## Pedro Lorite

## List of Publications by Year in descending order

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Version: 2024-02-01

|                | 331670          | 377865                  |
|----------------|-----------------|-------------------------|
| 1,445          | 21              | 34                      |
| citations      | h-index         | g-index                 |
|                |                 |                         |
|                |                 |                         |
| 80             | <b>9</b> 0      | 1.46.1                  |
| 80             | <b>6</b> U      | 1461                    |
| docs citations | times ranked    | citing authors          |
|                |                 |                         |
|                | citations<br>80 | 1,445 21 h-index  80 80 |

| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | High chromosomal mobility of r <scp>DNA</scp> clusters in holocentric chromosomes of Triatominae, vectors of Chagas disease ( <scp>Hemipteraâ€Reduviidae</scp> ). Medical and Veterinary Entomology, 2022, 36, 66-80.   | 1.5  | 16        |
| 2  | Multidisciplinary approach detects speciation within the kissing bug Panstrongylus rufotuberculatus populations (Hemiptera, Heteroptera, Reduviidae). Memorias Do Instituto Oswaldo Cruz, 2022, 116, e210259.           | 1.6  | 4         |
| 3  | The Complete Nucleotide Sequence and Gene Organization of the Mitochondrial Genome of Triatoma boliviana (Hemiptera, Reduviidae, Triatominae) and Phylogenetic Comparisons., 2022, 1, 2-10.                             |      | 2         |
| 4  | Satellitome of the Red Palm Weevil, Rhynchophorus ferrugineus (Coleoptera: Curculionidae), the Most Diverse Among Insects. Frontiers in Ecology and Evolution, 2022, 10, .  | 2.2  | 15        |
| 5  | Complete mitochondrial genome of the blister beetle <i>Hycleus scutellatus</i> Rosenhauer, 1856 (Coleoptera, Meloidae). Mitochondrial DNA Part B: Resources, 2022, 7, 986-988.  | 0.4  | 2         |
| 6  | Characterization of New Molecular Markers of Three Botflies Parasitizing Cervid Hosts. Journal of Medical Entomology, 2021, 58, 1463-1469.  | 1.8  | 4         |
| 7  | Cytogenetic Analysis, Heterochromatin Characterization and Location of the rDNA Genes of Hycleus scutellatus (Coleoptera, Meloidae); A Species with an Unexpected High Number of rDNA Clusters. Insects, 2021, 12, 385. | 2.2  | 1         |
| 8  | Significance of PD1 Alternative Splicing in Celiac Disease as a Novel Source for Diagnostic and Therapeutic Target. Frontiers in Immunology, 2021, 12, 678400.  | 4.8  | 5         |
| 9  | Satellitome Analysis of Rhodnius prolixus, One of the Main Chagas Disease Vector Species.<br>International Journal of Molecular Sciences, 2021, 22, 6052.   | 4.1  | 19        |
| 10 | Chromosome Structure and Evolution of Triatominae: A Review. True Bugs (Heteroptera) of the Neotropics, 2021, , 65-99.  | 1.2  | 10        |
| 11 | Aphids and Ants, Mutualistic Species, Share a Mariner Element with an Unusual Location on Aphid Chromosomes. Genes, 2021, 12, 1966.   | 2.4  | 7         |
| 12 | Satellitome Analysis in the Ladybird Beetle Hippodamia variegata (Coleoptera, Coccinellidae). Genes, 2020, 11, 783.   | 2.4  | 18        |
| 13 | Complete Mitochondrial Genome of Three Species of the Genus Microtus (Arvicolinae, Rodentia).<br>Animals, 2020, 10, 2130.   | 2.3  | 7         |
| 14 | The complete mitochondrial genome of Talpa aquitania (Talpidae; Insectivora), a mole species endemic to northern Spain and southern France. Molecular Biology Reports, 2020, 47, 2397-2403.                             | 2.3  | 6         |
| 15 | Complex Evolutionary History of Mboumar, a Mariner Element Widely Represented in Ant Genomes. Scientific Reports, 2020, 10, 2610.   | 3.3  | 9         |
| 16 | Isolation of a Pericentromeric Satellite DNA Family in Chnootriba argus (Henosepilachna argus) with an Unusual Short Repeat Unit (TTAAAA) for Beetles. Insects, 2019, 10, 306.  | 2.2  | 3         |
| 17 | Dysregulation of the PD-1/PD-L1 pathway contributes to the pathogenesis of celiac disease. Cellular and Molecular Immunology, 2019, 16, 777-779.  | 10.5 | 10        |

