

# Carmelo AndÃ³jar

## List of Publications by Year in descending order

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Version: 2024-02-01

44  
papers

1,338  
citations

361045

20  
h-index

377514

34  
g-index

51  
all docs

51  
docs citations

51  
times ranked

1784  
citing authors

#	ARTICLE	IF	CITATIONS
1	Why the COI barcode should be the community <scp>DNA</scp> metabarcode for the metazoa. <i>Molecular Ecology</i> , 2018, 27, 3968-3975.	2.0	131
2	Winding up the molecular clock in the genus <i>Carabus</i> (Coleoptera: Carabidae): assessment of methodological decisions on rate and node age estimation. <i>BMC Evolutionary Biology</i> , 2012, 12, 40.	3.2	106
3	Dispersal ability rather than ecological tolerance drives differences in range size between lentic and lotic water beetles (Coleoptera: Hydrophilidae). <i>Journal of Biogeography</i> , 2012, 39, 984-994.	1.4	94
4	Phylogenetic community ecology of soil biodiversity using mitochondrial metagenomics. <i>Molecular Ecology</i> , 2015, 24, 3603-3617.	2.0	93
5	Metabarcoding and mitochondrial metagenomics of endogean arthropods to unveil the mesofauna of the soil. <i>Methods in Ecology and Evolution</i> , 2016, 7, 1071-1081.	2.2	75
6	Lessons from genome skimming of arthropods—preserving ethanol. <i>Molecular Ecology Resources</i> , 2016, 16, 1365-1377.	2.2	59
7	The phylogeny of leaf beetles (Chrysomelidae) inferred from mitochondrial genomes. <i>Systematic Entomology</i> , 2020, 45, 188-204.	1.7	56
8	Metabarcoding of freshwater invertebrates to detect the effects of a pesticide spill. <i>Molecular Ecology</i> , 2018, 27, 146-166.	2.0	54
9	The limited spatial scale of dispersal in soil arthropods revealed with whole—community haplotype—level metabarcoding. <i>Molecular Ecology</i> , 2021, 30, 48-61.	2.0	49
10	Mitochondrial Metagenomics Reveals the Ancient Origin and Phylodiversity of Soil Mites and Provides a Phylogeny of the Acari. <i>Molecular Biology and Evolution</i> , 2020, 37, 683-694.	3.5	42
11	The contribution of mitochondrial metagenomics to large-scale data mining and phylogenetic analysis of Coleoptera. <i>Molecular Phylogenetics and Evolution</i> , 2018, 128, 1-11.	1.2	41
12	Late Miocene origin of an Ibero—Maghrebian clade of ground beetles with multiple colonizations of the subterranean environment. <i>Journal of Biogeography</i> , 2014, 41, 1979-1990.	1.4	40
13	Integrative taxonomy and conservation of cryptic beetles in the Mediterranean region (Hydrophilidae). <i>Zoologica Scripta</i> , 2013, 42, 182-200.	0.7	34
14	Integration of conflict into integrative taxonomy: fitting hybridization in species delimitation of <i><scp>M</scp>esocarabus</i> (<scp>C</scp>oleoptera: <scp>C</scp>arabidae). <i>Molecular Ecology</i> , 2014, 23, 4344-4361.	2.0	33
15	Tempo and mode of the multiple origins of salinity tolerance in a water beetle lineage. <i>Molecular Ecology</i> , 2014, 23, 360-373.	2.0	32
16	Validated removal of nuclear pseudogenes and sequencing artefacts from mitochondrial metabarcode data. <i>Molecular Ecology Resources</i> , 2021, 21, 1772-1787.	2.2	32
17	Genome sequencing of <i>Rhinorhipus</i> Lawrence exposes an early branch of the Coleoptera. <i>Frontiers in Zoology</i> , 2018, 15, 21.	0.9	30
18	A validated workflow for rapid taxonomic assignment and monitoring of a national fauna of bees (Apiformes) using high throughput DNA barcoding. <i>Molecular Ecology Resources</i> , 2020, 20, 40-53.	2.2	30

