# Roger Vila

List of Publications by Year
in descending order



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|---|---|---|---|
| 122 papers | 5,458 citations | 87888 38 h-index | 98798 67 g-index |
| 137 all docs | 137 docs citations | 137 times ranked | 5468 citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | DNA Barcodes Combined with Multilocus Data of Representative Taxa Can Generate Reliable Higher-Level Phylogenies. Systematic Biology, 2022, 71, 382-395. | 5.6 | 35 |
| 2 | The isolated Erebia pandrose Apennine population is genetically unique and endangered by climate change. Insect Conservation and Diversity, 2022, 15, 136-148. | 3.0 | 18 |
| 3 | Hybridization fuelled diversification in Spialia butterflies. Molecular Ecology, 2022, , . | 3.9 | 6 |
| 4 | The genome sequence of the lesser marbled fritillary, Brenthis ino, and evidence for a segregating neo-Z chromosome. G3: Genes, Genomes, Genetics, 2022, 12, . | 1.8 | 8 |
| 5 | The worrying arrival of the invasive Asian needle ant Brachyponera chinensis in Europe (Hymenoptera:) Tj ETQq1 1 0.784314 rgBT /Over | 0.5 | 4 |
| 6 | Integrative Taxonomy Reveals a New Melitaea (Lepidoptera: Nymphalidae) Species Widely Distributed in the Iberian Peninsula. Insect Systematics and Diversity, 2022, 6, . | 1.7 | 4 |
| 7 | Integrative taxonomy reveals cryptic diversity in North American Lasius ants, and an overlooked introduced species. Scientific Reports, 2022, 12, 5970. | 3.3 | 8 |
| 8 | Genetic assessment and climate modelling of the Iberian specialist butterfly Euchloe bazae (Lepidoptera: Pieridae). Insect Conservation and Diversity, 2022, 15, 594-605. | 3.0 | 2 |
| 9 | Erratic spatiotemporal vegetation growth anomalies drive population outbreaks in a trans-Saharan insect migrant. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2121249119. | 7.1 | 9 |
| 10 | Delimiting continuity: Comparison of target enrichment and double digest restrictionâ€site associated DNA sequencing for delineating admixing parapatric Melitaea butterflies. Systematic Entomology, 2022, 47, 637-654. | 3.9 | 2 |
| 11 | Linking largeâ€scale genetic structure of three Argynnini butterfly species to geography and environment. Molecular Ecology, 2022, 31, 4381-4401. | 3.9 | 7 |
| 12 | Host plant diet affects growth and induces altered gene expression and microbiome composition in the wood white ( Leptidea sinapis ) butterfly. Molecular Ecology, 2021, 30, 499-516. | 3.9 | 17 |
| 13 | Genetics and extreme confinement of three overlooked butterfly species in Romania call for immediate conservation actions. Journal of Insect Conservation, 2021, 25, 137-146. | 1.4 | 2 |
| 14 | Overlooked cryptic diversity inMuschampia(Lepidoptera: Hesperiidae) adds two species to the European butterfly fauna. Zoological Journal of the Linnean Society, 2021, 193, 847-859. | 2.3 | 9 |
| 15 | Two ways to be endemic. Alps and Apennines are different functional refugia during climatic cycles. Molecular Ecology, 2021, 30, 1297-1310. | 3.9 | 27 |
| 16 | High resolution DNA barcode library for European butterflies reveals continental patterns of mitochondrial genetic diversity. Communications Biology, 2021, 4, 315. | 4.4 | 57 |
| 17 | The Pleistocene species pump past its prime: Evidence from European butterfly sister species. Molecular Ecology, 2021, 30, 3575-3589. | 3.9 | 35 |
| 18 | The genome sequence of the large tortoiseshell, Nymphalis polychloros (Linnaeus, 1758). Wellcome Open Research, 2021, 6, 238. | 1.8 | 3 |

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|---|---|---|---|
| 19 | Conserved ancestral tropical niche but different continental histories explain the latitudinal diversity gradient in brush-footed butterflies. Nature Communications, 2021, 12, 5717. | 12.8 | 33 |
