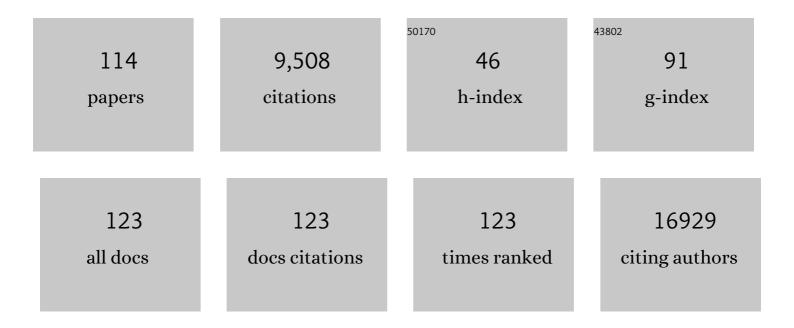
Yu Zhou

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Transcriptomic characteristics of bronchoalveolar lavage fluid and peripheral blood mononuclear cells in COVID-19 patients. Emerging Microbes and Infections, 2020, 9, 761-770.	3.0	994
2	Direct Conversion of Fibroblasts to Neurons by Reprogramming PTB-Regulated MicroRNA Circuits. Cell, 2013, 152, 82-96.	13.5	508
3	Genome-wide Analysis of PTB-RNA Interactions Reveals a Strategy Used by the General Splicing Repressor to Modulate Exon Inclusion or Skipping. Molecular Cell, 2009, 36, 996-1006.	4.5	429
4	MicroRNA Directly Enhances Mitochondrial Translation during Muscle Differentiation. Cell, 2014, 158, 607-619.	13.5	385
5	Pachytene piRNAs instruct massive mRNA elimination during late spermiogenesis. Cell Research, 2014, 24, 680-700.	5.7	344
6	SR Proteins Collaborate with 7SK and Promoter-Associated Nascent RNA to Release Paused Polymerase. Cell, 2013, 153, 855-868.	13.5	279
7	R-ChIP Using Inactive RNase H Reveals Dynamic Coupling of R-loops with Transcriptional Pausing at Gene Promoters. Molecular Cell, 2017, 68, 745-757.e5.	4.5	263
8	Genome-wide Analysis Reveals SR Protein Cooperation and Competition in Regulated Splicing. Molecular Cell, 2013, 50, 223-235.	4.5	261
9	Pre-mRNA splicing is facilitated by an optimal RNA polymerase II elongation rate. Genes and Development, 2014, 28, 2663-2676.	2.7	250
10	Pancreatic cancer-targeting exosomes for enhancing immunotherapy and reprogramming tumor microenvironment. Biomaterials, 2021, 268, 120546.	5.7	237
11	Pervasive Chromatin-RNA Binding Protein Interactions Enable RNA-Based Regulation of Transcription. Cell, 2019, 178, 107-121.e18.	13.5	224
12	The Akt-SRPK-SR Axis Constitutes a Major Pathway in Transducing EGF Signaling to Regulate Alternative Splicing in the Nucleus. Molecular Cell, 2012, 47, 422-433.	4.5	221
13	Enhanced Cardioprotection by Human Endometrium Mesenchymal Stem Cells Driven by Exosomal MicroRNA-21. Stem Cells Translational Medicine, 2017, 6, 209-222.	1.6	217
14	ALS-causative mutations in FUS/TLS confer gain and loss of function by altered association with SMN and U1-snRNP. Nature Communications, 2015, 6, 6171.	5.8	205
15	The Augmented R-Loop Is a Unifying Mechanism for Myelodysplastic Syndromes Induced by High-Risk Splicing Factor Mutations. Molecular Cell, 2018, 69, 412-425.e6.	4.5	203
16	Toxic gain of function from mutant <scp>FUS</scp> protein is crucial to trigger cell autonomous motor neuron loss. EMBO Journal, 2016, 35, 1077-1097.	3.5	187
17	Interactions between the NR2B Receptor and CaMKII Modulate Synaptic Plasticity and Spatial Learning. Journal of Neuroscience, 2007, 27, 13843-13853.	1.7	169
18	NEAT1 scaffolds RNA-binding proteins and the Microprocessor to globally enhance pri-miRNA processing. Nature Structural and Molecular Biology, 2017, 24, 816-824.	3.6	165

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19	Impact of Single-chain Fv Antibody Fragment Affinity on Nanoparticle Targeting of Epidermal Growth Factor Receptor-expressing Tumor Cells. Journal of Molecular Biology, 2007, 371, 934-947.	2.0	164
20	Coinfection with influenza A virus enhances SARS-CoV-2 infectivity. Cell Research, 2021, 31, 395-403.	5.7	164
21	Mechanisms for U2AF to define 3′ splice sites and regulate alternative splicing in the human genome. Nature Structural and Molecular Biology, 2014, 21, 997-1005.	3.6	150
22	Nuclear Matrix Factor hnRNP U/SAF-A Exerts a Global Control of Alternative Splicing by Regulating U2 snRNP Maturation. Molecular Cell, 2012, 45, 656-668.	4.5	146