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19	Climate drives community-wide divergence within species over a limited spatial scale: evidence from an oceanic island. <i>Ecology Letters</i> , 2020, 23, 305-315.	3.0	28
20	Connecting high-throughput biodiversity inventories: Opportunities for a site-based genomic framework for global integration and synthesis. <i>Molecular Ecology</i> , 2021, 30, 1120-1135.	2.0	26
21	Gondwanian relicts and oceanic dispersal in a cosmopolitan radiation of euedaphic ground beetles. <i>Molecular Phylogenetics and Evolution</i> , 2016, 99, 235-246.	1.2	25
22	Coming of age for COI metabarcoding of whole organism community DNA: Towards bioinformatic harmonisation. <i>Molecular Ecology Resources</i> , 2022, 22, 847-861.	2.2	22
23	Congruence test of molecular clock calibration hypotheses based on Bayesian factor comparisons. <i>Methods in Ecology and Evolution</i> , 2014, 5, 226-242.	2.2	21
24	Speciation below ground: Tempo and mode of diversification in a radiation of endogean ground beetles. <i>Molecular Ecology</i> , 2017, 26, 6053-6070.	2.0	17
25	Molecular systematics and evolution of the subgenus <i>Mesocarabus</i> Thomson, 1875 (Coleoptera: Carabidae). <i>Journal of Biogeography</i> , 2012, 39, 787-804.	1.0	16
26	The mitochondrial genome of <i>Iberobaenia</i> (Coleoptera: Iberobaeniidae): first rearrangement of protein-coding genes in the beetles. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2017, 28, 156-158.	0.7	16
27	<i>Terra incognita</i> of soil biodiversity: unseen invasions under our feet. <i>Molecular Ecology</i> , 2017, 26, 3087-3089.	2.0	16
28	Community metabarcoding reveals the relative role of environmental filtering and spatial processes in metacommunity dynamics of soil microarthropods across a mosaic of montane forests. <i>Molecular Ecology</i> , 2023, 32, 6110-6128.	2.0	15
29	Flightlessness in insects enhances diversification and determines assemblage structure across whole communities. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2021, 288, 20202646.	1.2	13
30	A new <i>Typhlocharis</i> (Coleoptera: Carabidae: Anillina) from Spain: combining adult and larval morphological data with DNA information. <i>Zootaxa</i> , 2010, 2485, 47.	0.2	11
31	The mitogenome of <i>Hydropsyche pellucidula</i> (Hydropsychidae): first gene arrangement in the insect order Trichoptera. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2017, 28, 71-72.	0.7	11
32	Hidden island endemic species and their implications for cryptic speciation within soil arthropods. <i>Journal of Biogeography</i> , 2022, 49, 1367-1380.	1.4	9
33	Community assembly and metaphylogeography of soil biodiversity: Insights from haplotype-level community DNA metabarcoding within an oceanic island. <i>Molecular Ecology</i> , 2022, 31, 4078-4094.	2.0	9
34	<i>Typhlocharis diecki</i> (Coleoptera: Carabidae, Anillini): a new species from the Iberian Peninsula, with notes about its relationships and the evolution of the diecki species group. <i>Zootaxa</i> , 2008, 1842, 35.	0.2	7
35	Dispersal limitations and long-term persistence drive differentiation from haplotypes to communities within a tropical sky-island: Evidence from community metabarcoding. <i>Molecular Ecology</i> , 2021, 30, 6611-6626.	2.0	6
36	Elemental composition, rare earths and minority elements in organic and conventional wines from volcanic areas: The Canary Islands (Spain). <i>PLoS ONE</i> , 2021, 16, e0258739.	1.1	6

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37	A new species of endogean, anophthalmous Parazuphium Jeannel from Northern Morocco (Coleoptera: Carabidae), with new molecular data for the tribe Zuphiini. ZooKeys, 2011, 103, 49-62.	0.5	4
38	Hidden biodiversity: total evidence phylogenetics and evolution of morphological traits in a highly diverse lineage of endogean ground beetles, <i>Typhlocharis</i> Dieck, 1869 (Carabidae, Trechinae). Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.2	4
39	New mitochondrial genomes of 39 soil dwelling Coleoptera from metagenome sequencing. Mitochondrial DNA Part B: Resources, 2019, 4, 2447-2450.	0.2	4
40	Endogean beetles (Coleoptera) of Madagascar: deep soil sampling and illustrated overview. Zootaxa, 2021, 4963, zootaxa.4963.2.4.	0.2	3
41	Mitogenomic phylogenetics of <i>Diochus occultus</i> n. sp., a palaeoendemic endogean species within the tribe Diochini (Coleoptera: Staphylinidae: Staphylininae). Journal of Zoological Systematics and Evolutionary Research, 2021, 59, 78-93.	0.6	2
42	Oromia orahan (Curculionidae, Molytinae), a new subterranean species for the Canarian underground biodiversity. Subterranean Biology, 0, 35, 1-14.	5.0	2
43	The discovery of Barretonus (Curculionidae: Cossoninae) in the Canary Islands: barcoding, morphology and description of new species. Acta Entomologica Musei Nationalis Pragae, 2019, 59, 443-452.	0.5	2
44	DNA barcoding reveals new records of invasive terrestrial flatworms (Platyhelminthes, Tricladida,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.2	0