

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/34418/publications.pdf Version: 2024-02-01



LIE LV

#	Article	IF	CITATIONS
1	Insights into bilaterian evolution from three spiralian genomes. Nature, 2013, 493, 526-531.	27.8	564
2	The First Myriapod Genome Sequence Reveals Conservative Arthropod Gene Content and Genome Organisation in the Centipede Strigamia maritima. PLoS Biology, 2014, 12, e1002005.	5.6	221
3	MLL4 Is Required to Maintain Broad H3K4me3 Peaks and Super-Enhancers at Tumor Suppressor Genes. Molecular Cell, 2018, 70, 825-841.e6.	9.7	123
4	ZMYND8 Reads the Dual Histone Mark H3K4me1-H3K14ac to Antagonize the Expression of Metastasis-Linked Genes. Molecular Cell, 2016, 63, 470-484.	9.7	112
5	AIBP-mediated cholesterol efflux instructs hematopoietic stem and progenitor cell fate. Science, 2019, 363, 1085-1088.	12.6	90
6	Joint assembly and genetic mapping of the Atlantic horseshoe crab genome reveals ancient whole genome duplication. GigaScience, 2014, 3, 9.	6.4	86
7	HP1Î ³ Promotes Lung Adenocarcinoma by Downregulating the Transcription-Repressive Regulators NCOR2 and ZBTB7A. Cancer Research, 2018, 78, 3834-3848.	0.9	63
8	Developing Spindlin1 small-molecule inhibitors by using protein microarrays. Nature Chemical Biology, 2017, 13, 750-756.	8.0	47
9	Reservoir of Fibroblasts Promotes Recovery From Limb Ischemia. Circulation, 2020, 142, 1647-1662.	1.6	33
10	TBX20 Regulates Angiogenesis Through the Prokineticin 2–Prokineticin Receptor 1 Pathway. Circulation, 2018, 138, 913-928.	1.6	31
11	Machine learning uncovers cell identity regulator by histone code. Nature Communications, 2020, 11, 2696.	12.8	25
12	Constraints on genes shape long-term conservation of macro-synteny in metazoan genomes. BMC Bioinformatics, 2011, 12, S11.	2.6	24
13	Lmo2 (LIM-Domain-Only 2) Modulates Sphk1 (Sphingosine Kinase) and Promotes Endothelial Cell Migration. Arteriosclerosis, Thrombosis, and Vascular Biology, 2017, 37, 1860-1868.	2.4	21
14	LIM Domain Only 2 Regulates Endothelial Proliferation, Angiogenesis, and Tissue Regeneration. Journal of the American Heart Association, 2016, 5, .	3.7	19
15	Association between the availability of environmental resources and the atomic composition of organismal proteomes: Evidence from Prochlorococcus strains living at different depths. Biochemical and Biophysical Research Communications, 2008, 375, 241-246.	2.1	18
16	Heterogeneous rates of genome rearrangement contributed to the disparity of species richness in Ascomycota. BMC Genomics, 2018, 19, 282.	2.8	17
17	Low Contents of Carbon and Nitrogen in Highly Abundant Proteins: Evidence of Selection for the Economy of Atomic Composition. Journal of Molecular Evolution, 2009, 68, 248-255.	1.8	15
18	Broad genic repression domains signify enhanced silencing of oncogenes. Nature Communications, 2020, 11, 5560.	12.8	10

Jie Lv

#	Article	IF	CITATIONS
19	Broad H3K4me3 as A Novel Epigenetic Signature for Normal Development and Disease. Genomics, Proteomics and Bioinformatics, 2016, 14, 262-264.	6.9	8
20	TADsplimer reveals splits and mergers of topologically associating domains for epigenetic regulation of transcription. Genome Biology, 2020, 21, 84.	8.8	6
21	MACMIC Reveals A Dual Role of CTCF in Epigenetic Regulation of Cell Identity Genes. Genomics, Proteomics and Bioinformatics, 2021, 19, 140-153.	6.9	4