

# Manuel Ruiz-Garcia

## List of Publications by Year in descending order

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Version: 2024-02-01

73

papers

1,829

citations

361413

20

h-index

289244

40

g-index

79

all docs

79

docs citations

79

times ranked

2033

citing authors

| #  | ARTICLE  | IF  | CITATIONS |
|----|--|-----|-----------|
| 1  | Phylogeography of the capybara, <i>Hydrochoerus hydrochaeris</i> , in a large portion of its distribution area in South America. <i>Journal of Mammalian Evolution</i> , 2022, 29, 191-206.  | 1.8 | 2         |
| 2  | Out of the shadows: Multilocus systematics and biogeography of night monkeys suggest a Central Amazonian origin and a very recent widespread southeastward expansion in South America. <i>Molecular Phylogenetics and Evolution</i> , 2022, 170, 107426.   | 2.7 | 4         |
| 3  | Phylogenetics and an updated taxonomic status of the Tamarins (Callitrichinae, Cebidae). <i>Molecular Phylogenetics and Evolution</i> , 2022, 173, 107504.   | 2.7 | 9         |
| 4  | Effects of Sample Size in the Determination of the True Number of Haplogroups or ESUs Within a Species with Phylogeographic and Conservation Purposes: The Case of <i>Cebus albifrons</i> in Ecuador, and the Kinkajous and Coatis Throughout Latin America. , 2021, , 101-148.  |     | 0         |
| 5  | Comparative mitogenome phylogeography of two anteater genera (&lt;i&gt; <i>Tamandua</i> &lt;/i&gt; and) <i>T. j. ETQq1</i> 1 0.784314 rgBT /Overlock traits. <i>Zoological Research</i> , 2021, 42, 525-547.   | 2.1 | 6         |
| 6  | Mitochondrial and karyotypic evidence reveals a lack of support for the genus <i>Nasuella</i> (Procyonidae,) <i>T. j. ETQq0</i> 0 0 rgBT /Overlock 10 T  | 1.0 | 2         |
| 7  | Molecular Phylogenetics of <i>Bradypus</i> (Three-Toed Sloth, Pilosa: Bradypodidae, Mammalia) and Phylogeography of <i>Bradypus variegatus</i> (Brown-Throated Three-Toed Sloth) with Mitochondrial Gene Sequences. <i>Journal of Mammalian Evolution</i> , 2020, 27, 461-482.   | 1.8 | 4         |