| 20 | The first known riodinid â€˜cuckooâ€™ butterfly reveals deep-time convergence and parallelism in ant social parasites. Zoological Journal of the Linnean Society, 2021, 193, 860-879. | 2.3 | 5 |
| 21 | The genome sequence of the heath fritillary, Melitaea athalia (Rottemburg, 1775). Wellcome Open Research, 2021, 6, 304. | 1.8 | 1 |
| 22 | Genomics Reveal Admixture and Unexpected Patterns of Diversity in a Parapatric Pair of Butterflies. Genes, 2021, 12, 2009. | 2.4 | 5 |
| 23 | Assigning occurrence data to cryptic taxa improves climatic niche assessments: Biodecrypt, a new tool tested on European butterflies. Global Ecology and Biogeography, 2020, 29, 1852-1865. | 5.8 | 11 |
| 24 | Incomplete Sterility of Chromosomal Hybrids: Implications for Karyotype Evolution and Homoploid Hybrid Speciation. Frontiers in Genetics, 2020, 11, 583827. | 2.3 | 24 |
| 25 | Rapid colour shift by reproductive character displacement in Cupido butterflies. Molecular Ecology, 2020, 29, 4942-4955. | 3.9 | 10 |
| 26 | Evolution of multiple sex-chromosomes associated with dynamic genome reshuffling in Leptidea wood-white butterflies. Heredity, 2020, 125, 138-154. | 2.6 | 35 |
| 27 | Integrative biodiversity inventory of ants from a Sicilian archipelago reveals high diversity on young volcanic islands (Hymenoptera: Formicidae). Organisms Diversity and Evolution, 2020, 20, 405-416. | 1.6 | 10 |
| 28 | How long is 3Âkm for a butterfly? Ecological constraints and functional traits explain high mitochondrial genetic diversity between Sicily and the Italian Peninsula. Journal of Animal Ecology, 2020, 89, 2013-2026. | 2.8 | 29 |
| 29 | Integrative analyses on Western PalearcticLasiommatareveal a mosaic of nascent butterfly species. Journal of Zoological Systematics and Evolutionary Research, 2020, 58, 809-822. | 1.4 | 12 |
| 30 | The determinants of genetic diversity in butterflies. Nature Communications, 2019, 10, 3466. | 12.8 | 80 |
| 31 | Integrating three comprehensive data sets shows that mitochondrial DNA variation is linked to species traits and paleogeographic events in European butterflies. Molecular Ecology Resources, 2019, 19, 1623-1636. | 4.8 | 66 |
| 32 | Lack of gene flow: Narrow and dispersed differentiation islands in a triplet of Leptidea butterfly species. Molecular Ecology, 2019, 28, 3756-3770. | 3.9 | 31 |
| 33 | A mirage of cryptic species: Genomics uncover striking mitonuclear discordance in the butterfly Thymelicus sylvestris. Molecular Ecology, 2019, 28, 3857-3868. | 3.9 | 75 |
| 34 | Dissecting the Effects of Selection and Mutation on Genetic Diversity in Three Wood White (Leptidea) Butterfly Species. Genome Biology and Evolution, 2019, 11, 2875-2886. | 2.5 | 18 |
| 35 | Exploitation of the invasive Asian HornetVespa velutinaby the European Honey BuzzardPernis apivorus. Bird Study, 2019, 66, 425-429. | 1.0 | 10 |
| 36 | The conundrum of species delimitation: a genomic perspective on a mitogenetically super-variable butterfly. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20191311. | 2.6 | 37 |

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|---|---|---|---|
| 37 | Global invasion history of the agricultural pest butterfly Pieris rapae revealed with genomics and citizen science. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 20015-20024. | 7.1 | 70 |
| 38 | Erebia epiphronandErebia orientalis: sibling butterfly species with contrasting histories. Biological Journal of the Linnean Society, 2019, 126, 338-348. | 1.6 | 12 |
| 39 | Two consecutive Wolbachiaâ€mediated mitochondrial introgressions obscure taxonomy in Palearctic swallowtail butterflies (Lepidoptera, Papilionidae). Zoologica Scripta, 2019, 48, 507-519. | 1.7 | 25 |
