

Fei Ji

List of Publications by Year in descending order

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

Version: 2024-02-01

38
papers

2,852
citations

331670

21
h-index

330143

37
g-index

41
all docs

41
docs citations

41
times ranked

5917
citing authors

#	ARTICLE	IF	CITATIONS
1	HERVH-derived lncRNAs negatively regulate chromatin targeting and remodeling mediated by CHD7. <i>Life Science Alliance</i> , 2022, 5, e202101127.	2.8	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. <i>Nature Genetics</i> , 2022, 54, 318-327.	21.4	38
4	Regulation of chromatin accessibility by the histone chaperone CAF-1 sustains lineage fidelity. <i>Nature Communications</i> , 2022, 13, 2350.	12.8	8
5	DEPCOD: a tool to detect and visualize co-evolution of protein domains. <i>Nucleic Acids Research</i> , 2022, , .	14.5	0
6	Phenotypic continuum between Waardenburg syndrome and idiopathic hypogonadotropic hypogonadism in humans with SOX10 variants. <i>Genetics in Medicine</i> , 2021, 23, 629-636.	2.4	9
7	RNA m6A reader IMP2/IGF2BP2 promotes pancreatic β -cell proliferation and insulin secretion by enhancing PDX1 expression. <i>Molecular Metabolism</i> , 2021, 48, 101209.	6.5	28
8	Dissecting dual roles of MyoD during lineage conversion to mature myocytes and myogenic stem cells. <i>Genes and Development</i> , 2021, 35, 1209-1228.	5.9	20
9	tiRNA signaling via stress-regulated vesicle transfer in the hematopoietic niche. <i>Cell Stem Cell</i> , 2021, 28, 2090-2103.e9.	11.1	20
10	Collective regulation of chromatin modifications predicts replication timing during cell cycle. <i>Cell Reports</i> , 2021, 37, 109799.	6.4	20
11	Histone Lysine Methylation Dynamics Control EGFR DNA Copy-Number Amplification. <i>Cancer Discovery</i> , 2020, 10, 306-325.	9.4	31
12	A MicroRNA Linking Human Positive Selection and Metabolic Disorders. <i>Cell</i> , 2020, 183, 684-701.e14.	28.9	46
13	The lysine demethylase KDM4A controls the cell-cycle expression of replicative canonical histone genes. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2020, 1863, 194624.	1.9	7
14	S-phase Enriched Non-coding RNAs Regulate Gene Expression and Cell Cycle Progression. <i>Cell Reports</i> , 2020, 31, 107629.	6.4	11
15	A post-transcriptional program of chemoresistance by AU-rich elements and TTP in quiescent leukemic cells. <i>Genome Biology</i> , 2020, 21, 33.	8.8	22
16	The Histone Deacetylase SIRT6 Restrains Transcription Elongation via Promoter-Proximal Pausing. <i>Molecular Cell</i> , 2019, 75, 683-699.e7.	9.7	50
17	Targeting FGFR overcomes EMT-mediated resistance in EGFR mutant non-small cell lung cancer. <i>Oncogene</i> , 2019, 38, 6399-6413.	5.9	160
18	Exercise reduces inflammatory cell production and cardiovascular inflammation via instruction of hematopoietic progenitor cells. <i>Nature Medicine</i> , 2019, 25, 1761-1771.	30.7	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. <i>Nature Cell Biology</i> , 2019, 21, 1449-1461.	10.3	40
20	The RNA Helicase DDX6 Controls Cellular Plasticity by Modulating P-Body Homeostasis. <i>Cell Stem Cell</i> , 2019, 25, 622-638.e13.	11.1	82
21	IMP2 Increases Mouse Skeletal Muscle Mass and Voluntary Activity by Enhancing Autocrine Insulin-Like Growth Factor 2 Production and Optimizing Muscle Metabolism. <i>Molecular and Cellular Biology</i> , 2019, 39, .	2.3	12
22	Single-Cell RNA-seq: Introduction to Bioinformatics Analysis. <i>Current Protocols in Molecular Biology</i> , 2019, 127, e92.	2.9	10
23	Sudden sensorineural hearing loss (SSHL) following a local anesthetic dental procedure. <i>Journal of Otology</i> , 2019, 14, 67-72.	1.0	1
24	Mitochondrial Dysfunction in <i>C. elegans</i> Activates Mitochondrial Relocalization and Nuclear Hormone Receptor-Dependent Detoxification Genes. <i>Cell Metabolism</i> , 2019, 29, 1182-1191.e4.	16.2	55
25	Nudt21 Controls Cell Fate by Connecting Alternative Polyadenylation to Chromatin Signaling. <i>Cell</i> , 2018, 172, 106-120.e21.	28.9	123
26	RNA-seq: Basic Bioinformatics Analysis. <i>Current Protocols in Molecular Biology</i> , 2018, 124, e68.	2.9	44
27	Mutant GNAS drives pancreatic tumorigenesis by inducing PKA-mediated SIK suppression and reprogramming lipid metabolism. <i>Nature Cell Biology</i> , 2018, 20, 811-822.	10.3	124
28	The Association of Obesity and Cardiometabolic Traits With Incident HFpEF and HFrEF. <i>JACC: Heart Failure</i> , 2018, 6, 701-709.	4.1	254
29	The surveillance of pre-mRNA splicing is an early step in <i>C. elegans</i> RNAi of endogenous genes. <i>Genes and Development</i> , 2018, 32, 670-681.	5.9	27
30	Next-Generation Sequencing for Identification of EMS-Induced Mutations in <i>Caenorhabditis elegans</i> . <i>Current Protocols in Molecular Biology</i> , 2017, 117, 7.29.1-7.29.12.	2.9	19
31	Polycomb Repressive Complex 1 Generates Discrete Compacted Domains that Change during Differentiation. <i>Molecular Cell</i> , 2017, 65, 432-446.e5.	9.7	287
32	Maintenance of macrophage transcriptional programs and intestinal homeostasis by epigenetic reader SP140. <i>Science Immunology</i> , 2017, 2, .	11.9	54
33	Unitary ototoxic gentamicin exposure may not disrupt the function of cochlear outer hair cells in mice. <i>Acta Oto-Laryngologica</i> , 2017, 137, 842-849.	0.9	1
34	PAR-TERRA directs homologous sex chromosome pairing. <i>Nature Structural and Molecular Biology</i> , 2017, 24, 620-631.	8.2	48
35	IGF2 mRNA binding protein-2 is a tumor promoter that drives cancer proliferation through its client mRNAs IGF2 and HMGA1. <i>ELife</i> , 2017, 6, .	6.0	77
36	Inhibiting fungal multidrug resistance by disrupting an activator-Mediator interaction. <i>Nature</i> , 2016, 530, 485-489.	27.8	120

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37	Tumor cells can follow distinct evolutionary paths to become resistant to epidermal growth factor receptor inhibition. <i>Nature Medicine</i> , 2016, 22, 262-269.	30.7	768
38	PhyloGene server for identification and visualization of co-evolving proteins using normalized phylogenetic profiles. <i>Nucleic Acids Research</i> , 2015, 43, W154-W159.	14.5	43