| 8  | The phylogeographic structure of the mountain coati ( <i>Nasuella olivacea</i> ; Procyonidae, Carnivora), and its phylogenetic relationships with other coati species ( <i>Nasua nasua</i> and <i>Nasua narica</i> ) as inferred by mitochondrial DNA. <i>Mammalian Biology</i> , 2020, 100, 521-548.                            | 1.5 | 4         |
| 9  | The genetic structure of the spectacled bear ( <i>Tremarctos ornatus</i> ; Ursidae, Carnivora) in Colombia by means of mitochondrial and microsatellite markers. <i>Journal of Mammalogy</i> , 2020, 101, 1072-1090.   | 1.3 | 39        |
| 10 | Genetics of the Andean bear ( <i>Tremarctos ornatus</i> ; Ursidae, Carnivora) in Ecuador: when the Andean Cordilleras are not an Obstacle. Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2020, 31, 190-208.   | 0.7 | 36        |
| 11 | Invalidation of taxa within the silvery wooly monkey ( <i>Lagothrix lagothricha poeppigii</i> , Atelidae,) <i>T. j. ETQq1</i> 1 0.784314 rgBT /Overlock 10 T   | 0.7 | 10        |
| 12 | Molecular Evolution (Mitochondrial and Nuclear Microsatellites Markers) in the Andean Bear ( <i>Tremarctos ornatus</i> ; Ursidae, Carnivora): How Many ESUs Are There?. , 2020, , 165-194.   |     | 5         |
| 13 | Systematics, Evolution, and Genetics of Bears. , 2020, , 3-20.   |     | 0         |
| 14 | Andean Bear ( <i>Tremarctos ornatus</i> ). , 2020, , 78-87.  |     | 1         |
| 15 | Mitochondrial phylogeography of kinkajous (Procyonidae, Carnivora): maybe not a single ESU. <i>Journal of Mammalogy</i> , 2019, 100, 1631-1652.  | 1.3 | 9         |
| 16 | First Molecular Phylogenetic Analysis of the <i>Lagothrix</i> Taxon Living in Southern Peru and Northern Bolivia: <i>Lagothrix lagothricha tschudii</i> (Atelidae, Primates), a New Subspecies. <i>Folia Primatologica</i> , 2019, 90, 215-239.  | 0.7 | 8         |
| 17 | The mystery of the origins of <i>Cebus albifrons malitiosus</i> and <i>Cebus albifrons hypoleucus</i> : mitogenomics and microsatellite analyses revealed an amazing evolutionary history of the Northern Colombian white-fronted capuchins. Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis. 2019, 30, 525-547. | 0.7 | 2         |
| 18 | Mitogenomics phylogenetic relationships of the current slothâ€™s genera and species (Bradypodidae and) <i>T. j. ETQq0</i> 0 0 rgBT /Overlock 0.7   | 0.7 | 9         |