23	Molecular basis for 5-carboxycytosine recognition by RNA polymerase II elongation complex. Nature, 2015, 523, 621-625.	13.7	141
24	A Translation-Activating Function of MIWI/piRNA during Mouse Spermiogenesis. Cell, 2019, 179, 1566-1581.e16.	13.5	136
25	A novel class of microRNA-recognition elements that function only within open reading frames. Nature Structural and Molecular Biology, 2018, 25, 1019-1027.	3.6	134
26	RBFox1-mediated RNA splicing regulates cardiac hypertrophy and heart failure. Journal of Clinical Investigation, 2015, 126, 195-206.	3.9	114
27	Both Decreased and Increased SRPK1 Levels Promote Cancer by Interfering with PHLPP-Mediated Dephosphorylation of Akt. Molecular Cell, 2014, 54, 378-391.	4.5	105
28	Pooled CRISPR screening identifies m ⁶ A as a positive regulator of macrophage activation. Science Advances, 2021, 7, .	4.7	102
29	SRSF2 Is Essential for Hematopoiesis, and Its Myelodysplastic Syndrome-Related Mutations Dysregulate Alternative Pre-mRNA Splicing. Molecular and Cellular Biology, 2015, 35, 3071-3082.	1.1	92
30	TDP-43 aggregation induced by oxidative stress causes global mitochondrial imbalance in ALS. Nature Structural and Molecular Biology, 2021, 28, 132-142.	3.6	92
31	Sequential regulatory loops as key gatekeepers for neuronal reprogramming in human cells. Nature Neuroscience, 2016, 19, 807-815.	7.1	88
32	GISSD: Group I Intron Sequence and Structure Database. Nucleic Acids Research, 2008, 36, D31-D37.	6.5	78
33	Bone marrow mesenchymal stem cells-derived exosomes for penetrating and targeted chemotherapy of pancreatic cancer. Acta Pharmaceutica Sinica B, 2020, 10, 1563-1575.	5.7	78
34	Repression of the Central Splicing Regulator RBFox2 Is Functionally Linked to Pressure Overload-Induced Heart Failure. Cell Reports, 2015, 10, 1521-1533.	2.9	74
35	The SARS-CoV-2 subgenome landscape and its novel regulatory features. Molecular Cell, 2021, 81, 2135-2147.e5.	4.5	72
36	Multi-strategic RNA-seq analysis reveals a high-resolution transcriptional landscape in cotton. Nature Communications, 2019, 10, 4714.	5.8	70

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37	A missense variant in FGD6 confers increased risk of polypoidal choroidal vasculopathy. Nature Genetics, 2016, 48, 640-647.	9.4	68
38	Caffeic acid ameliorates early and delayed brain injuries after focal cerebral ischemia in rats. Acta Pharmacologica Sinica, 2006, 27, 1103-1110.	2.8	66
39	RBFox2 Binds Nascent RNA to Globally Regulate Polycomb Complex 2 Targeting in Mammalian Genomes. Molecular Cell, 2016, 62, 875-889.	4.5	66
40	Versatile pathway-centric approach based on high-throughput sequencing to anticancer drug discovery. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 4609-4614.	3.3	63
41	A multiplex RNA-seq strategy to profile poly(A+) RNA: Application to analysis of transcription response and 3′ end formation. Genomics, 2011, 98, 266-271.	1.3	61
42	Spatio-temporal properties of 5-lipoxygenase expression and activation in the brain after focal cerebral ischemia in rats. Life Sciences, 2006, 79, 1645-1656.	2.0	59
43	De Novo Prediction of PTBP1 Binding and Splicing Targets Reveals Unexpected Features of Its RNA Recognition and Function. PLoS Computational Biology, 2014, 10, e1003442.	1.5	56
44	Impact of Intrinsic Affinity on Functional Binding and Biological Activity of EGFR Antibodies. Molecular Cancer Therapeutics, 2012, 11, 1467-1476.	1.9	54
45	Cyclin E in normal physiology and disease states. Trends in Cell Biology, 2021, 31, 732-746.	3.6	54
46	Internalizing Cancer Antibodies from Phage Libraries Selected on Tumor Cells and Yeast-Displayed Tumor Antigens. Journal of Molecular Biology, 2010, 404, 88-99.	2.0	53
