

Andrés López-Cortés

List of Publications by Year in descending order

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Version: 2024-02-01

83
papers

1,948
citations

361045

20
h-index

301761

39
g-index

103
all docs

103
docs citations

103
times ranked

3159
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of Key Proteins from the Alternative Lengthening of Telomeres-Associated Promyelocytic Leukemia Nuclear Bodies Pathway. <i>Biology</i> , 2022, 11, 185.	1.3	1
2	Pulmonary Inflammatory Response in Lethal COVID-19 Reveals Potential Therapeutic Targets and Drugs in Phases III/IV Clinical Trials. <i>Frontiers in Pharmacology</i> , 2022, 13, 833174.	1.6	6
3	Integrated In Silico Analyses Identify PUF60 and SF3A3 as New Spliceosome-Related Breast Cancer RNA-Binding Proteins. <i>Biology</i> , 2022, 11, 481.	1.3	3
4	Molecular Pathogenesis and New Therapeutic Dimensions for Spinal Muscular Atrophy. <i>Biology</i> , 2022, 11, 894.	1.3	1
5	Genetic Variations of the DPYD Gene and Its Relationship with Ancestry Proportions in Different Ecuadorian Trihybrid Populations. <i>Journal of Personalized Medicine</i> , 2022, 12, 950.	1.1	1
6	The close interaction between hypoxia-related proteins and metastasis in pancarcinomas. <i>Scientific Reports</i> , 2022, 12, .	1.6	4
7	Acute respiratory distress syndrome (ARDS) caused by the novel coronavirus disease (COVID-19): a practical comprehensive literature review. <i>Expert Review of Respiratory Medicine</i> , 2021, 15, 183-195.	1.0	36
8	Epidemiological, socio-demographic and clinical features of the early phase of the COVID-19 epidemic in Ecuador. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0008958.	1.3	94
9	In silico Analyses of Immune System Protein Interactome Network, Single-Cell RNA Sequencing of Human Tissues, and Artificial Neural Networks Reveal Potential Therapeutic Targets for Drug Repurposing Against COVID-19. <i>Frontiers in Pharmacology</i> , 2021, 12, 598925.	1.6	16
10	SARS-CoV-2 vaccines strategies: a comprehensive review of phase 3 candidates. <i>Npj Vaccines</i> , 2021, 6, 28.	2.9	507
11	A New Insight for the Identification of Oncogenic Variants in Breast and Prostate Cancers in Diverse Human Populations, With a Focus on Latinos. <i>Frontiers in Pharmacology</i> , 2021, 12, 630658.	1.6	3
12	Vaccine market and production capabilities in the Americas. <i>Tropical Diseases, Travel Medicine and Vaccines</i> , 2021, 7, 11.	0.9	13
13	15q Duplication Syndrome: Report on the First Patient from Ecuador with an Unusual Clinical Presentation. <i>Case Reports in Medicine</i> , 2021, 2021, 1-9.	0.3	1
14	Identification of key proteins in the signaling crossroads between wound healing and cancer hallmark phenotypes. <i>Scientific Reports</i> , 2021, 11, 17245.	1.6	7
15	Manifestaciones neurológicas del lupus eritematoso sistémico: Revisión de literatura.. <i>Revista Ecuatoriana De Neurología</i> , 2021, 30, 76-82.	0.1	1
16	Tracking SARS-CoV-2: Novel Trends and Diagnostic Strategies. <i>Diagnostics</i> , 2021, 11, 1981.	1.3	13
17	Editorial: Pharmacogenetics and Pharmacogenomics in Latin America: Ethnic Variability, New Insights in Advances and Perspectives: A RELIVAF-CYTED Initiative. <i>Frontiers in Pharmacology</i> , 2021, 12, 833000.	1.6	3
18	Title is missing!. , 2021, 15, e0008958.		0

