Jacobo J Pardo-Seco

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6252401/publications.pdf

Version: 2024-02-01

59 1,455 22 34 g-index

63 63 63 63 2784

times ranked

citing authors

docs citations

all docs

| # | Article | lF | CITATIONS |
|----------|---|-----|-----------|
| 1 | CovidPhy: A tool for phylogeographic analysis of SARS-CoV-2 variation. Environmental Research, 2022, 204, 111909. | 3.7 | 5 |
| 2 | Evaluation of BNT162b2 Vaccine Effectiveness in Galicia, Northwest Spain. International Journal of Environmental Research and Public Health, 2022, 19, 4039. | 1.2 | 4 |
| 3 | A multi-tissue study of immune gene expression profiling highlights the key role of the nasal epithelium in COVID-19 severity. Environmental Research, 2022, 210, 112890. | 3.7 | 23 |
| 4 | Risk Analysis by Age on the Burden of Meningococcal Disease in Spain. Vaccines, 2022, 10, 592. | 2.1 | 4 |
| 5 | Pitfalls of barcodes in the study of worldwide SARS-CoV-2 variation and phylodynamics. Zoological Research, 2021, 42, 87-93. | 0.9 | 7 |
| 6 | BCG vaccination improves DTaP immune responses in mice and is associated with lower pertussis incidence in ecological epidemiological studies. EBioMedicine, 2021, 65, 103254. | 2.7 | 10 |
| 7 | Identification of a Minimal 3-Transcript Signature to Differentiate Viral from Bacterial Infection from Best Genome-Wide Host RNA Biomarkers: A Multi-Cohort Analysis. International Journal of Molecular Sciences, 2021, 22, 3148. | 1.8 | 6 |
| 8 | Case Report: Two Monochorionic Twins With a Critically Different Course of Progressive Osseous Heteroplasia. Frontiers in Pediatrics, 2021, 9, 662669. | 0.9 | 3 |
| 9 | Development and Evaluation of the Ancestry Informative Marker Panel of the VISAGE Basic Tool. Genes, 2021, 12, 1284. | 1.0 | 20 |
| 10 | Superspreading in the emergence of COVID-19 variants. Trends in Genetics, 2021, 37, 1069-1080. | 2.9 | 31 |
| 11 | Routine infant vaccination of pneumococcal conjugate vaccines has decreased pneumonia across all age groups in Northern Spain. Human Vaccines and Immunotherapeutics, 2020, 16, 1446-1453. | 1.4 | 5 |
| 12 | Seroprevalence of SARS-CoV-2 Among Pediatric Healthcare Workers in Spain. Frontiers in Pediatrics, 2020, 8, 547. | 0.9 | 19 |
| 13 | Mapping genome variation of SARS-CoV-2 worldwide highlights the impact of COVID-19 super-spreaders. Genome Research, 2020, 30, 1434-1448. | 2.4 | 91 |
| 14 | Phylogeography of SARS-CoV-2 pandemic in Spain: a story of multiple introductions, micro-geographic | | 34 |
| | stratification, founder effects, and super-spreaders. Zoológical Research, 2020, 41, 605-620. | 0.9 | |
| 15 | stratification, founder effects, and super-spreaders. Zoological Research, 2020, 41, 605-620. Rotavirus infection beyond the gut. Infection and Drug Resistance, 2019, Volume 12, 55-64. | 1.1 | 32 |
| 15 16 | stratification, founder effects, and super-spreaders. Zoological Research, 2020, 41, 605-620. | | |
| | Rotavirus infection beyond the gut. Infection and Drug Resistance, 2019, Volume 12, 55-64. A qPCR expression assay of IFI44L gene differentiates viral from bacterial infections in febrile | 1.1 | 32 |

