

Douglas Londono

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

807
citations

933447

10
h-index

996975

15
g-index

15
all docs

15
docs citations

15
times ranked

1090
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic variants in GPR126 are associated with adolescent idiopathic scoliosis. <i>Nature Genetics</i> , 2013, 45, 676-679.	21.4	240
2	Genome-wide association studies of adolescent idiopathic scoliosis suggest candidate susceptibility genes. <i>Human Molecular Genetics</i> , 2011, 20, 1456-1466.	2.9	172
3	A PAX1 enhancer locus is associated with susceptibility to idiopathic scoliosis in females. <i>Nature Communications</i> , 2015, 6, 6452.	12.8	122
4	A meta-analysis identifies adolescent idiopathic scoliosis association with <i>LBX1</i> locus in multiple ethnic groups. <i>Journal of Medical Genetics</i> , 2014, 51, 401-406.	3.2	79
5	Chromatin Profiling Reveals Regulatory Network Shifts and a Protective Role for Hepatocyte Nuclear Factor 4 μ during Colitis. <i>Molecular and Cellular Biology</i> , 2014, 34, 3291-3304.	2.3	41
6	<i>EDNRA</i> variants associate with smooth muscle mRNA levels, cell proliferation rates, and cystic fibrosis pulmonary disease severity. <i>Physiological Genomics</i> , 2010, 41, 71-77.	2.3	34
7	TDT-HET: A new transmission disequilibrium test that incorporates locus heterogeneity into the analysis of family-based association data. <i>BMC Bioinformatics</i> , 2012, 13, 13.	2.6	27
8	Investigation of previously implicated genetic variants in chronic tic disorders: a transmission disequilibrium test approach. <i>European Archives of Psychiatry and Clinical Neuroscience</i> , 2018, 268, 301-316.	3.2	23
9	Beta 2 adrenergic receptor polymorphisms in cystic fibrosis. <i>Pediatric Pulmonology</i> , 2005, 39, 544-550.	2.0	16
10	A novel method for analyzing genetic association with longitudinal phenotypes. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2013, 12, 241-61.	0.6	16
11	Single-Variant and Multi-Variant Trend Tests for Genetic Association with Next-Generation Sequencing That Are Robust to Sequencing Error. <i>Human Heredity</i> , 2012, 74, 172-183.	0.8	10
12	A Cost-Effective Statistical Method to Correct for Differential Genotype Misclassification When Performing Case-Control Genetic Association. <i>Human Heredity</i> , 2010, 70, 102-108.	0.8	9
13	Bone marrow-derived epithelial cells and hair follicle stem cells contribute to development of chronic cutaneous neoplasms. <i>Nature Communications</i> , 2018, 9, 5293.	12.8	9
14	Mapping genes with longitudinal phenotypes via Bayesian posterior probabilities. <i>BMC Proceedings</i> , 2014, 8, S81.	1.6	6
15	An Analytic Solution to the Computation of Power and Sample Size for Genetic Association Studies under a Pleiotropic Mode of Inheritance. <i>Human Heredity</i> , 2016, 81, 194-209.	0.8	3