| 40 | Flight over the Proto-Caribbean seaway: Phylogeny and macroevolution of Neotropical Anaeini leafwing butterflies. Molecular Phylogenetics and Evolution, 2019, 137, 86-103. | 2.7 | 14 |
| 41 | Out of the Orient: Postâ€Tethyan transoceanic and transâ€Arabian routes fostered the spread of Baorini skippers in the Afrotropics. Systematic Entomology, 2019, 44, 926-938. | 3.9 | 16 |
| 42 | Bacterial communities within Phengaris (Maculinea) alcon caterpillars are shifted following transition from solitary living to social parasitism of Myrmica ant colonies. Ecology and Evolution, 2019, 9, 4452-4464. | 1.9 | 10 |
| 43 | Phenotypic biomarkers of climatic impacts on declining insect populations: A key role for decadal drought, thermal buffering and amplification effects and host plant dynamics. Journal of Animal Ecology, 2019, 88, 376-391. | 2.8 | 21 |
| 44 | Pollen metabarcoding as a tool for tracking longâ€distance insect migrations. Molecular Ecology Resources, 2019, 19, 149-162. | 4.8 | 52 |
| 45 | Use of genetic, climatic, and microbiological data to inform reintroduction of a regionally extinct butterfly. Conservation Biology, 2018, 32, 828-837. | 4.7 | 26 |
| 46 | A Comprehensive and Dated Phylogenomic Analysis of Butterflies. Current Biology, 2018, 28, 770-778.e5. | 3.9 | 249 |
| 47 | Gene expression profiling across ontogenetic stages in the wood white (Leptidea sinapis) reveals pathways linked to butterfly diapause regulation. Molecular Ecology, 2018, 27, 935-948. | 3.9 | 16 |
| 48 | An updated checklist of the European Butterflies (Lepidoptera, Papilionoidea). ZooKeys, 2018, 811, 9-45. | 1.1 | 90 |
| 49 | Versatility of multivalent orientation, inverted meiosis, and rescued fitness in holocentric chromosomal hybrids. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E9610-E9619. | 7.1 | 62 |
| 50 | Molecular evidence of hybridization in sympatric populations of the Enantia jethys complex (Lepidoptera: Pieridae). PLoS ONE, 2018, 13, e0197116. | 2.5 | 10 |
| 51 | Ecological specialization is associated with genetic structure in the ant-associated butterfly family Lycaenidae. Proceedings of the Royal Society B: Biological Sciences, 2018, 285, 20181158. | 2.6 | 9 |
| 52 | Historical and current patterns of gene flow in the butterfly Pararge aegeria. Journal of Biogeography, 2018, 45, 1628-1639. | 3.0 | 18 |
| 53 | Do Holarctic ant species exist? Transâ€Beringian dispersal and homoplasy in the Formicidae. Journal of Biogeography, 2018, 45, 1917-1928. | 3.0 | 33 |
| 54 | Round-trip across the Sahara: Afrotropical Painted Lady butterflies recolonize the Mediterranean in early spring. Biology Letters, 2018, 14, 20180274. | 2.3 | 34 |

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| 55 | Natural history and immature stage morphology of Spialia Swinhoe, 1912 in the Iberian Peninsula (Lepidoptera, Hesperiidae). Nota Lepidopterologica, 2018, 41, 1-22. | 0.6 | 5 |
| 56 | DNA Barcoding of an Assembly of Montane Andean Butterflies (Satyrinae): Geographical Scale and Identification Performance. Neotropical Entomology, 2017, 46, 514-523. | 1.2 | 13 |
| 57 | Climatic niche evolution is faster in sympatric than allopatric lineages of the butterfly genus Pyrgus. Proceedings of the Royal Society B: Biological Sciences, 2017, 284, 20170208. | 2.6 | 21 |
| 58 | Molecular substitution rate increases with latitude in butterflies: evidence for a transâ€glacial latitudinal layering of populations?. Ecography, 2017, 40, 930-935. | 4.5 | 4 |
| 59 | Genomics of extreme ecological specialists: multiple convergent evolution but no genetic divergence between ecotypes of Maculinea alcon butterflies. Scientific Reports, 2017, 7, 13752. | 3.3 | 13 |