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 19 | Biodiversity in the Amazon: Origin Hypotheses, Intrinsic Capacity of Species Colonization, and Comparative Phylogeography of River Otters ( <i>Lontra longicaudis</i> and <i>Pteronura brasiliensis</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tff Evolution, 2018, 25, 213-240.   | 1.8 | 11        |
| 20 | Small spotted bodies with multiple specific mitochondrial DNAs: existence of diverse and differentiated tigrina lineages or species (<i>Leopardus</i>spp: Felidae, Mammalia) throughout Latin America. Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2018, 29, 993-1014.   | 0.7 | 11        |
| 21 | How Many Species, Taxa, or Lineages of <i>Cebus albifrons</i> (Platyrrhini, Primates) Inhabit Ecuador? Insights from Mitogenomics. International Journal of Primatology, 2018, 39, 1068-1104.   | 1.9 | 4         |
| 22 | Mitogenomics of the jaguarundi ( <i>Puma yagouaroundi</i> , Felidae, Carnivora): Disagreement between morphological subspecies and molecular data. Mammalian Biology, 2018, 93, 153-168.  | 1.5 | 8         |
| 23 | DNA Polymerase Sequences of New World Monkey Cytomegaloviruses: Another Molecular Marker with Which To Infer Platyrhini Systematics. Journal of Virology, 2018, 92, .   | 3.4 | 5         |
| 24 | Contrasting Rates of LINE-1 Amplification among New World Primates of the Atelidae Family. Cytogenetic and Genome Research, 2018, 154, 217-228.   | 1.1 | 5         |
| 25 | Phylogeography of the Mantled Howler Monkey ( <i>Alouatta palliata</i> ; Atelidae, Primates) across Its Geographical Range by Means of Mitochondrial Genetic Analyses and New Insights about the Phylogeny of <i>Alouatta</i> . Folia Primatologica, 2017, 88, 421-454.   | 0.7 | 11        |
| 26 | Continuous Miocene, Pliocene and Pleistocene Influences on Mitochondrial Diversification of the Capybara ( <i>Hydrochoerus hydrochaeris</i> ; Hydrochoeridae, Rodentia): Incapacity to Determine Exclusive Hypotheses on the Origins of the Amazon and Orinoco Diversity for This Species. Journal of Phylogenetics & Evolutionary Biology, 2016, 04, . | 0.2 | 7         |
| 27 | Phylogeography and spatial structure of the lowland tapir ( <i>Tapirus terrestris</i> , Perissodactyla) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tff 27, 2334-2342.  | 0.7 | 13        |
| 28 | Mitogenomics of the mountain tapir ( <i>Tapirus pinchaque</i> , Tapiridae, Perissodactyla, Mammalia) in Colombia and Ecuador: Phylogeography and insights into the origin and systematics of the South American tapirs. Mammalian Biology, 2016, 81, 163-175.   | 1.5 | 17        |
| 29 | Population genetics of the endangered Wattled Curassow ( <i>Crax globulosa</i> , Cracidae, Aves) of the Colombian-Peruvian Amazon using DNA microsatellites and ND2 mitochondrial sequences. Studies on Neotropical Fauna and Environment, 2015, 50, 80-95.   | 1.0 | 1         |
| 30 | Molecular phylogenetics and phylogeography of all the Saimiri taxa (Cebidae, Primates) inferred from mt COI and COII gene sequences. Primates, 2015, 56, 145-161.   | 1.1 | 13        |
| 31 | How many genera and species of woolly monkeys (Atelidae, Platyrrhine, Primates) are there? The first molecular analysis of <i>Lagothrix flavicauda</i> , an endemic Peruvian primate species. Molecular Phylogenetics and Evolution, 2014, 79, 179-198.   | 2.7 | 29        |
| 32 | Genetic characterization and structure of the endemic Colombian silvery brown bare-face tamarin, <i>Saguinus leucopus</i> (Callitrichinae, Cebidae, Primates). Primates, 2014, 55, 415-435.   | 1.1 | 10        |
| 33 | Molecular Identification and Historic Demography of the Marine Tucuxi ( <i>Sotalia guianensis</i> ) at the Amazon River's Mouth by Means of Mitochondrial Control Region Gene Sequences and Implications for Conservation. Diversity, 2013, 5, 703-723.   | 1.7 | 1         |
| 34 | Molecular Relationships and Classification of Several Tufted Capuchin Lineages (Cebus apella, Cebus) Tj ETQq0 0 0 rgBT /Overlock 10 Tff Sequences. Folia Primatologica, 2012, 83, 100-125.  | 0.7 | 24        |
| 35 | Pattern and timing of diversification of Cetartiodactyla (Mammalia, Laurasiatheria), as revealed by a comprehensive analysis of mitochondrial genomes. Comptes Rendus - Biologies, 2012, 335, 32-50.  | 0.2 | 448       |
| 36 | Molecular systematics and phylogeography of <i>Cebus capucinus</i> (Cebidae, Primates) in Colombia and Costa Rica by means of the mitochondrial COII gene. American Journal of Primatology, 2012, 74, 366-380.  | 1.7 | 25        |