#	ARTICLE	IF	CITATIONS
19	Title is missing!. , 2021, 15, e0008958.		0
20	Title is missing!. , 2021, 15, e0008958.		0
21	Title is missing!. , 2021, 15, e0008958.		0
22	Title is missing!. , 2021, 15, e0008958.		0
23	Title is missing!. , 2021, 15, e0008958.		0
24	Title is missing!. , 2021, 15, e0008958.		0
25	Title is missing!. , 2021, 15, e0008958.		0
26	Pharmacogenomics, biomarker network, and allele frequencies in colorectal cancer. Pharmacogenomics Journal, 2020, 20, 136-158.	0.9	15
27	Multi-institutional experience of genetic diagnosis in Ecuador: National registry of chromosome alterations and polymorphisms. Molecular Genetics & Genomic Medicine, 2020, 8, e1087.	0.6	3
28	Cytogenetic and genomic analysis of a patient with turner syndrome and t(2;12): a case report. Molecular Cytogenetics, 2020, 13, 46.	0.4	4
29	Drugs Repurposing Using QSAR, Docking and Molecular Dynamics for Possible Inhibitors of the SARS-CoV-2 Mpro Protease. Molecules, 2020, 25, 5172.	1.7	42
30	A Multi-Objective Approach for Anti-Osteosarcoma Cancer Agents Discovery through Drug Repurposing. Pharmaceuticals, 2020, 13, 409.	1.7	6
31	Oncology and Pharmacogenomics Insights in Polycystic Ovary Syndrome: An Integrative Analysis. Frontiers in Endocrinology, 2020, 11, 585130.	1.5	16
32	TCGA Pan-Cancer Genomic Analysis of Alternative Lengthening of Telomeres (ALT) Related Genes. Genes, 2020, 11, 834.	1.0	8
33	Clinical, genomics and networking analyses of a high-altitude native American Ecuadorian patient with congenital insensitivity to pain with anhidrosis: a case report. BMC Medical Genomics, 2020, 13, 113.	0.7	5
34	A deep analysis using panel-based next-generation sequencing in an Ecuadorian pediatric patient with anaplastic astrocytoma: a case report. Journal of Medical Case Reports, 2020, 14, 136.	0.4	1
35	Perturbation-Theory Machine Learning (PTML) Multilabel Model of the ChEMBL Dataset of Preclinical Assays for Antisarcoma Compounds. ACS Omega, 2020, 5, 27211-27220.	1.6	4
36	Clinical, molecular, and epidemiological characterization of the SARS-CoV-2 virus and the Coronavirus Disease 2019 (COVID-19), a comprehensive literature review. Diagnostic Microbiology and Infectious Disease, 2020, 98, 115094.	0.8	293