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|----|---|-----------|-----------|
| 19 | Lifestyle and comorbid conditions as risk factors for community-acquired pneumonia in outpatient adults (NEUMO-ES-RISK project). BMJ Open Respiratory Research, 2019, 6, e000359. | 1.2 | 19 |
| 20 | Impact of rotavirus vaccination on childhood hospitalizations for seizures: Heterologous or unforeseen direct vaccine effects?. Vaccine, 2019, 37, 3362-3368. | 1.7 | 11 |
| 21 | Biogeographical origin and timing of the founder ichthyosis TGM1 c.1187G > A mutation in an isolate Ecuadorian population. Scientific Reports, 2019, 9, 7175. | ed 1.6 | 7 |
| 22 | Ancestry patterns inferred from massive RNA-seq data. Rna, 2019, 25, 857-868. | 1.6 | 16 |
| 23 | The geographic mosaic of Ecuadorian Y-chromosome ancestry. Forensic Science International: Genetics, 2018, 33, 59-65. | 1.6 | 19 |
| 24 | Whole Exome Sequencing Identifies New Host Genomic Susceptibility Factors in Empyema Caused by Streptococcus pneumoniae in Children: A Pilot Study. Genes, 2018, 9, 240. | 1.0 | 9 |
| 25 | Rotavirus intestinal infection induces an oral mucosa cytokine response. PLoS ONE, 2018, 13, e0195314. | 1.1 | 5 |
| 26 | A 2-transcript host cell signature distinguishes viral from bacterial diarrhea and it is influenced by the severity of symptoms. Scientific Reports, 2018, 8, 8043. | 1.6 | 20 |
| 27 | Life-threatening infections in children in Europe (the EUCLIDS Project): a prospective cohort study. The Lancet Child and Adolescent Health, 2018, 2, 404-414. | 2.7 | 69 |
| 28 | The peopling of South America and the trans-Andean gene flow of the first settlers. Genome Research, 2018, 28, 767-779. | 2.4 | 59 |
| 29 | Y-chromosome Peruvian origin of the 500-year-old Inca child mummy sacrificed in Cerro Aconcagua (Argentina). Science Bulletin, 2018, 63, 1457-1459. | 4.3 | 5 |
| 30 | Phylogenetic and population-based approaches to mitogenome variation do not support association with male infertility. Journal of Human Genetics, 2017, 62, 361-371. | 1.1 | 3 |
| 31 | Phylogeographic and genome-wide investigations of Vietnam ethnic groups reveal signatures of complex historical demographic movements. Scientific Reports, 2017, 7, 12630. | 1.6 | 17 |
| 32 | Clinical respiratory scales: which one should we use?. Expert Review of Respiratory Medicine, 2017, 11, 1-19. | 1.0 | 13 |
| 33 | Whole Exome Sequencing reveals new candidate genes in host genomic susceptibility to Respiratory Syncytial Virus Disease. Scientific Reports, 2017, 7, 15888. | 1.6 | 29 |
| 34 | Salivary epidermal growth factor correlates with hospitalization length in rotavirus infection. BMC Infectious Diseases, 2017, 17, 370. | 1.3 | 4 |
| 35 | Bacteremia in Children Hospitalized with Respiratory Syncytial Virus Infection. PLoS ONE, 2016, 11, e0146599. | 1.1 | 36 |
| 36 | Development and Validation of a New Clinical Scale for Infants with Acute Respiratory Infection: The ReSVinet Scale. PLoS ONE, 2016, 11, e0157665. | 1.1 | 41 |