| 60 | Rise and fall of island butterfly diversity: Understanding genetic differentiation and extinction in a highly diverse archipelago. Diversity and Distributions, 2017, 23, 1169-1181. | 4.1 | 32 |
| 61 | Differentiation in the marbled white butterfly species complex driven by multiple evolutionary forces. Journal of Biogeography, 2017, 44, 433-445. | 3.0 | 16 |
| 62 | Rapid Increase in Genome Size as a Consequence of Transposable Element Hyperactivity in Wood-White (Leptidea) Butterflies. Genome Biology and Evolution, 2017, 9, 2491-2505. | 2.5 | 94 |
| 63 | Oneâ€note samba: the biogeographical history of the relict Brazilian butterfly Elkalyce cogina. Journal of Biogeography, 2016, 43, 727-737. | 3.0 | 5 |
| 64 | Long-distance autumn migration across the Sahara by painted lady butterflies: exploiting resource pulses in the tropical savannah. Biology Letters, 2016, 12, 20160561. | 2.3 | 54 |
| 65 | Asymmetric constraints on limits to species ranges influence consumerâ€resource richness over an environmental gradient. Global Ecology and Biogeography, 2016, 25, 1477-1488. | 5.8 | 12 |
| 66 | Discovery of mass migration and breeding of the painted lady butterflyVanessa carduiin the Sub-Sahara: the Europe-Africa migration revisited. Biological Journal of the Linnean Society, 2016, , . | 1.6 | 19 |
| 67 | Integrative analyses unveil speciation linked to host plant shift in <scp>S</scp>pialia butterflies. Molecular Ecology, 2016, 25, 4267-4284. | 3.9 | 44 |
| 68 | Historical and contemporary factors generate unique butterfly communities on islands. Scientific Reports, 2016, 6, 28828. | 3.3 | 29 |
| 69 | Species-Level Para- and Polyphyly in DNA Barcode Gene Trees: Strong Operational Bias in European Lepidoptera. Systematic Biology, 2016, 65, 1024-1040. | 5.6 | 160 |
| 70 | The sibling species Leptidea juvernica and L. sinapis (Lepidoptera, Pieridae) in the Balkan Peninsula: ecology, genetic structure, and morphological variation. Zoology, 2016, 119, 11-20. | 1.2 | 6 |
| 71 | Why Do Cryptic Species Tend Not to Co-Occur? A Case Study on Two Cryptic Pairs of Butterflies. PLoS ONE, 2015, 10, e0117802. | 2.5 | 63 |
| 72 | Dynamic karyotype evolution and unique sex determination systems in Leptidea wood white butterflies. BMC Evolutionary Biology, 2015, 15, 89. | 3.2 | 51 |

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| 73 | DNA barcode reference library for Iberian butterflies enables a continental-scale preview of potential cryptic diversity. Scientific Reports, 2015, 5, 12395. | 3.3 | 110 |
| 74 | Ancient Neotropical origin and recent recolonisation: Phylogeny, biogeography and diversification of the Riodinidae (Lepidoptera: Papilionoidea). Molecular Phylogenetics and Evolution, 2015, 93, 296-306. | 2.7 | 72 |
| 75 | Cryptic matters: overlooked species generate most butterfly betaâ€diversity. Ecography, 2015, 38, 405-409. | 4.5 | 62 |
| 76 | Discovered just before extinction? The first endemic ant from the Balearic Islands ( Lasius balearicus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 | 3.0 | 18 |
| 77 | Biogeography of western Mediterranean butterflies: combining turnover and nestedness components of faunal dissimilarity. Journal of Biogeography, 2014, 41, 1639-1650. | 3.0 | 45 |
| 78 | Morphological and chemical analysis of male scent organs in the butterfly genus Pyrgus (Lepidoptera:) Tj ETQq0 0 0 rgBT /Overlock 10 T | 1.6 | 13 |
| 79 | Comparing population patterns for genetic and morphological markers with uneven sample sizes. An example for the butterfly Maniola jurtina. Methods in Ecology and Evolution, 2014, 5, 834-843. | 5.2 | 9 |
| 80 | A unified framework for diversity gradients: the adaptive trait continuum. Global Ecology and Biogeography, 2013, 22, 6-18. | 5.8 | 41 |