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|----|---|-----|-----------|
| 37 | Sexual isolation between North American and Bogota strains of <i>Drosophila pseudoobscura</i> . <i>Behavior Genetics</i> , 2012, 42, 472-482.   | 2.1 | 8         |
| 38 | Population structure and conservation of a high-altitude specialist, the Andean cat <i>Leopardus jacobita</i> . <i>Endangered Species Research</i> , 2012, 16, 283-294.   | 2.4 | 19        |
| 39 | Partial molecular characterisation of New World non-human primate lymphocryptoviruses. <i>Infection, Genetics and Evolution</i> , 2011, 11, 1782-1789.  | 2.3 | 5         |
| 40 | Molecular Phylogenetics of <i>Aotus</i> (Platyrrhini, Cebidae). <i>International Journal of Primatology</i> , 2011, 32, 1218-1241.  | 1.9 | 26        |
| 41 | Population history, phylogeography, and conservation genetics of the last Neotropical mega-herbivore, the lowland tapir ( <i>Tapirus terrestris</i> ). <i>BMC Evolutionary Biology</i> , 2010, 10, 278.             | 3.2 | 41        |
| 42 | Phylogeny and phylogeography of squirrel monkeys (genus <i>Saimiri</i> ) based on cytochrome b genetic analysis. <i>American Journal of Primatology</i> , 2010, 72, 242-253.  | 1.7 | 41        |
| 43 | Molecular phylogenetics and phylogeography of the white-fronted capuchin ( <i>Cebus albifrons</i> ). Tj ETQq1 1 0.784314 rgBT /Overlock 1075, 57, 1049-1061.  | 2.7 | 35        |
| 44 | Genetic Structure of <i>Anopheles (Nyssorhynchus) marajoara</i> (Diptera: Culicidae) in Colombia. <i>American Journal of Tropical Medicine and Hygiene</i> , 2010, 83, 585-595.                                     | 1.4 | 13        |
| 45 | Molecular Systematics and Phylogeography of the Genus <i>Lagothrix</i> (Atelidae, Primates) by Means of the Mitochondrial COI Gene. <i>Folia Primatologica</i> , 2010, 81, 109-128.                                 | 0.7 | 31        |
| 46 | Analysis of diversity among six populations of Colombian mango ( <i>Mangifera indica L. cvar. Hilacha</i> ) using RAPDs markers. <i>Electronic Journal of Biotechnology</i> , 2009, 12, .                           | 2.2 | 6         |
| 47 | Influence of ancient glacial periods on the Andean fauna: the case of the pampas cat ( <i>Leopardus jacobita</i> ). Tj ETQq1 1 0.784314 rgBT /Overlock 1075, 57, 1049-1061.   | 3.2 | 41        |
| 48 | Isoenzyme Polymorphism and Genetic Structure of <i>Ochlerotatus taeniorhynchus</i> (Diptera: Culicidae) in Populations from the Colombian Atlantic Coast. <i>Biochemical Genetics</i> , 2009, 47, 462-470.          | 1.7 | 3         |
| 49 | The Taxonomy and Conservation Status of <i>Saimiri Sciureus Albigena</i> : A Squirrel Monkey Endemic to Colombia. <i>Primate Conservation</i> , 2009, 24, 59-64.  | 0.6 | 60        |
| 50 | Genetic Variability in Four <i>Alouatta</i> Species Measured by Means of Nine DNA Microsatellite Markers: Genetic Structure and Recent Bottlenecks. <i>Folia Primatologica</i> , 2007, 78, 73-87.                   | 0.7 | 25        |
| 51 | DNA microsatellite characterization of the jaguar ( <i>Panthera onca</i> ) in Colombia. <i>Genes and Genetic Systems</i> , 2006, 81, 115-127.   | 0.7 | 40        |
| 52 | Morphological analysis of <i>Threemnia</i> (Cetacea: Iniidae) populations from Colombia and Bolivia. <i>Acta Theriologica</i> , 2006, 51, 411-426.  | 1.1 | 10        |
| 53 | Microsatellite Analysis of the Spectacled Bear ( <i>Tremarctos ornatus</i> ) Across its Range Distribution. <i>Genes and Genetic Systems</i> , 2005, 80, 57-69.   | 0.7 | 62        |
| 54 | Population genetic analysis of cat populations from Mexico, Colombia, Bolivia, and the Dominican Republic: Identification of different gene pools in Latin America. <i>Journal of Genetics</i> , 2005, 84, 147-171. | 0.7 | 8         |

