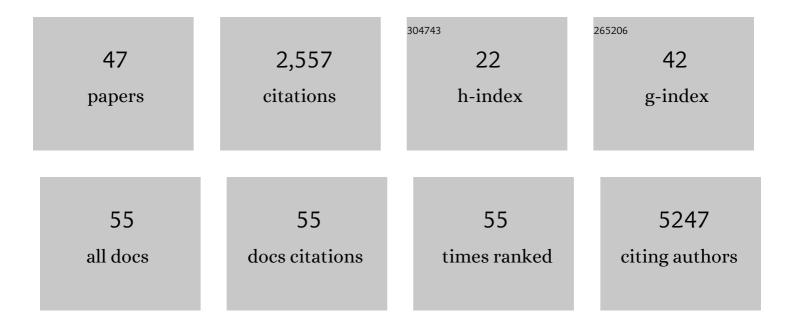
Yuliang Wang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Amino acid primed mTOR activity is essential for heart regeneration. IScience, 2022, 25, 103574.	4.1	15
2	The Alzheimer's gene SORL1 is a regulator of endosomal traffic and recycling in human neurons. Cellular and Molecular Life Sciences, 2022, 79, 162.	5.4	52
3	dCas9 fusion to computer-designed PRC2 inhibitor reveals functional TATA box in distal promoter region. Cell Reports, 2022, 38, 110457.	6.4	12
4	Interplay between orphan nuclear receptors and androgen receptor-dependent or-independent growth signalings in prostate cancer. Molecular Aspects of Medicine, 2021, 78, 100921.	6.4	7
5	Orphan nuclear receptors as regulators of intratumoral androgen biosynthesis in castration-resistant prostate cancer. Oncogene, 2021, 40, 2625-2634.	5.9	19
6	University of Washington Nathan Shock Center: innovation to advance aging research. GeroScience, 2021, 43, 2161-2165.	4.6	1
7	Podocyte Aging: Why and How Getting Old Matters. Journal of the American Society of Nephrology: JASN, 2021, , ASN.2021-05-0614.	6.1	1
8	Podocyte Aging: Why and How Getting Old Matters. Journal of the American Society of Nephrology: JASN, 2021, 32, 2697-2713.	6.1	28
9	Genetic Polymorphisms of Very Important Pharmacogene Variants in the Blang Population from Yunnan Province in China. Pharmacogenomics and Personalized Medicine, 2021, Volume 14, 1647-1660.	0.7	0
10	Conserved Epigenetic Regulatory Logic Infers Genes Governing Cell Identity. Cell Systems, 2020, 11, 625-639.e13.	6.2	31
11	Targeting prostate cancer stem-like cells by an immunotherapeutic platform based on immunogenic peptide-sensitized dendritic cells-cytokine-induced killer cells. Stem Cell Research and Therapy, 2020, 11, 123.	5.5	16
12	Global transcriptomic changes occur in aged mouse podocytes. Kidney International, 2020, 98, 1160-1173.	5.2	23
13	Spatial modeling of prostate cancer metabolic gene expression reveals extensive heterogeneity and selective vulnerabilities. Scientific Reports, 2020, 10, 3490.	3.3	43
14	Metabolic Control over mTOR-Dependent Diapause-like State. Developmental Cell, 2020, 52, 236-250.e7.	7.0	79
15	Nuclear receptor ERRα contributes to castration-resistant growth of prostate cancer via its regulation of intratumoral androgen biosynthesis. Theranostics, 2020, 10, 4201-4216.	10.0	20
16	Towards understanding androgen receptor-independent prostate cancer: an evolving paradigm. Translational Cancer Research, 2020, 9, 415-417.	1.0	2
17	TFPa/HADHA is required for fatty acid beta-oxidation and cardiolipin re-modeling in human cardiomyocytes. Nature Communications, 2019, 10, 4671.	12.8	77
18	Genome-bound enzymes as epigenetic drug targets in cancer. Epigenomics, 2019, 11, 1463-1467.	2.1	0

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19	Fatty Acids Enhance the Maturation of Cardiomyocytes Derived from Human Pluripotent Stem Cells. Stem Cell Reports, 2019, 13, 657-668.	4.8	187
20	PIXUL-ChIP: integrated high-throughput sample preparation and analytical platform for epigenetic studies. Nucleic Acids Research, 2019, 47, e69-e69.	14.5	16
21	Patterned human microvascular grafts enable rapid vascularization and increase perfusion in infarcted rat hearts. Nature Communications, 2019, 10, 584.	12.8	100
22	Metabolism as an early predictor of DPSCs aging. Scientific Reports, 2019, 9, 2195.	3.3	26
23	microRNAs Regulating Human and Mouse NaÃ ⁻ ve Pluripotency. International Journal of Molecular Sciences, 2019, 20, 5864.	4.1	11
