

# 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  
g-index

41  
all docs

41  
docs citations

41  
times ranked

7883  
citing authors

#	ARTICLE	IF	CITATIONS
1	Exosome and Exosomal MicroRNA: Trafficking, Sorting, and Function. Genomics, Proteomics and Bioinformatics, 2015, 13, 17-24.	6.9	1,466
2	Differential responses to lithium in hyperexcitable neurons from patients with bipolar disorder. Nature, 2015, 527, 95-99.	27.8	461
3	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
4	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
5	Identification of functional cooperative mutations of SETD2 in human acute leukemia. Nature Genetics, 2014, 46, 287-293.	21.4	213
6	Aberrant overexpression and function of the miR-17-92 cluster in <i>MLL</i>-rearranged acute leukemia. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 3710-3715.	7.1	141
7	CRISPR interference-based specific and efficient gene inactivation in the brain. Nature Neuroscience, 2018, 21, 447-454.	14.8	133
8	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
9	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
10	Consistent Deregulation of Gene Expression between Human and Murine <i>MLL</i> Rearrangement Leukemias. Cancer Research, 2009, 69, 1109-1116.	0.9	81
11	Population-specific genetic variants important in susceptibility to cytarabine arabinoside cytotoxicity. Blood, 2009, 113, 2145-2153.	1.4	81
12	Circulating MicroRNAs as Biomarkers for Inflammatory Diseases. MicroRNA (Shariqah, United Arab) Tj ETQqO 0 0 rgBT /Overlock 10 Tf 50	1.2	77
13	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
14	Platinum Sensitivityâ€‘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
15	Tumor-derived circulating exosomal miR-342-5p and miR-574-5p as promising diagnostic biomarkers for early-stage Lung Adenocarcinoma. International Journal of Medical Sciences, 2020, 17, 1428-1438.	2.5	33
16	Secretome of Activated Fibroblasts Induced by Exosomes for the Discovery of Biomarkers in Nonâ€‘Small Cell Lung Cancer. Small, 2021, 17, e2004750.	10.0	18
17	<i>miR</i> - <i>218</i> - <i>2</i> regulates cognitive functions in the hippocampus through complement component 3â€‘dependent modulation of synaptic vesicle release. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	17
18	An eQTL-based method identifies CTTN and ZMAT3 as pemetrexed susceptibility markers. Human Molecular Genetics, 2012, 21, 1470-1480.	2.9	16

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19	Synaptotagmin-7 is a key factor for bipolar-like behavioral abnormalities in mice. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 4392-4399.	7.1	15
20	Expression and alternative splicing of folate pathway genes in HapMap lymphoblastoid cell lines. Pharmacogenomics, 2009, 10, 549-563.	1.3	13
21	Dietary Intervention With $\alpha$ -Amylase Inhibitor in White Kidney Beans Added Yogurt Modulated Gut Microbiota to Adjust Blood Glucose in Mice. Frontiers in Nutrition, 2021, 8, 664976.	3.7	12
22	Heterogeneous Nuclear Ribonucleoprotein A1 Loads Batched Tumor-Promoting MicroRNAs Into Small Extracellular Vesicles With the Assist of Caveolin-1 in A549 Cells. Frontiers in Cell and Developmental Biology, 2021, 9, 687912.	3.7	9
23	The long noncoding RNA HOTAIRM1 controlled by AML1 enhances glucocorticoid resistance by activating RHOA/ROCK1 pathway through suppressing ARHGAP18. Cell Death and Disease, 2021, 12, 702.	6.3	8
24	Disequilibrium in chicken gut microflora with avian colibacillosis is related to microenvironment damaged by antibiotics. Science of the Total Environment, 2021, 762, 143058.	8.0	6
25	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
26	Base-excision repair and beyond "A short summary attributed to scientific achievements of Tomas Lindahl, Nobel Prize Laureate in Chemistry 2015. Science China Life Sciences, 2016, 59, 89-92.	4.9	3
27	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
28	RBUD: A New Functional Potential Analysis Approach for Whole Microbial Genome Shotgun Sequencing. Microorganisms, 2020, 8, 1563.	3.6	1
29	A promising iPS-based single-cell cloning strategy revealing signatures of somatic mutations in heterogeneous normal cells. Computational and Structural Biotechnology Journal, 2020, 18, 2326-2335.	4.1	0
30	Identification of Potential Key lncRNAs in the Context of Mouse Myeloid Differentiation by Systematic Transcriptomics Analysis. Genes, 2021, 12, 630.	2.4	0
31	Identification of Genes Abnormally Expressed in Human MLL-AF4 Leukemia.. Blood, 2006, 108, 4314-4314.	1.4	0
32	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
33	MicroRNA Expression Signatures Accurately Discriminate Acute Lymphoblastic Leukemia from Acute Myeloid Leukemia.. Blood, 2007, 110, 231-231.	1.4	0
34	MicroRNA Expression Profiles in Acute Myeloid Leukemia with Common Translocations.. Blood, 2007, 110, 3181-3181.	1.4	0
35	Identification of Genes Deregulated in Both Human and Murine MLLRearrangement Leukemias.. Blood, 2008, 112, 3359-3359.	1.4	0
36	MLL Fusion Proteins Directly Regulate a Small Set of Wild Type MLL Target Genes.. Blood, 2009, 114, 1279-1279.	1.4	0

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37	Genome Wide Location Analysis Reveals Deregulated MicroRNA Genes In MLL Rearranged Leukemic Genome. Blood, 2010, 116, 2507-2507.	1.4	0
38	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
39	Early Different Downstream Target of Glucocorticoid Receptor Contributing to Glucocorticoids Sensitivity in Kasumi-1 Cells. Blood, 2016, 128, 5132-5132.	1.4	0