| #  | ARTICLE  | IF  | CITATIONS |
|----|--|-----|-----------|
| 55 | Mutant Allele Frequencies in Domestic Cat Populations in Arkansas and Tennessee. <i>Journal of Heredity</i> , 2005, 96, 557-565.   | 2.4 | 5         |
| 56 | COLONIZATION OF OCHLEROTATUS TAENIORHYNCHUS FROM RIOHACHA, COLOMBIA. <i>Journal of the American Mosquito Control Association</i> , 2005, 21, 28-32.  | 0.7 | 5         |
| 57 | Molecular population genetic analysis of the spectacled bear ( <i>Tremarctos ornatus</i> ) in the northern Andean area. <i>Hereditas</i> , 2003, 138, 81-93.   | 1.4 | 65        |
| 58 | Molecular systematics and biogeography of the Neotropical monkey genus, <i>Alouatta</i> . <i>Molecular Phylogenetics and Evolution</i> , 2003, 26, 64-81.  | 2.7 | 265       |
| 59 | RFLP Analysis of mtDNA from Six Platyrhine Genera: Phylogenetic Inferences. <i>Folia Primatologica</i> , 2003, 74, 59-70.  | 0.7 | 11        |
| 60 | Psorophora columbiae and Psorophora toltecum (Diptera: Culicidae) Colombian populations cannot be differentiated by Isoenzymes. <i>Genetics and Molecular Research</i> , 2003, 2, 229-59.  | 0.2 | 4         |
| 61 | Molecular Identification of Evolutionarily Significant Units in the Amazon River Dolphin <i>Inia</i> sp. (Cetacea: Iniidae). , 2002, 93, 312-322.  |     | 68        |
| 62 | Population genetic analysis of Colombian Trypanosoma cruzi isolates revealed by enzyme electrophoretic profiles. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2001, 96, 31-51.  | 1.6 | 1         |
| 63 | Genetic population history relationships of the population of BogotÃ¡, Colombia, by using the D1S80, VWA, and TH01 molecular markers. <i>American Journal of Human Biology</i> , 2001, 13, 374-383.  | 1.6 | 4         |
| 64 | Population Genetic Analysis of the Genes APOE, APOB (3â€²VNTR) and ACE in Some Black and Amerindian Communities from Colombia. <i>Human Heredity</i> , 2001, 52, 14-33.  | 0.8 | 25        |
| 65 | Genetic microstructure in two spanish cat populations. I. Genic diversity, gene flow and selection.. <i>Genes and Genetic Systems</i> , 2000, 75, 269-280.   | 0.7 | 14        |
| 66 | Genetic microsturture in two spanish cat populations. II. Gametic disequilibrium and spatial autocorrelation.. <i>Genes and Genetic Systems</i> , 2000, 75, 281-292.   | 0.7 | 6         |
| 67 | Lack of Evolutionary Divergence in Courtship Songs of <i>Drosophila pseudoobscura</i> Subspecies. <i>Journal of Insect Behavior</i> , 2000, 13, 255-262.   | 0.7 | 7         |
| 68 | Genetic structure of different cat populations in Europe and South America at a microgeographic level: importance of the choice of an adequate sampling level in the accuracy of population genetics interpretations. <i>Genetics and Molecular Biology</i> , 1999, 22, 493-505. | 1.3 | 10        |
| 69 | Genetic structure of different populations of domestic cat in Spain, Italy and Argentina at a micro-geographic level. <i>Acta Theriologica</i> , 1998, 43, 39-66.  | 1.1 | 11        |
| 70 | Genetic structure of populations of the domestic cat in Catalonia (Spain) and upper midwestern USA: A microgeographic and macrogeographic study. <i>Journal of Genetics</i> , 1997, 76, 99-115.  | 0.7 | 13        |
| 71 | Genetic relationships among some new cat populations sampled in Europe: A spatial autocorrelation analysis. <i>Journal of Genetics</i> , 1997, 76, 1-24.   | 0.7 | 32        |
| 72 | Mutant allele frequencies in domestic cat populations in Catalonia, Spain, and their genetic relationships with Spanish and English colonial cat populations. <i>Genetica</i> , 1990, 82, 209-214.   | 1.1 | 18        |

# ARTICLE

IF CITATIONS

73

Mitogenomic phylogenetics and population genetics of several taxa of agouties (*Dasyprocta* sp.,) Tj ETQq1 1 0.784314 rgBT /Overlock  
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