Phylogenetic relationships between the slave-making ants<i>Rossomyrmex</i>and their<i>Proformica</i>hosts in relation to other genera of the ant tribe Formicini (Hymenoptera:) Tj ETQq0 0 0 rgBTL/Overlocks 10 Tf 50 5

| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 19 | Celiac Disease Autoimmunity. Archivum Immunologiae Et Therapiae Experimentalis, 2018, 66, 423-430.  | 2.3  | 30        |
| 20 | Comparative Analysis of Repetitive DNA between the Main Vectors of Chagas Disease: Triatoma infestans and Rhodnius prolixus. International Journal of Molecular Sciences, 2018, 19, 1277.                                   | 4.1  | 14        |
| 21 | Complete mitochondrial genome of the Iberian Mole Talpa occidentalis (Talpidae, Insectivora) and comparison with Talpa europaea. Genetica, 2018, 146, 415-423.  | 1.1  | 15        |
| 22 | Differentiating Iberoformica and Formica (Serviformica) with Description of the Sexual Castes of Formica (Serviformica) gerardi Bondroit, 1917 stat. rev Sociobiology, 2018, 65, 463.                                       | 0.5  | 2         |
| 23 | Potential role of the IL-33/ST2 axis in celiac disease. Cellular and Molecular Immunology, 2017, 14, 285-292.   | 10.5 | 23        |
| 24 | Concerted evolution, a slow process for ant satellite DNA: study of the satellite DNA in the Aphaenogaster genus (Hymenoptera, Formicidae). Organisms Diversity and Evolution, 2017, 17, 595-606.                           | 1.6  | 21        |
| 25 | Complete mitochondrial genome of Triatoma infestans (Hemiptera, Reduviidae, Triatominae), main vector of Chagas disease. Infection, Genetics and Evolution, 2017, 54, 158-163.  | 2.3  | 17        |
| 26 | Chromosome Painting in Triatomine Insects Reveals Shared Sequences Between X Chromosomes and Autosomes. Journal of Medical Entomology, 2017, 54, 44-49.   | 1.8  | 6         |
| 27 | Comparative repeatome analysis on Triatoma infestans Andean and Non-Andean lineages, main vector of Chagas disease. PLoS ONE, 2017, 12, e0181635.   | 2.5  | 46        |
| 28 | Holocentric chromosome evolution in kissing bugs (Hemiptera: Reduviidae: Triatominae): diversification of repeated sequences. Parasites and Vectors, 2017, 10, 410.   | 2.5  | 9         |
| 29 | Physiology and Pathology of Immune Dysregulation: Regulatory T Cells and Anergy. , 2017, , .  |      | 1         |
| 30 | New arrangements on several species subcomplexes of Triatoma genus based on the chromosomal position of ribosomal genes (Hemiptera - Triatominae). Infection, Genetics and Evolution, 2016, 43, 225-231.                    | 2.3  | 44        |
| 31 | The presence of the ancestral insect telomeric motif in kissing bugs (Triatominae) rules out the hypothesis of its loss in evolutionarily advanced Heteroptera (Cimicomorpha). Comparative Cytogenetics, 2016, 10, 427-437. | 0.8  | 16        |
| 32 | A PCR-RFLP method for detection of the LNPEP encoding human insulin-regulated aminopeptidase (IRAP) rs4869317 polymorphism. Indian Journal of Medical Research, 2016, 144, 120.   | 1.0  | 6         |
| 33 | USING COOPERATIVE LEARNING TO IMPROVE GENERIC SKILLS ACQUISITION IN UNIVERSITY STUDENTS. , 2016, ,  |      | 0         |
| 34 | Celiac Disease and Other Autoimmune Disorders. , 2015, , .  |      | 3         |
| 35 | Evolutionary history of the Azteca-like mariner transposons and their host ants. Die Naturwissenschaften, 2015, 102, 44.  | 1.6  | 7         |

Characterisation of an Iberian population of Rhyssocolpus iuventutis Andrássy, 1971 (Dorylaimida:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5

