i>Parasitology</i> , 2013, 140, 604-616.	0.7	11
94	A molecular and morphological reassessment of the phylogeny of the subfamily Ophioninae (Hymenoptera: Ichneumonidae). <i>Zoological Journal of the Linnean Society</i> , 2016, 178, 128-148.	1.0	11
95	Lack of taxonomic differentiation in an apparently widespread freshwater isopod morphotype (Phreatoicidea: Mesamphisopidae: <i>Mesamphisopus</i> ) from South Africa. <i>Molecular Phylogenetics and Evolution</i> , 2005, 37, 289-305.	1.2	10
96	Exploring the Diversity and Molecular Evolution of Shrews (Family Soricidae) using mtDNA Cytochrome <i>b</i> Data. <i>African Zoology</i> , 2011, 46, 246-262.	0.2	10
97	The influence of life history and climate driven diversification on the mtDNA phylogeographic structures of two southern African <i>Mastomys</i> species (Rodentia: Muridae: Murinae). <i>Biological Journal of the Linnean Society</i> , 2015, 114, 58-68.	0.7	10
98	Moonshine worms ( <i>Diopatra aciculata</i> : Onuphidae, Annelida) in the Knysna Estuary, South Africa; taxonomy and distribution. <i>Journal of the Marine Biological Association of the United Kingdom</i> , 2020, 100, 897-907.	0.4	10
99	A novel categorisation system to organise a large photo identification database for white sharks <i>Carcharodon carcharias</i> . <i>African Journal of Marine Science</i> , 2014, 36, 59-67.	0.4	9
100	Community structure of fleas within and among populations of three closely related rodent hosts: nestedness and beta-diversity. <i>Parasitology</i> , 2016, 143, 1268-1278.	0.7	9
101	The evolutionary history of parasitic sucking lice and their rodent hosts: A case of evolutionary co-divergences. <i>Zoologica Scripta</i> , 2020, 49, 72-85.	0.7	9
102	Clarifying the cryptogenic species <i>Polydora neocaeca</i> Williams & Radashevsky, 1999 (Annelida: Tj ETQq0 0 0 rgBT /Overlock 10 Tf	0.3	9
103	Genetic Diversity Levels in Fishery-Exploited Spiny Lobsters of the Genus <i>Palinurus</i> (Decapoda: Tj ETQq1 1 0.784314 rgBT /Oyerlock 1	0.3	8
104	Genetic population structure and recruitment patterns of three sympatric shallow-water penaeid prawns in Ungwana Bay, Kenya, with implication for fisheries management. <i>Marine and Freshwater Research</i> , 2014, 65, 255.	0.7	8
105	Viruses as indicators of contemporary host dispersal and phylogeography: an example of feline immunodeficiency virus ( <i>FIV<sub>P</sub></i> ) in free-ranging African lion ( <i>Panthera leo</i> ). <i>Journal of Evolutionary Biology</i> , 2018, 31, 1529-1543.	0.8	8
106	Phylogeny and vicariant speciation of the Grey Rhebok, <i>Pelea capreolus</i> . <i>Heredity</i> , 2014, 112, 325-332.	1.2	7
107	A genetic perspective on the taxonomy and evolution of the medically important flea, <i>Dinopsyllus ellobius</i> (Siphonaptera: Dinopsyllinae), and the resurrection of <i>Dinopsyllus abaris</i> .	0.7	7
108	Comparative mtDNA phylogeographic patterns reveal marked differences in population genetic structure between generalist and specialist ectoparasites of the African penguin ( <i>Spheniscus</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 57 T	0.6	7

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109	The influence of host dispersal on the gene flow and genetic diversity of generalist and specialist ectoparasites. <i>African Zoology</i> , 2020, 55, 119-126.	0.2	7
110	Delimitation of morphologically similar sponge crab species of the genus <i>Pseudodromia</i> (Crustacea). <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5</i>	0.7	6
111	Effective number of white shark ( <i>Carcharodon carcharias</i> , Linnaeus) breeders is stable over four successive years in the population adjacent to eastern Australia and New Zealand. <i>Ecology and Evolution</i> , 2021, 11, 186-198.	0.8	6
112	High population connectivity and Pleistocene range expansion in the direct-developing plough shell <i>Bullia rhodostoma</i> along the South African coast. <i>African Journal of Marine Science</i> , 2015, 37, 21-31.	0.4	5