47	Adding pyrrolysine to the <i>Escherichia coli</i> genetic code. FEBS Letters, 2007, 581, 5282-5288.	1.3	52
48	Wounding triggers MIRO-1 dependent mitochondrial fragmentation that accelerates epidermal wound closure through oxidative signaling. Nature Communications, 2020, 11, 1050.	5.8	44
49	Fluorogenic labeling and single-base resolution analysis of 5-formylcytosine in DNA. Chemical Science, 2017, 8, 7443-7447.	3.7	42
50	Distinct splicing signatures affect converged pathways in myelodysplastic syndrome patients carrying mutations in different splicing regulators. Rna, 2016, 22, 1535-1549.	1.6	40
51	JMJD6 and U2AF65 co-regulate alternative splicing in both JMJD6 enzymatic activity dependent and independent manner. Nucleic Acids Research, 2017, 45, 3503-3518.	6.5	40
52	Antitumour activity and tolerability of an EphA2-targeted nanotherapeutic in multiple mouse models. Nature Biomedical Engineering, 2019, 3, 264-280.	11.6	40
53	Transcriptional upregulation of MT2â€MMP in response to hypoxia is promoted by HIFâ€1α in cancer cells. Molecular Carcinogenesis, 2011, 50, 770-780.	1.3	37
54	Oleoylethanolamide inhibits glial activation via moudulating PPARα and promotes motor function recovery after brain ischemia. Pharmacological Research, 2019, 141, 530-540.	3.1	37

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55	Precision Engineering of an Anti-HLA-A2 Chimeric Antigen Receptor in Regulatory T Cells for Transplant Immune Tolerance. Frontiers in Immunology, 2021, 12, 686439.	2.2	37
56	PsORF: a database of small ORFs in plants. Plant Biotechnology Journal, 2020, 18, 2158-2160.	4.1	35
57	NRDE2 negatively regulates exosome functions by inhibiting MTR4 recruitment and exosome interaction. Genes and Development, 2019, 33, 536-549.	2.7	34
58	NEPdb: A Database of T-Cell Experimentally-Validated Neoantigens and Pan-Cancer Predicted Neoepitopes for Cancer Immunotherapy. Frontiers in Immunology, 2021, 12, 644637.	2.2	34
59	RBFox2-miR-34a-Jph2 axis contributes to cardiac decompensation during heart failure. Proceedings of the United States of America, 2019, 116, 6172-6180.	3.3	32
60	Exosomes derived from immunogenically dying tumor cells as a versatile tool for vaccination against pancreatic cancer. Biomaterials, 2022, 280, 121306.	5.7	32
61	GRID-seq for comprehensive analysis of global RNA–chromatin interactions. Nature Protocols, 2019, 14, 2036-2068.	5.5	31
62	Type I Interferon Regulates a Coordinated Gene Network to Enhance Cytotoxic T Cell–Mediated Tumor Killing. Cancer Discovery, 2020, 10, 382-393.	7.7	31
63	<scp>ALYREF</scp> links 3′â€end processing to nuclear export of nonâ€polyadenylated <scp>mRNA</scp> s. EMBO Journal, 2019, 38, .	3.5	30
64	Anti-MET ImmunoPET for Non–Small Cell Lung Cancer Using Novel Fully Human Antibody Fragments. Molecular Cancer Therapeutics, 2014, 13, 2607-2617.	1.9	29
65	CELF RNA binding proteins promote axon regeneration in C. elegans and mammals through alternative splicing of Syntaxins. ELife, 2016, 5, .	2.8	27
66	Atractylenolide III reduces depressive- and anxiogenic-like behaviors in rat depression models. Neuroscience Letters, 2021, 759, 136050.	1.0	27
67	Alternative polyadenylation by sequential activation of distal and proximal PolyA sites. Nature Structural and Molecular Biology, 2022, 29, 21-31.	3.6	27
68	The RNA binding protein EWS is broadly involved in the regulation of pri-miRNA processing in mammalian cells. Nucleic Acids Research, 2017, 45, 12481-12495.	6.5	26
69	The Paf1 complex transcriptionally regulates the mitochondrial-anchored protein Atg32 leading to activation of mitophagy. Autophagy, 2020, 16, 1366-1379.	4.3	26