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|----|---|------|-----------|
| 37 | A step to the gigantic genome of the desert locust: chromosome sizes and repeated DNAs. Chromosoma, 2015, 124, 263-275.   | 2.2  | 53        |
| 38 | Recombination, chromosome number and eusociality in the Hymenoptera. Journal of Evolutionary Biology, 2015, 28, 105-116.  | 1.7  | 29        |
| 39 | Molecular cytogenetic studies in the ladybird beetle Henosepilachna argus Geoffroy, 1762 (Coleoptera, Coccinellidae, Epilachninae). Comparative Cytogenetics, 2015, 9, 423-434.   | 0.8  | 6         |
| 40 | Identification and In Vitro Reactivity of Celiac Immunoactive Peptides in an Apparent Gluten-Free Beer. PLoS ONE, 2014, 9, e100917.   | 2.5  | 32        |
| 41 | HLA in Gastrointestinal Inflammatory Disorders. , 2014, , .   |      | 2         |
| 42 | Distribution and Evolution of Repeated Sequences in Genomes of Triatominae (Hemiptera-Reduviidae) Inferred from Genomic In Situ Hybridization. PLoS ONE, 2014, 9, e114298.  | 2.5  | 20        |
| 43 | Plasma renin–angiotensin system-regulating aminopeptidase activities are modified in early stage Alzheimer's disease and show gender differences but are not related to apolipoprotein E genotype. Experimental Gerontology, 2013, 48, 557-564. | 2.8  | 20        |
| 44 | Characterization of two unrelated satellite DNA families in the Colorado potato beetle < i > Leptinotarsa decemlineata < / i > (Coleoptera, Chrysomelidae). Bulletin of Entomological Research, 2013, 103, 538-546.                             | 1.0  | 8         |
| 45 | Significant differences in coeliac immunotoxicity of barley varieties. Molecular Nutrition and Food Research, 2012, 56, 1697-1707.  | 3.3  | 35        |
| 46 | The ant genomes have been invaded by several types of mariner transposable elements. Die Naturwissenschaften, 2012, 99, 1007-1020.  | 1.6  | 14        |
| 47 | The spatial distribution does not affect host–parasite coevolution in Rossomyrmex ants. Insectes Sociaux, 2012, 59, 361-368.  | 1.2  | 4         |
| 48 | A new taxonomic status for Iberoformica (Hymenoptera, Formicidae) based on the use of molecular markers. Journal of Zoological Systematics and Evolutionary Research, 2012, 50, 30-37.  | 1.4  | 5         |
| 49 | Diversity in oat potential immunogenicity: basis for the selection of oat varieties with no toxicity in coeliac disease. Gut, 2011, 60, 915-922.  | 12.1 | 130       |
| 50 | Evaluation of HLAâ€G5 Plasmatic Levels During Pregnancy and Relationship with the 14â€bp Polymorphism. American Journal of Reproductive Immunology, 2010, 64, 367-374.  | 1.2  | 7         |
| 51 | 14–Base pair polymorphism of human leukocyte antigen–G as genetic determinant in heart transplantation and cyclosporine therapy monitoring. Human Immunology, 2009, 70, 830-835.  | 2.4  | 29        |
| 52 | Satellite DNA in insects: a review. Heredity, 2008, 100, 564-573.   | 2.6  | 114       |
| 53 | Transposition of Mboumar-9: Identification of a New Naturally Active mariner-Family Transposon.<br>Journal of Molecular Biology, 2008, 382, 567-572.  | 4.2  | 45        |
| 54 | A new approach using tissue alkaline phosphatase histochemistry to identify Crohn's disease. Pathology Research and Practice, 2007, 203, 485-487.   | 2.3  | 11        |