| 81 | recluster: an unbiased clustering procedure for betaâ€diversity turnover. Ecography, 2013, 36, 1070-1075. | 4.5 | 71 |
| 82 | Reproductive isolation and patterns of genetic differentiation in a cryptic butterfly species complex. Journal of Evolutionary Biology, 2013, 26, 2095-2106. | 1.7 | 60 |
| 83 | Factors affecting species delimitations with the <scp>GMYC</scp> model: insights from a butterfly survey. Methods in Ecology and Evolution, 2013, 4, 1101-1110. | 5.2 | 271 |
| 84 | In the shadow of phylogenetic uncertainty: The recent diversification of Lysandra butterflies through chromosomal change. Molecular Phylogenetics and Evolution, 2013, 69, 469-478. | 2.7 | 81 |
| 85 | Dispersal, fragmentation, and isolation shape the phylogeography of the European lineages ofPolyommatus(Agrodiaetus)ripartii(Lepidoptera: Lycaenidae). Biological Journal of the Linnean Society, 2013, 109, 817-829. | 1.6 | 23 |
| 86 | Biogeography and systematics of Aricia butterflies (Lepidoptera, Lycaenidae). Molecular Phylogenetics and Evolution, 2013, 66, 369-379. | 2.7 | 16 |
| 87 | Establishing criteria for higherâ€level classification using molecular data: the systematics of Polyommatus blue butterflies (Lepidoptera, Lycaenidae). Cladistics, 2013, 29, 166-192. | 3.3 | 84 |
| 88 | Ithomiini Butterflies (Lepidoptera: Nymphalidae) of Antioquia, Colombia. Neotropical Entomology, 2013, 42, 146-157. | 1.2 | 2 |
| 89 | Identifying zones of phenetic compression in West Mediterranean butterflies (Satyrinae): refugia, invasion and hybridization. Diversity and Distributions, 2012, 18, 1066-1076. | 4.1 | 21 |
| 90 | Unexpected layers of cryptic diversity in wood white Leptidea butterflies. Nature Communications, 2011, 2, 324. | 12.8 | 131 |

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|---|---|---|---|
| 91 | Phylogeny and palaeoecology of Polyommatus blue butterflies show Beringia was a climate-regulated gateway to the New World. Proceedings of the Royal Society B: Biological Sciences, 2011, 278, 2737-2744. | 2.6 | 98 |
| 92 | Phylogenetic island disequilibrium: evidence for ongoing long-term population dynamics in two Mediterranean butterflies. Journal of Biogeography, 2011, 38, 854-867. | 3.0 | 18 |
| 93 | Tracing the origin of disjunct distributions: a case of biogeographical convergence in Pyrgus butterflies. Journal of Biogeography, 2011, 38, 2006-2020. | 3.0 | 3 |
| 94 | A combined geneticâ€morphometric analysis unravels the complex biogeographical history of Polyommatus icarus and Polyommatus celina Common Blue butterflies. Molecular Ecology, 2011, 20, 3921-3935. | 3.9 | 62 |
| 95 | Unprecedented within-species chromosome number cline in the Wood White butterfly Leptidea sinapis and its significance for karyotype evolution and speciation. BMC Evolutionary Biology, 2011, 11, 109. | 3.2 | 74 |
| 96 | What is the phylogenetic signal limit from mitogenomes? The reconciliation between mitochondrial and nuclear data in the Insecta class phylogeny. BMC Evolutionary Biology, 2011, 11, 315. | 3.2 | 87 |
| 97 | A phylogenetic revision of the Glaucopsyche section (Lepidoptera: Lycaenidae), with special focus on the Phengarisâ€“Maculinea clade. Molecular Phylogenetics and Evolution, 2011, 61, 237-243. | 2.7 | 33 |
| 98 | Complete DNA barcode reference library for a country's butterfly fauna reveals high performance for temperate Europe. Proceedings of the Royal Society B: Biological Sciences, 2011, 278, 347-355. | 2.6 | 135 |
| 99 | How common are dot-like distributions? Taxonomical oversplitting in western European Agrodiaetus (Lepidoptera: Lycaenidae) revealed by chromosomal and molecular markers. Biological Journal of the Linnean Society, 2010, 101, 130-154. | 1.6 | 43 |
