## Jose Luis Royo

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8150381/publications.pdf

Version: 2024-02-01

54 papers

2,060 citations

19 h-index 276875 41 g-index

54 all docs 54 docs citations

54 times ranked

3079 citing authors

#	Article	IF	CITATIONS
1	New insights into the genetic etiology of Alzheimer's disease and related dementias. Nature Genetics, 2022, 54, 412-436.	21.4	700
2	Common variants in Alzheimer's disease and risk stratification by polygenic risk scores. Nature Communications, 2021, 12, 3417.	12.8	140
3	Genomeâ€wide association analysis of dementia and its clinical endophenotypes reveal novel loci associated with Alzheimer's disease and three causality networks: The GR@ACE project. Alzheimer's and Dementia, 2019, 15, 1333-1347.	0.8	111
4	Pyrosequencing protocol using a universal biotinylated primer for mutation detection and SNP genotyping. Nature Protocols, 2007, 2, 1734-1739.	12.0	97
5	Transphyletic conservation of developmental regulatory state in animal evolution. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 14186-14191.	7.1	94
6	In vivo gene regulation in Salmonella spp. by a salicylate-dependent control circuit. Nature Methods, 2007, 4, 937-942.	19.0	84
7	CAPN10 Alleles Are Associated with Polycystic Ovary Syndrome. Journal of Clinical Endocrinology and Metabolism, 2002, 87, 3971-3976.	3.6	76
8	A method for detecting epistasis in genome-wide studies using case-control multi-locus association analysis. BMC Genomics, 2008, 9, 360.	2.8	76
9	Vitamin D Receptor polymorphisms and risk of enveloped virus infection: A meta-analysis. Gene, 2018, 678, 384-394.	2.2	69
10	Identification of the first recurrent PAR1 deletion in LÃ $@$ ri-Weill dyschondrosteosis and idiopathic short stature reveals the presence of a novel <i>SHOX</i> enhancer. Journal of Medical Genetics, 2012, 49, 442-450.	3.2	63
11	Genetic Structure of the Spanish Population. BMC Genomics, 2010, 11, 326.	2.8	49
12	Association of genetic markers within the KIT and KITLG genes with human male infertility. Human Reproduction, 2006, 21, 3185-3192.	0.9	40
13	Estrogen receptor alpha gene variants are associated with Alzheimer's disease. Neurobiology of Aging, 2012, 33, 198.e15-198.e24.	3.1	36
14	Comprehensive Toxicity Assessment of PEGylated Magnetic Nanoparticles for in vivo applications. Colloids and Surfaces B: Biointerfaces, 2019, 177, 253-259.	5.0	33
15	A new generation of vectors with increased induction ratios by overimposing a second regulatory level by attenuation. Nucleic Acids Research, 2005, 33, e169-e169.	14.5	30
16	Stable long-term indigo production by overexpression of dioxygenase genes using a chromosomal integrated cascade expression circuit. Journal of Biotechnology, 2005, 116, 113-124.	3.8	30
17	Synthesis and Characterization of Elongated-Shaped Silver Nanoparticles as a Biocompatible Anisotropic SERS Probe for Intracellular Imaging: Theoretical Modeling and Experimental Verification. Nanomaterials, 2019, 9, 256.	4.1	27
18	Improvement of Recombinant Protein Yield by a Combination of Transcriptional Amplification and Stabilization of Gene Expression. Applied and Environmental Microbiology, 2002, 68, 5034-5041.	3.1	24