113	Comparative genetic structure in two high-dispersal prawn species from the south-west Indian Ocean. <i>African Journal of Marine Science</i> , 2017, 39, 467-474.	0.4	5
114	Semi-automated software for dorsal fin photographic identification of marine species: application to <i>Carcharodon carcharias</i> . <i>Marine Biodiversity</i> , 2018, 48, 1655-1660.	0.3	5
115	Genetic structure of bloodworm, <i>Arenicola loveni</i> (Annelida; Arenicolidae) suggests risk of local extinction in the face of overexploitation is lower than expected. <i>African Zoology</i> , 2020, 55, 175-183.	0.2	5
116	Reeling them in: taxonomy of marine annelids used as bait by anglers in the Western Cape Province, South Africa. <i>PeerJ</i> , 2021, 9, e11847.	0.9	5
117	Genetic assessment of seasonal alongshore migration in <i>Merluccius capensis</i> in the Benguela region. <i>Fisheries Research</i> , 2022, 250, 106293.	0.9	4
118	A microsatellite perspective on the reproductive success of subordinate male honey badgers, <i>Mellivora capensis</i> . <i>African Zoology</i> , 2004, 39, 305-308.	0.2	3
119	Comparative phylogeography between parasitic sucking lice and their host the Namaqua rock mouse, <i>Micaelamys namaquensis</i> (Rodentia: Muridae). <i>Zoological Journal of the Linnean Society</i> , 2021, 192, 1017-1028.	1.0	3
120	Molecular genetic relationships of the extinct ostrich, <i>Struthio camelus syriacus</i> : consequences for ostrich introductions into Saudi Arabia. <i>Animal Conservation</i> , 1999, 2, 165-171.	1.5	3
121	The role of controlled human-animal interactions in changing the negative perceptions towards white sharks, in a sample of White Shark cage diving tours participants. <i>Marine Policy</i> , 2022, 143, 105130.	1.5	3
122	The influence of kelp density on white shark presence within the Dyer Island nature reserve, South Africa. <i>Ocean and Coastal Management</i> , 2019, 179, 104819.	2.0	2
123	Erring on the side of caution: Reply to Irion et al. (2017). <i>Marine Ecology - Progress Series</i> , 2017, 577, 257-262.	0.9	2
124	The GDR1 191 (2007-2010): Biodiversity and global change in southern Africa. <i>Acta Oecologica</i> , 2012, 42, 1-2.	0.5	1
125	Species-landscape interactions drive divergent population trajectories in four forest-dependent Afromontane forest songbird species within a biodiversity hotspot in South Africa. <i>Evolutionary Applications</i> , 2021, 14, 2680-2697.	1.5	1
126	Two New Species of Sucking Lice (Phthiraptera: Anoplura: Hoplopleuridae and Polyplacidae) from Grant's Rock Mouse, <i>Micaelamys granti</i> , in South Africa. <i>Journal of Parasitology</i> , 2020, 106, 478.	0.3	1



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127	Climate refugia for three Afromontane forest-dependent bird species in south-eastern South Africa. <i>Journal of Biogeography</i> , 2022, 49, 1352-1366.	1.4	1
128	Their young bite better: On- and off-host selection pressure as drivers for evolutionary-developmental modification in <i>Rhipicephalus</i> ticks. <i>Arthropod Structure and Development</i> , 2022, 70, 101189.	0.8	1
129	The importance of correctly identifying the process responsible for spatial genetic structure in Leopard: A response to McManus and Smuts (2016). <i>Comptes Rendus - Biologies</i> , 2016, 339, 439-441.	0.1	0
130	Evaluating the Diversity of the Feline Immunodeficiency Virus (FIV): A Leopard Perspective. <i>African Journal of Wildlife Research</i> , 2017, 47, 92-105.	0.2	0
131	Conservation Genetics of the Critically Endangered Riverine Rabbit, <i>Bunolagus monticularis</i> : Structured Populations and High mtDNA Genetic Diversity. <i>Journal of Mammalian Evolution</i> , 2022, 29, 137-147.	1.0	0
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133	Artiodactyl Nuclear DNA Study. <i>Science</i> , 1999, 285, 1355-1355.	6.0	0