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|----|---|-----|-----------|
| 37 | The Burden of Pediatric Invasive Meningococcal Disease in Spain (2008–2013). Pediatric Infectious Disease Journal, 2016, 35, 407-413. | 1.1 | 27 |
| 38 | Role of Vitamin D in Hospitalized Children With Lower Tract Acute Respiratory Infections. Journal of Pediatric Gastroenterology and Nutrition, 2016, 62, 479-485. | 0.9 | 12 |
| 39 | Natural resistance to Meningococcal Disease related to CFH loci: Meta-analysis of genome-wide association studies. Scientific Reports, 2016, 6, 35842. | 1.6 | 33 |
| 40 | Incidence and risk factor prevalence of community-acquired pneumonia in adults in primary care in Spain (NEUMO-ES-RISK project). BMC Infectious Diseases, 2016, 16, 645. | 1.3 | 64 |
| 41 | Genomic continuity of Argentinean Mennonites. Scientific Reports, 2016, 6, 36392. | 1.6 | 4 |
| 42 | Mapping the genomic mosaic of two â€~Afro-Bolivians' from the isolated Yungas valleys. BMC Genomics, 2016, 17, 207. | 1.2 | 9 |
| 43 | Charting the Y-chromosome ancestry of present-day Argentinean Mennonites. Journal of Human Genetics, 2016, 61, 507-513. | 1.1 | 10 |
| 44 | Analysis of Y-chromosome STRs in Chile confirms an extensive introgression of European male lineages in urban populations. Forensic Science International: Genetics, 2016, 21, 76-80. | 1.6 | 12 |
| 45 | A comprehensive Y-STR portrait of Argentinean populations. Forensic Science International: Genetics, 2016, 20, 1-5. | 1.6 | 9 |
| 46 | Does Viral Co-Infection Influence the Severity of Acute Respiratory Infection in Children?. PLoS ONE, 2016, 11, e0152481. | 1.1 | 46 |
| 47 | The complete mitogenome of a 500-year-old Inca child mummy. Scientific Reports, 2015, 5, 16462. | 1.6 | 31 |
| 48 | Impact of Rotavirus Vaccination on Childhood Hospitalization for Seizures. Pediatric Infectious Disease Journal, 2015, 34, 769-773. | 1.1 | 40 |
| 49 | Nonspecific (Heterologous) Protection of Neonatal BCG Vaccination Against Hospitalization Due to Respiratory Infection and Sepsis. Clinical Infectious Diseases, 2015, 60, 1611-1619. | 2.9 | 173 |
| 50 | The multiethnic ancestry of Bolivians as revealed by the analysis of Y-chromosome markers. Forensic Science International: Genetics, 2015, 14, 210-218. | 1.6 | 18 |
| 51 | Mitogenomes from The 1000 Genome Project Reveal New Near Eastern Features in Present-Day Tuscans. PLoS ONE, 2015, 10, e0119242. | 1.1 | 15 |
| 52 | The Genomic Legacy of the Transatlantic Slave Trade in the Yungas Valley of Bolivia. PLoS ONE, 2015, 10, e0134129. | 1.1 | 8 |
| 53 | Viral Co-Infections in Pediatric Patients Hospitalized with Lower Tract Acute Respiratory Infections. PLoS ONE, 2015, 10, e0136526. | 1.1 | 67 |
| 54 | Evaluating the accuracy of AIM panels at quantifying genome ancestry. BMC Genomics, 2014, 15, 543. | 1.2 | 29 |

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|----|--|-----|-----------|
| 55 | A Genome-Wide Study of Modern-Day Tuscans: Revisiting Herodotus's Theory on the Origin of the Etruscans. PLoS ONE, 2014, 9, e105920. | 1.1 | 23 |
| 56 | Ancestry analysis reveals a predominant Native American component with moderate European admixture in Bolivians. Forensic Science International: Genetics, 2013, 7, 537-542. | 1.6 | 26 |
| 57 | Indian Signatures in the Westernmost Edge of the European Romani Diaspora: New Insight from Mitogenomes. PLoS ONE, 2013, 8, e75397. | 1.1 | 24 |
| 58 | The Genetic Legacy of the Pre-Colonial Period in Contemporary Bolivians. PLoS ONE, 2013, 8, e58980. | 1.1 | 42 |
| 59 | A Generalized Model to Estimate the Statistical Power in Mitochondrial Disease Studies Involving 2×k Tables. PLoS ONE, 2013, 8, e73567. | 1.1 | 11 |