

Shuangli Mi

List of Publications by Year in descending order

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39
papers

4,011
citations

567281
15
h-index

454955
30
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41
all docs

41
docs citations

41
times ranked

7883
citing authors

#	ARTICLE	IF	CITATIONS
1	Disequilibrium in chicken gut microflora with avian colibacillosis is related to microenvironment damaged by antibiotics. <i>Science of the Total Environment</i> , 2021, 762, 143058.	8.0	6
2	miR-218-2 regulates cognitive functions in the hippocampus through complement component 3-dependent modulation of synaptic vesicle release. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	17
3	Identification of Potential Key lncRNAs in the Context of Mouse Myeloid Differentiation by Systematic Transcriptomics Analysis. <i>Genes</i> , 2021, 12, 630.	2.4	0
4	Heterogeneous Nuclear Ribonucleoprotein A1 Loads Batched Tumor-Promoting MicroRNAs Into Small Extracellular Vesicles With the Assist of Caveolin-1 in A549 Cells. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 687912.	3.7	9
5	The long noncoding RNA HOTAIRM1 controlled by AML1 enhances glucocorticoid resistance by activating RHOA/ROCK1 pathway through suppressing ARHGAP18. <i>Cell Death and Disease</i> , 2021, 12, 702.	6.3	8
6	Secretome of Activated Fibroblasts Induced by Exosomes for the Discovery of Biomarkers in Non-Small Cell Lung Cancer. <i>Small</i> , 2021, 17, e2004750.	10.0	18
7	Dietary Intervention With α -Amylase Inhibitor in White Kidney Beans Added Yogurt Modulated Gut Microbiota to Adjust Blood Glucose in Mice. <i>Frontiers in Nutrition</i> , 2021, 8, 664976.	3.7	12
8	A promising iPS-based single-cell cloning strategy revealing signatures of somatic mutations in heterogeneous normal cells. <i>Computational and Structural Biotechnology Journal</i> , 2020, 18, 2326-2335.	4.1	0
9	RBUD: A New Functional Potential Analysis Approach for Whole Microbial Genome Shotgun Sequencing. <i>Microorganisms</i> , 2020, 8, 1563.	3.6	1
10	Tumor-derived circulating exosomal miR-342-5p and miR-574-5p as promising diagnostic biomarkers for early-stage Lung Adenocarcinoma. <i>International Journal of Medical Sciences</i> , 2020, 17, 1428-1438.	2.5	33
11	Synaptotagmin-7 is a key factor for bipolar-like behavioral abnormalities in mice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 4392-4399.	7.1	15
12	CRISPR interference-based specific and efficient gene inactivation in the brain. <i>Nature Neuroscience</i> , 2018, 21, 447-454.	14.8	133
13	Base-excision repair and beyond – A short summary attributed to scientific achievements of Tomas Lindahl, Nobel Prize Laureate in Chemistry 2015. <i>Science China Life Sciences</i> , 2016, 59, 89-92.	4.9	3
14	Early Different Downstream Target of Glucocorticoid Receptor Contributing to Glucocorticoids Sensitivity in Kasumi-1 Cells. <i>Blood</i> , 2016, 128, 5132-5132.	1.4	0
15	Exosome and Exosomal MicroRNA: Trafficking, Sorting, and Function. <i>Genomics, Proteomics and Bioinformatics</i> , 2015, 13, 17-24.	6.9	1,466
16	Differential responses to lithium in hyperexcitable neurons from patients with bipolar disorder. <i>Nature</i> , 2015, 527, 95-99.	27.8	461
17	Identification of functional cooperative mutations of SETD2 in human acute leukemia. <i>Nature Genetics</i> , 2014, 46, 287-293.	21.4	213
18	Circulating MicroRNAs as Biomarkers for Inflammatory Diseases. <i>MicroRNA (Shariqah, United Arab)</i> Tj ETQqO 0 0 rgBT /Overlock 10 Tf 50	1.2	77

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19	An eQTL-based method identifies CTTN and ZMAT3 as pemetrexed susceptibility markers. Human Molecular Genetics, 2012, 21, 1470-1480.	2.9	16
20	Whole-Genome Sequencing of a Monozygotic Twin Pair Reveals Functional Cooperative Mutations of SETD2 in Acute Leukemia. Blood, 2012, 120, 781-781.	1.4	0
21	MLL fusion proteins preferentially regulate a subset of wild-type MLL target genes in the leukemic genome. Blood, 2011, 117, 6895-6905.	1.4	103
22	Platinum Sensitivity-Related Germline Polymorphism Discovered via a Cell-Based Approach and Analysis of Its Association with Outcome in Ovarian Cancer Patients. Clinical Cancer Research, 2011, 17, 5490-5500.	7.0	57
23	Rapid growth of a hepatocellular carcinoma and the driving mutations revealed by cell-population genetic analysis of whole-genome data. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 12042-12047.	7.1	117
24	Aberrant overexpression and function of the miR-17-92 cluster in MLL-rearranged acute leukemia. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 3710-3715.	7.1	141
25	Genome Wide Location Analysis Reveals Deregulated MicroRNA Genes In MLL Rearranged Leukemic Genome. Blood, 2010, 116, 2507-2507.	1.4	0
26	Comprehensive analysis of the impact of SNPs and CNVs on human microRNAs and their regulatory genes. RNA Biology, 2009, 6, 412-425.	3.1	58
27	Consistent Deregulation of Gene Expression between Human and Murine MLL Rearrangement Leukemias. Cancer Research, 2009, 69, 1109-1116.	0.9	81
28	Expression and alternative splicing of folate pathway genes in HapMap lymphoblastoid cell lines. Pharmacogenomics, 2009, 10, 549-563.	1.3	13
29	Population-specific genetic variants important in susceptibility to cytarabine arabinoside cytotoxicity. Blood, 2009, 113, 2145-2153.	1.4	81
30	MLL Fusion Proteins Directly Regulate a Small Set of Wild Type MLL Target Genes.. Blood, 2009, 114, 1279-1279.	1.4	0
31	Distinct microRNA expression profiles in acute myeloid leukemia with common translocations. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 15535-15540.	7.1	418
32	Identification of Genes Deregulated in Both Human and Murine MLLRearrangement Leukemias.. Blood, 2008, 112, 3359-3359.	1.4	0
33	MicroRNA expression signatures accurately discriminate acute lymphoblastic leukemia from acute myeloid leukemia. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 19971-19976.	7.1	435
34	Recent Patents on the Identification and Clinical Application of microRNAs and Target Genes. Recent Patents on DNA & Gene Sequences, 2007, 1, 116-24.	0.7	4
35	MicroRNA Expression Signatures Accurately Discriminate Acute Lymphoblastic Leukemia from Acute Myeloid Leukemia.. Blood, 2007, 110, 231-231.	1.4	0
36	MicroRNA Expression Profiles in Acute Myeloid Leukemia with Common Translocations.. Blood, 2007, 110, 3181-3181.	1.4	0

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37	Identification of Genes Abnormally Expressed in Human MLL-AF4 Leukemia.. Blood, 2006, 108, 4314-4314.	1.4	0
38	Identification of Genes Abnormally Expressed in Both Human and Murine MLL-ELL and/or MLL-ENL Leukemia.. Blood, 2006, 108, 2249-2249.	1.4	0
39	The sense and antisense expression of gibberellin 20-oxidase gene (rga5) in rice and its effects on GA1 level and agronomic traits. Science Bulletin, 2003, 48, 443-448.	1.7	1