

Elaine M Kenny

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

1,239
citations

567281

15
h-index

752698

20
g-index

20
all docs

20
docs citations

20
times ranked

3424
citing authors

#	ARTICLE	IF	CITATIONS
1	Targeted Sequencing of 10,198 Samples Confirms Abnormalities in Neuronal Activity and Implicates Voltage-Gated Sodium Channels in Schizophrenia Pathogenesis. <i>Biological Psychiatry</i> , 2019, 85, 554-562.	1.3	40
2	ltn2a silencing rescues lamin A mediated inhibition of 3T3-L1 adipocyte differentiation. <i>Adipocyte</i> , 2017, 6, 259-276.	2.8	5
3	Translation of 5' UTR leaders is pervasive in genes resistant to eIF2 repression. <i>ELife</i> , 2015, 4, e03971.	6.0	294
4	An inherited duplication at the gene p21 Protein-Activated Kinase 7 (PAK7) is a risk factor for psychosis. <i>Human Molecular Genetics</i> , 2014, 23, 3316-3326.	2.9	37
5	Identification of mechanosensitive genes during skeletal development: alteration of genes associated with cytoskeletal rearrangement and cell signalling pathways. <i>BMC Genomics</i> , 2014, 15, 48.	2.8	80
6	Transcriptome analysis of the mouse E14.5 (TS23) developing humerus and differential expression in muscle-less mutant embryos lacking mechanical stimulation. <i>Genomics Data</i> , 2014, 2, 32-36.	1.3	5
7	Delineating the genetic heterogeneity of ALS using targeted high-throughput sequencing. <i>Journal of Medical Genetics</i> , 2013, 50, 776-783.	3.2	151
8	Development of Strategies for SNP Detection in RNA-Seq Data: Application to Lymphoblastoid Cell Lines and Evaluation Using 1000 Genomes Data. <i>PLoS ONE</i> , 2013, 8, e58815.	2.5	108
9	Global endometrial transcriptomic profiling: transient immune activation precedes tissue proliferation and repair in healthy beef cows. <i>BMC Genomics</i> , 2012, 13, 489.	2.8	26
10	Avian Resistance to <i>Campylobacter jejuni</i> Colonization Is Associated with an Intestinal Immunogene Expression Signature Identified by mRNA Sequencing. <i>PLoS ONE</i> , 2012, 7, e40409.	2.5	46
11	Assessment of Inactivating Stop Codon Mutations in Forty <i>Saccharomyces cerevisiae</i> Strains: Implications for [PSI ⁺] Prion-Mediated Phenotypes. <i>PLoS ONE</i> , 2011, 6, e28684.	2.5	13
12	Gene-ontology enrichment analysis in two independent family-based samples highlights biologically plausible processes for autism spectrum disorders. <i>European Journal of Human Genetics</i> , 2011, 19, 1082-1089.	2.8	39
13	First implication of STRA6 mutations in isolated anophthalmia, microphthalmia, and coloboma: A new dimension to the STRA6 phenotype. <i>Human Mutation</i> , 2011, 32, 1417-1426.	2.5	66
14	Allelic expression imbalance of the schizophrenia susceptibility gene CHI3L1. <i>Psychiatric Genetics</i> , 2011, 21, 281-286.	1.1	2
15	Multiplex Target Enrichment Using DNA Indexing for Ultra-High Throughput SNP Detection. <i>DNA Research</i> , 2011, 18, 31-38.	3.4	41
16	The SNP ratio test: pathway analysis of genome-wide association datasets. <i>Bioinformatics</i> , 2009, 25, 2762-2763.	4.1	125
17	Non-random error in genotype calling procedures: Implications for family-based and case-control genome-wide association studies. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008, 147B, 1379-1386.	1.7	11
18	Conduct disorder and ADHD: Evaluation of conduct problems as a categorical and quantitative trait in the international multicentre ADHD genetics study. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008, 147B, 1369-1378.	1.7	106

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19	Chitinase-3-Like 1 (CHI3L1) Gene and Schizophrenia: Genetic Association and a Potential Functional Mechanism. <i>Biological Psychiatry</i> , 2008, 64, 98-103.	1.3	28
20	Feasibility and relevance of global expression profiling of gene transcripts in serum from breast cancer patients using whole genome microarrays and quantitative RT-PCR. <i>Cancer Genomics and Proteomics</i> , 2008, 5, 94-104.	2.0	16