24	Inducible CRISPR genome editing platform in naive human embryonic stem cells reveals JARID2 function in self-renewal. Cell Cycle, 2018, 17, 00-00.	2.6	13
25	Orphan nuclear receptor TLX contributes to androgen insensitivity in castration-resistant prostate cancer via its repression of androgen receptor transcription. Oncogene, 2018, 37, 3340-3355.	5.9	20
26	Single-Cell Transcriptomic Analysis of Cardiac Differentiation from Human PSCs Reveals HOPX-Dependent Cardiomyocyte Maturation. Cell Stem Cell, 2018, 23, 586-598.e8.	11.1	215
27	Development of a novel and economical agar-based non-adherent three-dimensional culture method for enrichment of cancer stem-like cells. Stem Cell Research and Therapy, 2018, 9, 243.	5.5	48
28	DNA methylation yields epigenetic clues into the diabetic nephropathy of Pima Indians. Kidney International, 2018, 93, 1272-1275.	5.2	9
29	Human Organ-Specific Endothelial Cell Heterogeneity. IScience, 2018, 4, 20-35.	4.1	181
30	Nuclear receptor ERRα and transcription factor ERG form a reciprocal loop in the regulation of TMPRSS2:ERG fusion gene in prostate cancer. Oncogene, 2018, 37, 6259-6274.	5.9	36
31	LRH-1 drives hepatocellular carcinoma partially through induction of c-myc and cyclin E1, and suppression of p21. Cancer Management and Research, 2018, Volume 10, 2389-2400.	1.9	24
32	Sex differences in transcriptomic profiles in aged kidney cells of renin lineage. Aging, 2018, 10, 606-621.	3.1	12
33	Transcriptomic, proteomic, and metabolomic landscape of positional memory in the caudal fin of zebrafish. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E717-E726.	7.1	81
34	Gene-Edited Human Kidney Organoids Reveal Mechanisms of Disease in Podocyte Development. Stem Cells, 2017, 35, 2366-2378.	3.2	101
35	First critical repressive H3K27me3 marks in embryonic stem cells identified using designed protein inhibitor. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 10125-10130.	7.1	39
36	Chromatin and Transcriptional Analysis of Mesoderm Progenitor Cells Identifies HOPX as a Regulator of Primitive Hematopoiesis. Cell Reports, 2017, 20, 1597-1608.	6.4	50

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37	Nkx2.5 marks angioblasts that contribute to hemogenic endothelium of the endocardium and dorsal aorta. ELife, 2017, 6, .	6.0	27
38	Metabolic remodeling in early development and cardiomyocyte maturation. Seminars in Cell and Developmental Biology, 2016, 52, 84-92.	5.0	62
39	Integrated Genomic Analysis of Diverse Induced Pluripotent Stem Cells from the Progenitor Cell Biology Consortium. Stem Cell Reports, 2016, 7, 110-125.	4.8	101
40	The emerging roles of orphan nuclear receptors in prostate cancer. Biochimica Et Biophysica Acta: Reviews on Cancer, 2016, 1866, 23-36.	7.4	23
41	The metabolome regulates the epigenetic landscape during naive-to-primed human embryonic stem cellÂtransition. Nature Cell Biology, 2015, 17, 1523-1535.	10.3	360
42	Measuring the Effect of Inter-Study Variability on Estimating Prediction Error. PLoS ONE, 2014, 9, e110840.	2.5	19
43	Multi-study Integration of Brain Cancer Transcriptomes Reveals Organ-Level Molecular Signatures. PLoS Computational Biology, 2013, 9, e1003148.	3.2	16
44	Reconstruction of genome-scale metabolic models for 126 human tissues using mCADRE. BMC Systems Biology, 2012, 6, 153.	3.0	239
45	Molecular signatures from omics data: From chaos to consensus. Biotechnology Journal, 2012, 7, 946-957.	3.5	101
46	Cardiac Directed Differentiation Using Small Molecule WNT Modulation at Single-Cell Resolution. SSRN Electronic Journal, 0, , .	0.4	0
47	Computer Designed PRC2 Inhibitor, EBdCas9, Reveals Functional TATA Boxes in Distal Promoter	0.4	0