70	SHQ1 regulation of RNA splicing is required for T-lymphoblastic leukemia cell survival. Nature Communications, 2018, 9, 4281.	5.8	24
71	MaGenDB: a functional genomics hub for Malvaceae plants. Nucleic Acids Research, 2020, 48, D1076-D1084.	6.5	23
72	Moringa oleifera seed extract protects against brain damage in both the acute and delayed stages of ischemic stroke. Experimental Gerontology, 2019, 122, 99-108.	1.2	23

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73	A fully human scFv phage display library for rapid antibody fragment reformatting. Protein Engineering, Design and Selection, 2015, 28, 307-316.	1.0	22
74	Recent advances in RNA structurome. Science China Life Sciences, 2022, 65, 1285-1324.	2.3	22
75	Regulation of splicing enhancer activities by RNA secondary structures. FEBS Letters, 2010, 584, 4401-4407.	1.3	21
76	Oncogenic miR-17/20a Forms a Positive Feed-forward Loop with the p53 Kinase DAPK3 to Promote Tumorigenesis. Journal of Biological Chemistry, 2015, 290, 19967-19975.	1.6	21
77	GoldCLIP: Gel-omitted Ligation-dependent CLIP. Genomics, Proteomics and Bioinformatics, 2018, 16, 136-143.	3.0	21
78	Actin Polymerization and ESCRT Trigger Recruitment of the Fusogens Syntaxin-2 and EFF-1 to Promote Membrane Repair in C.Âelegans. Developmental Cell, 2020, 54, 624-638.e5.	3.1	20
79	Exome Sequencing Analysis Identifies Compound Heterozygous Mutation in ABCA4 in a Chinese Family with Stargardt Disease. PLoS ONE, 2014, 9, e91962.	1.1	17
80	Immune-based mutation classification enables neoantigen prioritization and immune feature discovery in cancer immunotherapy. Oncolmmunology, 2021, 10, 1868130.	2.1	17
81	LeafNet: a tool for segmenting and quantifying stomata and pavement cells. Plant Cell, 2022, 34, 1171-1188.	3.1	17
82	Mutant RAMP2 causes primary open-angle glaucoma via the CRLR-cAMP axis. Genetics in Medicine, 2019, 21, 2345-2354.	1.1	16
83	MVIP: multi-omics portal of viral infection. Nucleic Acids Research, 2022, 50, D817-D827.	6.5	16
84	A Genetic Variant in theSKIV2LGene Is Significantly Associated With Age-Related Macular Degeneration in a Han Chinese Population. , 2013, 54, 2911.		15
85	High Throughput Identification of Monoclonal Antibodies to Membrane Bound and Secreted Proteins Using Yeast and Phage Display. PLoS ONE, 2014, 9, e111339.	1.1	15
86	Oâ€GlcNAcylation of TDPâ€43 suppresses proteinopathies and promotes TDPâ€43's mRNA splicing activity. EMBO Reports, 2021, 22, e51649.	2.0	15
87	Full-length annotation with multistrategy RNA-seq uncovers transcriptional regulation of lncRNAs in cotton. Plant Physiology, 2021, 185, 179-195.	2.3	15
88	A U2-snRNP–independent role of SF3b in promoting mRNA export. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 7837-7846.	3.3	13
89	Whole-exome sequencing reveals a novel frameshift mutation in the FAM161A gene causing autosomal recessive retinitis pigmentosa in the Indian population. Journal of Human Genetics, 2015, 60, 625-630.	1.1	12
90	Hypercholesterolemia risk-associated GPR146 is an orphan G-protein coupled receptor that regulates blood cholesterol levels in humans and mice. Cell Research, 2020, 30, 363-365.	5.7	12

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91	Exome Sequencing Identified a Recessive <i>RDH12</i> Mutation in a Family with Severe Early-Onset Retinitis Pigmentosa. Journal of Ophthalmology, 2015, 2015, 1-7.	0.6	11
92	Multiplex Analysis of PolyA-Linked Sequences (MAPS): An RNA-Seq Strategy to Profile Poly(A+) RNA. Methods in Molecular Biology, 2014, 1125, 169-178.	0.4	10
