Jiaqian Wu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10723474/publications.pdf

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11 papers	5,581 citations	9 h-index	11 g-index
12	12	12	10329
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Identification and analysis of functional elements in 1% of the human genome by the ENCODE pilot project. Nature, 2007, 447, 799-816.	27.8	4,709
2	The Status, Quality, and Expansion of the NIH Full-Length cDNA Project: The Mammalian Gene Collection (MGC). Genome Research, 2004, 14, 2121-2127.	5.5	486
3	The completion of the Mammalian Gene Collection (MGC). Genome Research, 2009, 19, 2324-2333.	5.5	125
4	Exosomes Mediate Epithelium–Mesenchyme Crosstalk in Organ Development. ACS Nano, 2017, 11, 7736-7746.	14.6	100
5	Parenchymal and stromal tissue regeneration of tooth organ by pivotal signals reinstated in decellularized matrix. Nature Materials, 2019, 18, 627-637.	27.5	53
6	Integrated analysis of experimental data sets reveals many novel promoters in 1% of the human genome. Genome Research, 2007, 17, 720-731.	5.5	31
7	Angiogenic gene networks are dysregulated in opioid use disorder: evidence from multi-omics and imaging of postmortem human brain. Molecular Psychiatry, 2021, 26, 7803-7812.	7.9	31
8	The DART classification of unannotated transcription within the ENCODE regions: Associating transcription with known and novel loci. Genome Research, 2007, 17, 732-745.	5.5	25
9	Hemogenic Endothelial Cells Can Transition to Hematopoietic Stem Cells through a B-1 Lymphocyte-Biased State during Maturation in the Mouse Embryo. Stem Cell Reports, 2019, 13, 21-30.	4.8	14
10	Long Noncoding RNAs: Critical Regulators for Cell Lineage Commitment in the Central Nervous System. Translational Bioinformatics, 2016, , 73-97.	0.0	3
11	The Analyses of Global Gene Expression and Transcription Factor Regulation. Translational Bioinformatics, 2016, , 1-35.	0.0	2