

Veronique Lisi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/184254/publications.pdf>

Version: 2024-02-01

14
papers

698
citations

932766

10
h-index

1281420

11
g-index

15
all docs

15
docs citations

15
times ranked

1230
citing authors

#	ARTICLE	IF	CITATIONS
1	Histone H3.3G34-Mutant Interneuron Progenitors Co-opt PDGFRA for Gliomagenesis. <i>Cell</i> , 2020, 183, 1617-1633.e22.	13.5	93
2	H3.3 G34W Promotes Growth and Impedes Differentiation of Osteoblast-Like Mesenchymal Progenitors in Giant Cell Tumor of Bone. <i>Cancer Discovery</i> , 2020, 10, 1968-1987.	7.7	40
3	The dual function of LMO2 in driving erythroid cell fate. <i>Experimental Hematology</i> , 2017, 53, S98-S99.	0.2	0
4	Enhanced Neuronal Regeneration in the CAST/Ei Mouse Strain Is Linked to Expression of Differentiation Markers after Injury. <i>Cell Reports</i> , 2017, 20, 1136-1147.	2.9	26
5	A Primate lncRNA Mediates Notch Signaling during Neuronal Development by Sequestering miRNA. <i>Neuron</i> , 2016, 90, 1174-1188.	3.8	115
6	The LMO2 oncogene regulates DNA replication in hematopoietic cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 1393-1398.	3.3	24
7	Robust Axonal Regeneration Occurs in the Injured CAST/Ei Mouse CNS. <i>Neuron</i> , 2015, 86, 1215-1227.	3.8	87
8	MiRBooking simulates the stoichiometric mode of action of microRNAs. <i>Nucleic Acids Research</i> , 2015, 43, 6730-6738.	6.5	18
9	LMO2 Regulates DNA Replication in Hematopoietic Cells. <i>Blood</i> , 2015, 126, 553-553.	0.6	0
10	Computational identification of RNA functional determinants by three-dimensional quantitative structure-activity relationships. <i>Nucleic Acids Research</i> , 2014, 42, 11261-11271.	6.5	4
11	<i>RNA-Puzzles</i> : A CASP-like evaluation of RNA three-dimensional structure prediction. <i>Rna</i> , 2012, 18, 610-625.	1.6	241
12	Role of SLV in SLI substrate recognition by the <i>Neurospora</i> VS ribozyme. <i>Rna</i> , 2008, 14, 736-748.	1.6	24
13	A comparative analysis of the triloops in all high-resolution RNA structures reveals sequence-structure relationships. <i>Rna</i> , 2007, 13, 1537-1545.	1.6	25
14	Identification of SCL and LMO Oncogenes as Transcriptional Repressors of HEB/E2A in the Thymus through Functional Genomic/Genetic Approaches.. <i>Blood</i> , 2007, 110, 2649-2649.	0.6	0