

Bruno Cavadas

List of Publications by Year in descending order

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Version: 2024-02-01

30
papers

849
citations

567247

15
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501174

28
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33
all docs

33
docs citations

33
times ranked

1649
citing authors

#	ARTICLE	IF	CITATIONS
1	TERT, BRAF, and NRAS in Primary Thyroid Cancer and Metastatic Disease. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2017, 102, 1898-1907.	3.6	113
2	Resolving the ancestry of Austronesian-speaking populations. <i>Human Genetics</i> , 2016, 135, 309-326.	3.8	71
3	Characterization of microbiota in male infertility cases uncovers differences in seminal hyperviscosity and oligoasthenoteratozoospermia possibly correlated with increased prevalence of infectious bacteria. <i>American Journal of Reproductive Immunology</i> , 2018, 79, e12838.	1.2	70
4	Profiling of lung microbiota discloses differences in adenocarcinoma and squamous cell carcinoma. <i>Scientific Reports</i> , 2019, 9, 12838.	3.3	64
5	NIS expression in thyroid tumors, relation with prognosis clinicopathological and molecular features. <i>Endocrine Connections</i> , 2018, 7, 78-90.	1.9	56
6	Chitinase 3-like-1 and fibronectin in the cargo of extracellular vesicles shed by human macrophages influence pancreatic cancer cellular response to gemcitabine. <i>Cancer Letters</i> , 2021, 501, 210-223.	7.2	51
7	OSBPL10, RXRA and lipid metabolism confer African-ancestry protection against dengue haemorrhagic fever in admixed Cubans. <i>PLoS Pathogens</i> , 2017, 13, e1006220.	4.7	51
8	Genome-Wide Characterization of Arabian Peninsula Populations: Shedding Light on the History of a Fundamental Bridge between Continents. <i>Molecular Biology and Evolution</i> , 2019, 36, 575-586.	8.9	45
9	Genetic Stratigraphy of Key Demographic Events in Arabia. <i>PLoS ONE</i> , 2015, 10, e0118625.	2.5	40
10	Quantifying the legacy of the Chinese Neolithic on the maternal genetic heritage of Taiwan and Island Southeast Asia. <i>Human Genetics</i> , 2016, 135, 363-376.	3.8	28
11	CRABP1, C1QL1 and LCN2 are biomarkers of differentiated thyroid carcinoma, and predict extrathyroidal extension. <i>BMC Cancer</i> , 2018, 18, 68.	2.6	26
12	Afadin Downregulation by Helicobacter pylori Induces Epithelial to Mesenchymal Transition in Gastric Cells. <i>Frontiers in Microbiology</i> , 2018, 9, 2712.	3.5	22
13	Helicobacter Pylori Targets the EPHA2 Receptor Tyrosine Kinase in Gastric Cells Modulating Key Cellular Functions. <i>Cells</i> , 2020, 9, 513.	4.1	19
14	Expression and Clinical Relevance of SOX9 in Gastric Cancer. <i>Disease Markers</i> , 2019, 2019, 1-11.	1.3	18
15	Joint ancestry and association test indicate two distinct pathogenic pathways involved in classical dengue fever and dengue shock syndrome. <i>PLoS Neglected Tropical Diseases</i> , 2018, 12, e0006202.	3.0	17
16	Population genetics-informed meta-analysis in seven genes associated with risk to dengue fever disease. <i>Infection, Genetics and Evolution</i> , 2018, 62, 60-72.	2.3	16
17	A panel of intestinal differentiation markers (CDX2, GPA33, and LI-cadherin) identifies gastric cancer patients with favourable prognosis. <i>Gastric Cancer</i> , 2020, 23, 811-823.	5.3	16
18	Association of Leukotriene A4 Hydrolase with Tuberculosis Susceptibility Using Genomic Data in Portugal. <i>Microorganisms</i> , 2019, 7, 650.	3.6	14

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19	Gastric Microbiome Diversities in Gastric Cancer Patients from Europe and Asia Mimic the Human Population Structure and Are Partly Driven by Microbiome Quantitative Trait Loci. <i>Microorganisms</i> , 2020, 8, 1196.	3.6	14
20	A different view on fine-scale population structure in Western African populations. <i>Human Genetics</i> , 2020, 139, 45-59.	3.8	13
21	Fine Time Scaling of Purifying Selection on Human Nonsynonymous mtDNA Mutations Based on the Worldwide Population Tree and Mother-Child Pairs. <i>Human Mutation</i> , 2015, 36, 1100-1111.	2.5	11
22	Human Genomic Diversity Where the Mediterranean Joins the Atlantic. <i>Molecular Biology and Evolution</i> , 2020, 37, 1041-1055.	8.9	11
23	Crime Prediction Using Regression and Resources Optimization. <i>Lecture Notes in Computer Science</i> , 2015, , 513-524.	1.3	10
24	Genomic and transcriptomic characterization of the mitochondrial-rich oncocytic phenotype on a thyroid carcinoma background. <i>Mitochondrion</i> , 2019, 46, 123-133.	3.4	10
25	Shedding Light on the African Enigma: In Vitro Testing of Homo sapiens-Helicobacter pylori Coevolution. <i>Microorganisms</i> , 2021, 9, 240.	3.6	10
26	The mitogenome portrait of Umbria in Central Italy as depicted by contemporary inhabitants and pre-Roman remains. <i>Scientific Reports</i> , 2020, 10, 10700.	3.3	9
27	Hypoxia and Macrophages Act in Concert Towards a Beneficial Outcome in Colon Cancer. <i>Cancers</i> , 2020, 12, 818.	3.7	9
28	Projecting Ancient Ancestry in Modern-Day Arabians and Iranians: A Key Role of the Past Exposed Arabo-Persian Gulf on Human Migrations. <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	6
29	Sub-Saharan African information potential to unveil adaptations to infectious disease. <i>Human Molecular Genetics</i> , 2021, 30, R138-R145.	2.9	2
30	Multi-Tissue Transcriptomic-Informed In Silico Investigation of Drugs for the Treatment of Dengue Fever Disease. <i>Viruses</i> , 2021, 13, 1540.	3.3	1