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37	OncoOmics approaches to reveal essential genes in breast cancer: a panoramic view from pathogenesis to precision medicine. <i>Scientific Reports</i> , 2020, 10, 5285.	1.6	36
38	Characterization of Ancestral Origin of Cystic Fibrosis of Patients with New Reported Mutations in CFTR. <i>BioMed Research International</i> , 2020, 2020, 1-6.	0.9	1
39	Gene Prioritization through Consensus Strategy, Enrichment Methodologies Analysis, and Networking for Osteosarcoma Pathogenesis. <i>International Journal of Molecular Sciences</i> , 2020, 21, 1053.	1.8	13
40	De Novo Duplication of Chromosome 9p in a Female Infant: Phenotype and Genotype Correlation. <i>Journal of Pediatric Genetics</i> , 2020, 09, 069-075.	0.3	3
41	Prediction of breast cancer proteins involved in immunotherapy, metastasis, and RNA-binding using molecular descriptors and artificial neural networks. <i>Scientific Reports</i> , 2020, 10, 8515.	1.6	29
42	Análisis del potencial genotípico y carcinógeno asociado a los cigarrillos electrónicos. <i>Revista Ecuatoriana De Medicina Y Ciencias Biológicas</i> , 2020, 41, .	0.1	0
43	Análisis del potencial genotípico y carcinógeno asociado a los cigarrillos electrónicos. <i>Revista Ecuatoriana De Medicina Y Ciencias Biológicas</i> , 2020, 41, .	0.1	0
44	Genetic variation of high-altitude Ecuadorian population using autosomal STR markers. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 62-64.	0.1	1
45	Allele frequency data for 15 autosomal strs and ancestral proportions using aims-indels in the shuar ethnic group from Ecuador. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 65-67.	0.1	0
46	Post-transcriptional Regulation of Colorectal Cancer: A Focus on RNA-Binding Proteins. <i>Frontiers in Molecular Biosciences</i> , 2019, 6, 65.	1.6	39
47	Ancestral analysis of a Native American Ecuadorian family with congenital insensitivity to pain with anhidrosis. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 126-128.	0.1	0
48	The three-hybrid genetic composition of an Ecuadorian population using AIMS-InDels compared with autosomes, mitochondrial DNA and Y chromosome data. <i>Scientific Reports</i> , 2019, 9, 9247.	1.6	31
49	A quick guide for using Microsoft OneNote as an electronic laboratory notebook. <i>PLoS Computational Biology</i> , 2019, 15, e1006918.	1.5	14
50	Genotoxic and Carcinogenic Potential of Compounds Associated with Electronic Cigarettes: A Systematic Review. <i>BioMed Research International</i> , 2019, 2019, 1-8.	0.9	15
51	Genes involved in damage response caused by UV radiation in Ecuadorian population of different altitude regions. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 140-141.	0.1	0
52	Mitochondrial DNA study in the Shuar ethnic group from Ecuador. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 142-143.	0.1	1
53	Molecular variants associated with flavor perceptions and ancestral proportions of Ecuadorian populations. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 59-61.	0.1	0
54	Breast Cancer Risk Associated with Genotype Polymorphisms of the Aurora Kinase a Gene (AURKA): a Case-Control Study in a High Altitude Ecuadorian Mestizo Population. <i>Pathology and Oncology Research</i> , 2018, 24, 457-465.	0.9	11

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55	Salivary MicroRNAs for Early Detection of Head and Neck Squamous Cell Carcinoma: A Case-Control Study in the High Altitude Mestizo Ecuadorian Population. <i>BioMed Research International</i> , 2018, 2018, 1-9.	0.9	38
56	Gene prioritization, communality analysis, networking and metabolic integrated pathway to better understand breast cancer pathogenesis. <i>Scientific Reports</i> , 2018, 8, 16679.	1.6	29
57	Analysis of Racial/Ethnic Representation in Select Basic and Applied Cancer Research Studies. <i>Scientific Reports</i> , 2018, 8, 13978.	1.6	105
58	Mutational Analysis of Oncogenic AKT1 Gene Associated with Breast Cancer Risk in the High Altitude Ecuadorian Mestizo Population. <i>BioMed Research International</i> , 2018, 2018, 1-10.	0.9	28
59	Study of the Huntington's disease <i>HTT</i> gene in different ethnic groups in Ecuador. <i>Clinical Genetics</i> , 2017, 92, 544-547.	1.0	8
60	Molecular analysis of ancestry informative markers (AIMs-INDELs) in a high altitude Ecuadorian mestizo population affected with breast cancer. <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e231-e232.	0.1	3
61	Ancestry study in Ecuadorian population with multiple myeloma. <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e435-e436.	0.1	0
62	Ancestry characterization of Ecuador's Highland mestizo population using autosomal AIM-INDELs. <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e477-e478.	0.1	10
63	Evaluation of ancestral membership proportions and genotype distribution in the perception of Umami taste in Ecuadorian mestizos. <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e171-e172.	0.1	1
64	Genotyping the High Altitude Mestizo Ecuadorian Population Affected with Prostate Cancer. <i>BioMed Research International</i> , 2017, 2017, 1-10.	0.9	7
65	State of Art of Cancer Pharmacogenomics in Latin American Populations. <i>International Journal of Molecular Sciences</i> , 2017, 18, 639.	1.8	25
66	A study of the molecular variants associated with lactase persistence in different Ecuadorian ethnic groups. <i>American Journal of Human Biology</i> , 2016, 28, 774-781.	0.8	7
67	Positive association of the androgen receptor CAG repeat length polymorphism with the risk of prostate cancer. <i>Molecular Medicine Reports</i> , 2016, 14, 1791-1798.	1.1	14
68	Positive Association of the Cathepsin D Ala224Val Gene Polymorphism With the Risk of Alzheimer's Disease. <i>American Journal of the Medical Sciences</i> , 2015, 350, 296-301.	0.4	24
69	Breast cancer risk associated with gene expression and genotype polymorphisms of the folate-metabolizing MTHFR gene: a case-control study in a high altitude Ecuadorian mestizo population. <i>Tumor Biology</i> , 2015, 36, 6451-6461.	0.8	31
70	Development of a multiplex system for identifying individuals of Andean Condor (<i>Vultur gryphus</i>). <i>Forensic Science International: Genetics Supplement Series</i> , 2015, 5, e228-e230.	0.1	2
71	Positive Association between the Polymorphic Variant CCND1 A870G and Colorectal Cancer in Ecuadorian Mestizo Population. <i>Journal of Cancer Research Updates</i> , 2015, 4, 163-170.	0.3	3
72	Association of genetic variants of membrane receptors related to recognition and induction of immune response with <i>Helicobacter pylori</i> infection in Ecuadorian individuals. <i>International Journal of Immunogenetics</i> , 2014, 41, 281-288.	0.8	8

