Fei Ji

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10981166/publications.pdf

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38	2,852	21 h-index	37
papers	citations		g-index
41	41	41	5917 citing authors
all docs	docs citations	times ranked	

#	Article	IF	Citations
1	HERVH-derived IncRNAs negatively regulate chromatin targeting and remodeling mediated by CHD7. Life Science Alliance, 2022, 5, e202101127.	1.3	3
2	Bone marrow endothelial dysfunction promotes myeloid cell expansion in cardiovascular disease. , 2022, 1, 28-44.		32
3	DNA replication fork speed underlies cell fate changes and promotes reprogramming. Nature Genetics, 2022, 54, 318-327.	9.4	38
4	Regulation of chromatin accessibility by the histone chaperone CAF-1 sustains lineage fidelity. Nature Communications, 2022, 13, 2350.	5.8	8
5	DEPCOD: a tool to detect and visualize co-evolution of protein domains. Nucleic Acids Research, 2022,	6.5	O
6	Phenotypic continuum between Waardenburg syndrome and idiopathic hypogonadotropic hypogonadism in humans with SOX10 variants. Genetics in Medicine, 2021, 23, 629-636.	1,1	9
7	RNA m6A reader IMP2/IGF2BP2 promotes pancreatic \hat{l}^2 -cell proliferation and insulin secretion by enhancing PDX1 expression. Molecular Metabolism, 2021, 48, 101209.	3.0	28
8	Dissecting dual roles of MyoD during lineage conversion to mature myocytes and myogenic stem cells. Genes and Development, 2021, 35, 1209-1228.	2.7	20
9	tiRNA signaling via stress-regulated vesicle transfer in the hematopoietic niche. Cell Stem Cell, 2021, 28, 2090-2103.e9.	5.2	20
10	Collective regulation of chromatin modifications predicts replication timing during cell cycle. Cell Reports, 2021, 37, 109799.	2.9	20
11	Histone Lysine Methylation Dynamics Control <i>EGFR</i> DNA Copy-Number Amplification. Cancer Discovery, 2020, 10, 306-325.	7.7	31
12	A MicroRNA Linking Human Positive Selection and Metabolic Disorders. Cell, 2020, 183, 684-701.e14.	13.5	46
13	The lysine demethylase KDM4A controls the cell-cycle expression of replicative canonical histone genes. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2020, 1863, 194624.	0.9	7
14	S-phase Enriched Non-coding RNAs Regulate Gene Expression and Cell Cycle Progression. Cell Reports, 2020, 31, 107629.	2.9	11
15	A post-transcriptional program of chemoresistance by AU-rich elements and TTP in quiescent leukemic cells. Genome Biology, 2020, 21, 33.	3.8	22
16	The Histone Deacetylase SIRT6 Restrains Transcription Elongation via Promoter-Proximal Pausing. Molecular Cell, 2019, 75, 683-699.e7.	4.5	50
17	Targeting FGFR overcomes EMT-mediated resistance in EGFR mutant non-small cell lung cancer. Oncogene, 2019, 38, 6399-6413.	2.6	160
18	Exercise reduces inflammatory cell production and cardiovascular inflammation via instruction of hematopoietic progenitor cells. Nature Medicine, 2019, 25, 1761-1771.	15.2	157

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19	Inducible histone K-to-M mutations are dynamic tools to probe the physiological role of site-specific histone methylation in vitro and in vivo. Nature Cell Biology, 2019, 21, 1449-1461.	4.6	40
20	The RNA Helicase DDX6 Controls Cellular Plasticity by Modulating P-Body Homeostasis. Cell Stem Cell, 2019, 25, 622-638.e13.	5.2	82
21	IMP2 Increases Mouse Skeletal Muscle Mass and Voluntary Activity by Enhancing Autocrine Insulin-Like Growth Factor 2 Production and Optimizing Muscle Metabolism. Molecular and Cellular Biology, 2019, 39, .	1.1	12
22	Singleâ€Cell RNAâ€seq: Introduction to Bioinformatics Analysis. Current Protocols in Molecular Biology, 2019, 127, e92.	2.9	10
23	Sudden sensorineural hearing loss (SSHL) following a local anesthetic dental procedure. Journal of Otology, 2019, 14, 67-72.	0.4	1
24	Mitochondrial Dysfunction in C.Âelegans Activates Mitochondrial Relocalization and Nuclear Hormone Receptor-Dependent Detoxification Genes. Cell Metabolism, 2019, 29, 1182-1191.e4.	7. 2	55
25	Nudt21 Controls Cell Fate by Connecting Alternative Polyadenylation to Chromatin Signaling. Cell, 2018, 172, 106-120.e21.	13.5	123
26	RNAâ€seq: Basic Bioinformatics Analysis. Current Protocols in Molecular Biology, 2018, 124, e68.	2.9	44
27	Mutant GNAS drives pancreatic tumourigenesis by inducing PKA-mediated SIK suppression and reprogramming lipid metabolism. Nature Cell Biology, 2018, 20, 811-822.	4.6	124
28	The Association of Obesity and Cardiometabolic Traits With IncidentÂHFpEF and HFrEF. JACC: Heart Failure, 2018, 6, 701-709.	1.9	254
29	The surveillance of pre-mRNA splicing is an early step in <i>C. elegans</i> RNAi of endogenous genes. Genes and Development, 2018, 32, 670-681.	2.7	27
30	Nextâ€Generation Sequencing for Identification of EMSâ€Induced Mutations in <i>Caenorhabditis elegans</i> . Current Protocols in Molecular Biology, 2017, 117, 7.29.1-7.29.12.	2.9	19
31	Polycomb Repressive Complex 1 Generates Discrete Compacted Domains that Change during Differentiation. Molecular Cell, 2017, 65, 432-446.e5.	4.5	287
32	Maintenance of macrophage transcriptional programs and intestinal homeostasis by epigenetic reader SP140. Science Immunology, 2017, 2 , .	5.6	54
33	Unitary ototoxic gentamicin exposure may not disrupt the function of cochlear outer hair cells in mice. Acta Oto-Laryngologica, 2017, 137, 842-849.	0.3	1
34	PAR-TERRA directs homologous sex chromosome pairing. Nature Structural and Molecular Biology, 2017, 24, 620-631.	3.6	48
35	IGF2 mRNA binding protein-2 is a tumor promoter that drives cancer proliferation through its client mRNAs IGF2 and HMGA1. ELife, 2017, 6, .	2.8	77
36	Inhibiting fungal multidrug resistance by disrupting an activator–Mediator interaction. Nature, 2016, 530, 485-489.	13.7	120

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37	Tumor cells can follow distinct evolutionary paths to become resistant to epidermal growth factor receptor inhibition. Nature Medicine, 2016, 22, 262-269.	15.2	768
38	PhyloGene server for identification and visualization of co-evolving proteins using normalized phylogenetic profiles. Nucleic Acids Research, 2015, 43, W154-W159.	6.5	43