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19	Lack of Association Between NOS3 Glu298Asp and Breast Cancer Risk: a Case–ontrol Study. Breast Cancer Research and Treatment, 2006, 100, 331-333.	2.5	21
20	Identification and Analysis of Conserved cis-Regulatory Regions of the MEIS1 Gene. PLoS ONE, 2012, 7, e33617.	2.5	20
21	Genetic analysis of caveolin-1 and eNOS genes in colorectal cancer. Oncology Reports, 2006, 16, 353-9.	2.6	18
22	<i><scp>SIRPB1</scp></i> copyâ€number polymorphism as candidate quantitative trait locus for impulsiveâ€disinhibited personality. Genes, Brain and Behavior, 2014, 13, 653-662.	2.2	17
23	Spectrofluorimetric Analysis of CCR5-Δ32 Allele Using Real-Time Polymerase Chain Reaction: Prevalence in Southern Spanish HIV+ Patients and Noninfected Population. AIDS Research and Human Retroviruses, 2001, 17, 191-193.	1.1	16
24	Pyrosequencing for SNP Genotyping. Methods in Molecular Biology, 2009, 578, 123-133.	0.9	16
25	Fluorescence Resonance Energy Transfer Analysis of CCR2-V64I and SDF1-3′A Polymorphisms: Prevalence in Southern Spain HIV Type 1+Cohort and Noninfected Population. AIDS Research and Human Retroviruses, 2001, 17, 663-666.	1.1	14
26	Identification of a protective haplogenotype within CAPN10 gene influencing colorectal cancer susceptibility. Journal of Gastroenterology and Hepatology (Australia), 2007, 22, 2298-2302.	2.8	14
27	Dissecting the Transcriptional Regulatory Properties of Human Chromosome 16 Highly Conserved Non-Coding Regions. PLoS ONE, 2011, 6, e24824.	2.5	13
28	Zebrafish as a model organism to study host–pathogen interactions. Methods, 2013, 62, 241-245.	3.8	13
29	White matter lesions and temporal atrophy are associated with cognitive and neuropsychiatric symptoms in patients with hypertension and Alzheimer's disease. International Journal of Geriatric Psychiatry, 2020, 35, 1292-1300.	2.7	11
30	Absence of Substantial Copy Number Differences in a Pair of Monozygotic Twins Discordant for Features of Autism Spectrum Disorder. Case Reports in Genetics, 2014, 2014, 1-9.	0.2	10
31	Genetic analysis of CAV1 gene in hypertension and metabolic syndrome. Thrombosis and Haemostasis, 2006, 95, 696-701.	3.4	9
32	Exploring allelic imbalance within paraffin-embedded tumor biopsies using pyrosequencing technology. Clinical Chemistry and Laboratory Medicine, 2006, 44, 1076-81.	2.3	7
33	Identification of a 2244 base pair interstitial deletion within the human ESR1 gene in the Spanish population. Journal of Medical Genetics, 2008, 45, 420-424.	3.2	7
34	Dysmorphic contribution of neurotransmitter and neuroendocrine system polymorphisms to subtherapeutic mood states. Brain and Behavior, 2019, 9, e01140.	2.2	7
35	Absence of allelic imbalance involving EMSY, CAPN5, and PAK1 genes in papillary thyroid carcinoma. Journal of Endocrinological Investigation, 2008, 31, 618-623.	3.3	6
36	A common copy-number variant within SIRPB1 correlates with human Out-of-Africa migration after genetic drift correction. PLoS ONE, 2018, 13, e0193614.	2.5	6

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37	Long runs of homozygosity are associated with Alzheimer's disease. Translational Psychiatry, 2021, 11, 142.	4.8	6
38	Hardy Weinberg Equilibrium Disturbances in Case-Control Studies Lead to Non-Conclusive Results. Cell Journal, 2021, 22, 572-574.	0.2	6
39	Pyrosequencing protocol requiring a unique biotinylated primer. Clinical Chemistry and Laboratory Medicine, 2006, 44, 435-41.	2.3	5
40	Engineered Salmonella allows real-time heterologous gene expression monitoring within infected zebrafish embryos. Journal of Biotechnology, 2012, 157, 413-416.	3.8	5
41	A Knockout IFNL4 Variant Is Associated With Protection From Sexually Transmitted HIV-1 Infection. Journal of Infectious Diseases, 2019, 219, 772-776.	4.0	5
42	Monoamino oxidase alleles correlate with the presence of essential hypertension among hypogonadic patients. Molecular Genetics & Enomic Medicine, 2020, 8, e1040.	1.2	5
43	Methylation alterations are not a major cause of PTTG1 missregulation. BMC Cancer, 2008, 8, 110.	2.6	4
44	Discordance in TLR2 (â~196 to â~174) polymorphism effect on HIV infection risk. Journal of Gene Medicine, 2018, 20, e3051.	2.8	4
45	Toll-Like Receptor 2 Promoter -196 to -174 Deletion Affects CD4 Levels Along Human Immunodeficiency Virus Infection Progression. Journal of Infectious Diseases, 2020, 222, 2007-2011.	4.0	4
46	Differential metabolic profiles associated to movement behaviour of stream-resident brown trout (Salmo trutta). PLoS ONE, 2017, 12, e0181697.	2.5	4
47	Fetal alpha 5-reductase Val89Leu mutation is associated with late miscarriage. Reproductive BioMedicine Online, 2017, 34, 653-658.	2.4	3
48	MAOB rs3027452 Modifies Mood Improvement After Tryptophan Supplementation. International Journal of General Medicine, 2021, Volume 14, 1751-1756.	1.8	3
49	Genotyping of common SIRPB1 copy number variant using Paralogue Ratio Test coupled to MALDI-MS quantification. Molecular and Cellular Probes, 2015, 29, 517-521.	2.1	2
50	Genetic Association Studies in Host–Pathogen Interaction Analysis. Methods in Molecular Biology, 2018, 1734, 1-11.	0.9	0
51	Straightforward protocol for allele-specific chromatin conformation capture. Gene, 2021, 767, 145185.	2.2	0
52	The Genetic Research in Alzheimer Disease (GERALD) Initiative Finds rs9320913 as a Neural eQTL of lincRNA AL589740.1. International Journal of Alzheimer's Disease, 2021, 2021, 1-7.	2.0	0
53	A polygenic risk score for mosaic loss of chromosome Y susceptibility is associated with higher risk of MCI to AD conversion Alzheimer's and Dementia, 2021, 17 Suppl 3, e053745.	0.8	0
54	White matter lesions and hippocampal atrophy are associated with cognitive and behavioral and psychological symptoms in patients with Alzheimer's disease and hypertension Alzheimer's and Dementia, 2021, 17 Suppl 3, e054134.	0.8	O