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73	Frequency of GJB2 and del(GJB6-D13S1830) mutations among an Ecuadorian mestizo population. <i>International Journal of Pediatric Otorhinolaryngology</i> , 2014, 78, 1648-1654.	0.4	11
74	Perception of the Usefulness of Drug/Gene Pairs and Barriers for Pharmacogenomics in Latin America. <i>Current Drug Metabolism</i> , 2014, 15, 202-208.	0.7	31
75	Estado de la mutación del gen IT-15 (HTT) en familias ecuatorianas con enfermedad de Huntington. <i>Archivos - Instituto Nacional De Neurología Y Neurocirugía</i> , 2014, 19, 73-78.	0.1	1
76	Genetic Polymorphisms in MTHFR (C677T, A1298C), MTR (A2756G) and MTRR (A66G) Genes Associated With Pathological Characteristics of Prostate Cancer in the Ecuadorian Population. <i>American Journal of the Medical Sciences</i> , 2013, 346, 447-454.	0.4	38
77	Baseline determination in social, health, and genetic areas in communities affected by glyphosate aerial spraying on the northeastern Ecuadorian border. <i>Reviews on Environmental Health</i> , 2011, 26, 45-51.	1.1	19
78	Genetic Polymorphisms in Apolipoprotein E and Glutathione Peroxidase 1 Genes in the Ecuadorian Population Affected With Alzheimer's Disease. <i>American Journal of the Medical Sciences</i> , 2010, 340, 373-377.	0.4	29
79	Incidence of the L858R and G719S mutations of the epidermal growth factor receptor oncogene in an Ecuadorian population with lung cancer. <i>Cancer Genetics and Cytogenetics</i> , 2010, 196, 201-203.	1.0	12
80	Relationship of an hRAD54 gene polymorphism (2290 C/T) in an Ecuadorian population with chronic myelogenous leukemia. <i>Genetics and Molecular Biology</i> , 2010, 33, 646-649.	0.6	7
81	Frequency of Polymorphisms pro198leu in $GPX-1$ Gene and ile58thr in $MnSOD$ Gene in the Altitude Ecuadorian Population With Bladder Cancer. <i>Oncology Research</i> , 2009, 18, 395-400.	0.6	39
82	Monitoring of DNA Damage in Individuals Exposed to Petroleum Hydrocarbons in Ecuador. <i>Annals of the New York Academy of Sciences</i> , 2008, 1140, 121-128.	1.8	26
83	Osteosarcoma gene prioritization through combined bioinformatics analysis. , 0, , .		0