93	Knockout of glutathione peroxidase 5 down-regulates the piRNAs in the caput epididymidis of aged mice. Asian Journal of Andrology, 2020, 22, 590.	0.8	10
94	Overexpression of membrane-type 2 matrix metalloproteinase induced by hypoxia-inducible factor-1α in pancreatic cancer: Implications for tumor progression and prognosis. Molecular and Clinical Oncology, 2014, 2, 973-981.	0.4	9
95	An accurate quantitative method for screening effective siRNA probes targeting a Hepatitis B virus transcript in single living cells. Biochemical and Biophysical Research Communications, 2008, 367, 866-873.	1.0	8
96	Defective minor spliceosomes induce SMA-associated phenotypes through sensitive intron-containing neural genes in Drosophila. Nature Communications, 2020, 11, 5608.	5.8	8
97	SLC35B2 Acts in a Dual Role in the Host Sulfation Required for EV71 Infection. Journal of Virology, 2022, 96, e0204221.	1.5	8
98	Context-dependent modulation of Pol II CTD phosphatase SSUP-72 regulates alternative polyadenylation in neuronal development. Genes and Development, 2015, 29, 2377-2390.	2.7	7
99	A Novel CRYBB2 Stopgain Mutation Causing Congenital Autosomal Dominant Cataract in a Chinese Family. Journal of Ophthalmology, 2016, 2016, 1-8.	0.6	7
100	Effect of Moringa oleifera stem extract on hydrogen peroxide-induced opacity of cultured mouse lens. BMC Complementary and Alternative Medicine, 2019, 19, 144.	3.7	7
101	Biomarkers and Immune Repertoire Metrics Identified by Peripheral Blood Transcriptomic Sequencing Reveal the Pathogenesis of COVID-19. Frontiers in Immunology, 2021, 12, 677025.	2.2	7
102	Whole exome sequencing identified novel CRB1 mutations in Chinese and Indian populations with autosomal recessive retinitis pigmentosa. Scientific Reports, 2016, 6, 33681.	1.6	6
103	HCV dsRNA-Activated Macrophages Inhibit HCV Replication in Hepatocytes. Hepatitis Monthly, 2015, 15, e29282.	0.1	6
104	Phytochemical wedelolactone reverses obesity by prompting adipose browning through SIRT1/AMPK/ PPARα pathway via targeting nicotinamide N-methyltransferase. Phytomedicine, 2022, 94, 153843.	2.3	6
105	Characterization of the replicon of a 51-kb native plasmid from the gram-positive bacterium Leifsonia xyli subsp. cynodontis. FEMS Microbiology Letters, 2004, 236, 33-39.	0.7	6
106	Flexible RNA design under structure and sequence constraints using formal languages. , 2013, , .		5
107	Identification of Key Genes and Pathways Associated with Age-Related Macular Degeneration. Journal of Ophthalmology, 2020, 2020, 1-10.	0.6	5
108	ASER: Animal Sex Reversal Database. Genomics, Proteomics and Bioinformatics, 2021, 19, 873-881.	3.0	5

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109	Efficacy of amiodarone and lidocaine for preventing ventricular fibrillation after aortic cross-clamp release in open heart surgery: a meta-analysis of randomized controlled trials. Journal of Zhejiang University: Science B, 2017, 18, 1113-1122.	1.3	4
110	Discovery of internalizing antibodies to basal breast cancer cells. Protein Engineering, Design and Selection, 2018, 31, 17-28.	1.0	4
111	Deletion of Asrgl1 Leads to Photoreceptor Degeneration in Mice. Frontiers in Cell and Developmental Biology, 2021, 9, 783547.	1.8	4
112	Role of Apg-1 in HSF1 activation and bortezomib sensitivity in myeloma cells. Experimental Hematology, 2020, 81, 50-59.	0.2	2
113	ILF3 represses repeat-derived microRNAs targeting RIG-I mediated type I interferon response. Journal of Molecular Biology, 2022, 434, 167469.	2.0	2
114	Coupling of Transcription with mRNA Processing in time and Space. FASEB Journal, 2015, 29, 238.1.	0.2	0