| 100 | The uneven phylogeny and biogeography of Erodium (Geraniaceae): radiations in the Mediterranean and recent recurrent intercontinental colonization. Annals of Botany, 2010, 106, 871-884. | 2.9 | 55 |
| 101 | Biogeography, ecology and conservation ofErebia oeme(HÃ¼bner) in the Carpathians (Lepidoptera: Nymphalidae: Satyrinae). Annales De La Societe Entomologique De France, 2010, 46, 486-498. | 0.9 | 8 |
| 102 | An inducible helix-Gly-Gly-helix motif in the N-terminal domain of histone H1e: A CD and NMR study. Protein Science, 2009, 11, 214-220. | 7.6 | 28 |
| 103 | Phylogeny and historical biogeography of the subtribe Aporiina (Lepidoptera: Pieridae): implications for the origin of Australian butterflies. Biological Journal of the Linnean Society, 2007, 90, 413-440. | 1.6 | 26 |
| 104 | Phylogeny of the Ants: Diversification in the Age of Angiosperms. Science, 2006, 312, 101-104. | 12.6 | 684 |
| 105 | Molecular phylogeny and systematics of the Pieridae (Lepidoptera: Papilionoidea): higher classification and biogeography. Zoological Journal of the Linnean Society, 2006, 147, 239-275. | 2.3 | 138 |
| 106 | Molecular phylogeny and systematics of the Pieridae (Lepidoptera: Papilionoidea): higher classification and biogeography. Zoological Journal of the Linnean Society, 2006, 147, 417-417. | 2.3 | 9 |
| 107 | Rearrangement of the Agrodiaetus dolus species group (Lepidoptera, Lycaenidae) using a new cytological approach and molecular data. Insect Systematics and Evolution, 2006, 37, 325-334. | 0.7 | 36 |
| 108 | Synergistic effects of combining morphological and molecular data in resolving the phylogeny of butterflies and skippers. Proceedings of the Royal Society B: Biological Sciences, 2005, 272, 1577-1586. | 2.6 | 228 |

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|---|---|---|---|
| 109 | The evolution of alternative parasitic life histories in large blue butterflies. Nature, 2004, 432, 386-390. | 27.8 | 163 |
| 110 | Sequence Complexity of Histone H1 Subtypes. Molecular Biology and Evolution, 2003, 20, 371-380. | 8.9 | 35 |
| 111 | DNA-induced Î±-Helical Structure in the NH2-terminal Domain of Histone H1. Journal of Biological Chemistry, 2001, 276, 46429-46435. | 3.4 | 57 |
| 112 | Induction of Secondary Structure in a COOH-terminal Peptide of Histone H1 by Interaction with the DNA. Journal of Biological Chemistry, 2001, 276, 30898-30903. | 3.4 | 63 |
| 113 | A helixâ€turn motif in the Câ€terminal domain of histone H1. Protein Science, 2000, 9, 627-636. | 7.6 | 38 |
| 114 | The genome sequence of the small tortoiseshell butterfly, Aglais urticae (Linnaeus, 1758). Wellcome Open Research, 0, 6, 233. | 1.8 | 4 |
| 115 | The genome sequence of the green-underside blue, Glaucopsyche alexis (Poda, 1761). Wellcome Open Research, 0, 6, 274. | 1.8 | 2 |
| 116 | The genome sequence of the Glanville fritillary, Melitaea cinxia (Linnaeus, 1758). Wellcome Open Research, 0, 6, 266. | 1.8 | 1 |
| 117 | The genome sequence of the small white, Pieris rapae (Linnaeus, 1758). Wellcome Open Research, 0, 6, 273. | 1.8 | 2 |
| 118 | The genome sequence of the European peacock butterfly, Aglais io (Linnaeus, 1758). Wellcome Open Research, 0, 6, 258. | 1.8 | 4 |
| 119 | An updated checklist of the European Butterflies (Lepidoptera, Papilionoidea). ZooKeys, 0, 811, 9-45. | 1.1 | 3 |
| 120 | The genome sequence of the small copper, Lycaena phlaeas (Linnaeus, 1760). Wellcome Open Research, 0, 6, 294. | 1.8 | 1 |
| 121 | The genome sequence of the grizzled skipper, Pyrgus malvae (Linnaeus, 1758). Wellcome Open Research, 0, 7, 114. | 1.8 | 0 |
| 122 | The genome sequence of the marbled white butterfly, Melanargia galathea (Linnaeus, 1758). Wellcome Open Research, 0, 7, 123. | 1.8 | 1 |