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|----|---|------------|----------------|
| 55 | Tryptophan metabolism and indoleamine 2,3-dioxygenase expression in coeliac disease. Clinical and Experimental Immunology, 2007, 148, 419-424.  | 2.6        | 55             |
| 56 | Detection of a mariner-like element and a miniature inverted-repeat transposable element (MITE) associated with the heterochromatin from ants of the genus Messor and their possible involvement for satellite DNA evolution. Gene, 2006, 371, 194-205. | 2.2        | 38             |
| 57 | Characterization and evolutionary dynamics of a complex family of satellite DNA in the leaf beetle Chrysolina carnifex (Coleoptera, Chrysomelidae). Chromosome Research, 2005, 13, 795-807.   | 2.2        | 30             |
| 58 | Expression of HLA-G in inflammatory bowel disease provides a potential way to distinguish between ulcerative colitis and Crohn's disease. International Immunology, 2004, 16, 579-583.  | 4.0        | 59             |
| 59 | Restriction Endonuclease Chromosome Banding in Tapinoma Nigerrimum (Hymenoptera, Formicidae)<br>Hereditas, 2004, 131, 197-201.  | 1.4        | 7              |
| 60 | A New Repetitive DNA Sequence Family in the Olive (Olea Europaea L.). Hereditas, 2004, 134, 73-78.  | 1.4        | 10             |
| 61 | Isolation and characterization of two families of satellite DNA with repetitive units of 135 bp and 2.5 kb in the ant <i>Monomorium subopacum</i> (Hymenoptera, Formicidae). Cytogenetic and Genome Research, 2004, 105, 83-92.                         | 1.1        | 18             |
| 62 | Evolutionary dynamics of satellite DNA in species of the Genus Formica (Hymenoptera, Formicidae). Gene, 2004, 332, 159-168.   | 2.2        | 27             |
| 63 | Conservation of (TTAGG)n Telomeric Sequences Among Ants (Hymenoptera, Formicidae). , 2002, 93, 282-285.   |            | 40             |
| 64 | Satellite DNA in the elm leaf beetle, <i>Xanthogaleruca luteola</i> (Coleoptera, Chrysomelidae): characterization, interpopulation analysis, and chromosome location. Cytogenetic and Genome Research, 2002, 98, 302-307.                               | 1,1        | 19             |
| 65 | Genomic organization and transcription of satellite DNA in the antAphaenogaster subterranea(Hymenoptera, Formicidae). Genome, 2002, 45, 609-616.  | 2.0        | 25             |
| 66 | Comparative study of satellite DNA in ants of the Messor genus. Gene, 2002, 297, 113-122.   | 2.2        | 25             |
| 67 | Characterization and chromosome location of satellite DNA in the leaf beetle Chrysolina americana (Coleoptera, Chrysomelidae). Genetica, 2000, 110, 143-150.  | 1.1        | 22             |
| 68 | Patterns of DNase I sensitivity in the chromosomes of the ant Tapinoma nigerrimum (Hymenoptera,) Tj ETQq0 0 (   | 0 rgBT /Ον | erlock 10 Tf 5 |
| 69 | Satellite DNA in the ant Messor structor (Hymenoptera, Formicidae). Genome, 1999, 42, 881-886.  | 2.0        | 6              |
| 70 | Satellite DNA in the ant <i>Messor structor </i> (Hymenoptera, Formicidae). Genome, 1999, 42, 881-886.  | 2.0        | 1              |
| 71 | Effects of restriction endonucleases on nucleolar organizing regions in the ant <i>Tapinoma nigerrimum</i> . Genome, 1998, 41, 872-875.   | 2.0        | 2              |
| 72 | Effects of restriction endonucleases on nucleolar organizing regions in the ant <i>Tapinoma nigerrimum &lt; <math>li</math> &gt;. Genome, 1998, 41, 872-875.</i>  | 2.0        | 2              |

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|----|--|----------------|-----------------------------|
| 73 | Analysis of the nucleolar organizing regions in the ant Tapinoma nigerrimum (Hymenoptera,) Tj ETQq1 1 0.7843   | 14.rgBT        | /Oveglock 10T               |
| 74 | Analysis of the nucleolar organizing regions in the ant Tapinoma nigerrimum (Hymenoptera,) Tj ETQq0 0 0 rgBT /   | Overloc<br>2.6 | k 10 <sub>7</sub> Tf 50 702 |
| 75 | Cytogenetic studies of antLinepithema humileShattuck (=Iridomyrmex humilisMayr) in European populations. Caryologia, 1996, 49, 199-205.  | 0.3            | 12                          |
| 76 | G-banding and chromosome condensation in the ant, Tapinoma nigerrimum. Chromosome Research, 1996, 4, 77-79.  | 2.2            | 17                          |
| 77 | Immune Checkpoints as a Novel Source for Diagnostic and Therapeutic Target in Celiac Disease. , 0, , .   |                | O                           |
| 78 | Characterization and transcriptional analysis of a subtelomeric satellite DNA family in the ladybird beetle Henosepilachna argus (Coleoptera: Coccinellidae). European Journal of Entomology, 0, 114, 481-487. | 1.2            | 4                           |
| 79 | Chromosome-level genome assembly and annotation of two lineages of the ant Cataglyphis hispanica: stepping stones towards genomic studies of hybridogenesis and thermal adaptation in desert ants., 0, 2, .    |                | 5                           |