Shengli Li

List of Publications by Year in descending order

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233421 304743 2,201 60 22 45 citations h-index g-index papers 63 63 63 3579 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Dominant mutations in the severe acute respiratory syndrome coronavirusâ€2 genome challenge polymerase chain reaction detection. Clinical and Translational Discovery, 2022, 2, e23.	0.5	O
2	LncPep: A Resource of Translational Evidences for IncRNAs. Frontiers in Cell and Developmental Biology, 2022, 10, 795084.	3.7	7
3	Editorial: Transcriptional Regulation in Metabolism and Immunology. Frontiers in Genetics, 2022, 13, 845697.	2.3	1
4	SRTdb: an omnibus for human tissue and cancer-specific RNA transcripts. Biomarker Research, 2022, 10, 27.	6.8	5
5	Docetaxel resistance-derived LINC01085 contributes to the immunotherapy of hormone-independent prostate cancer by activating the STING/MAVS signaling pathway. Cancer Letters, 2022, 545, 215829.	7.2	8
6	HeRA: an atlas of enhancer RNAs across human tissues. Nucleic Acids Research, 2021, 49, D932-D938.	14.5	27
7	Integrated Characterization of IncRNA-Immune Interactions in Prostate Cancer. Frontiers in Cell and Developmental Biology, 2021, 9, 641891.	3.7	14
8	Optimized protocol for an inducible rat model of liver tumor with chronic hepatocellular injury, inflammation, fibrosis, and cirrhosis. STAR Protocols, 2021, 2, 100353.	1.2	4
9	Single-cell transcriptomics reveals the landscape of intra-tumoral heterogeneity and transcriptional activities of ECs in CC. Molecular Therapy - Nucleic Acids, 2021, 24, 682-694.	5.1	51
10	Eyelid Lymphatics. Journal of Craniofacial Surgery, 2021, Publish Ahead of Print, 2878-2882.	0.7	0
11	Retrotransposons: Jump to Cancer?. Trends in Cancer, 2021, 7, 577-579.	7.4	2
12	Alternative polyadenylation-associated loci interpret human traits and diseases. Trends in Genetics, 2021, 37, 773-775.	6.7	4
13	Integrative analysis reveals clinically relevant molecular fingerprints in pancreatic cancer. Molecular Therapy - Nucleic Acids, 2021, 26, 11-21.	5.1	3
14	Maximizing the Utility of Transcriptomics Data in Inflammatory Skin Diseases. Frontiers in Immunology, 2021, 12, 761890.	4.8	9
15	Dissecting the single-cell transcriptome underlying chronic liver injury. Molecular Therapy - Nucleic Acids, 2021, 26, 1364-1373.	5.1	15
16	Y-Shaped En-Bloc Mortise-Tenon Rhinoplasty Technique. Aesthetic Surgery Journal, 2020, 40, NP8-NP20.	1.6	5
17	APAatlas: decoding alternative polyadenylation across human tissues. Nucleic Acids Research, 2020, 48, D34-D39.	14.5	41
18	Resolving Spliceosomal Malfunctions Advances RNA-Based Therapeutics. Trends in Molecular Medicine, 2020, 26, 135-137.	6.7	1

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19	Splicing Regulator p54nrb/Non–POU Domain–Containing Octamerâ€Binding Protein Enhances Carcinogenesis Through Oncogenic Isoform Switch of MYC Box–Dependent Interacting Protein 1 in Hepatocellular Carcinoma. Hepatology, 2020, 72, 548-568.	7.3	40
20	Characterization of the dual functional effects of heat shock proteins (HSPs) in cancer hallmarks to aid development of HSP inhibitors. Genome Medicine, 2020, 12, 101.	8.2	31
21	3D Spheroids Propel Tumor Characterization. Trends in Cancer, 2020, 6, 622-624.	7.4	6
22	The Mutational and Transcriptional Landscapes of Hepatocarcinogenesis in a Rat Model. IScience, 2020, 23, 101690.	4.1	12
23	Our Experiences with Plastic and Reconstructive Surgery Procedures during Coronavirus Disease 2019 Pandemic. Plastic and Reconstructive Surgery - Global Open, 2020, 8, e2868.	0.6	8
24	The Roles of Podoplanin-Positive/Podoplanin-Negative Cells from Adipose-Derived Stem Cells in Lymphatic Regeneration. Plastic and Reconstructive Surgery, 2020, 145, 420-431.	1.4	11
25	The genetic and pharmacogenomic landscape of snoRNAs in human cancer. Molecular Cancer, 2020, 19, 108.	19.2	17
26	RBP EIF2S2 Promotes Tumorigenesis and Progression by Regulating MYC-Mediated Inhibition via FHIT-Related Enhancers. Molecular Therapy, 2020, 28, 1105-1118.	8.2	37
27	The Potential Regulatory Roles of Circular RNAs in Tumor Immunology and Immunotherapy. Frontiers in Immunology, 2020, 11, 617583.	4.8	20
28	Tet2 Deficiency Rejuvenates Hematopoietic Stem and Progenitor Cells during Ageing. Blood, 2020, 136, 37-37.	1.4	0
29	Transcriptomeâ€Wide Analysis Reveals the Landscape of Aberrant Alternative Splicing Events in Liver Cancer. Hepatology, 2019, 69, 359-375.	7.3	86
30	Comprehensive characterization of circular RNAs in ~ 1000 human cancer cell lines. Genome Medicine, 2019, 11, 55.	8.2	116
31	ILâ€7 enhances the differentiation of adiposeâ€derived stem cells toward lymphatic endothelial cells through AKT signaling. Cell Biology International, 2019, 43, 394-401.	3.0	14
32	Molecular Treasures of Cancer Cell Lines. Trends in Molecular Medicine, 2019, 25, 657-659.	6.7	9
33	Circular RNAs as promising biomarkers in cancer: detection, function, and beyond. Genome Medicine, 2019, 11, 15.	8.2	57
34	Single-cell reconstruction of differentiation trajectory reveals a critical role of ETS1 in human cardiac lineage commitment. BMC Biology, 2019, 17, 89.	3.8	31
35	Novel Hybrid Dextran-Gadolinium Nanoparticles as High-relaxivity T1 Magnetic Resonance Imaging Contrast Agent for Mapping the Sentinel Lymph Node. Journal of Computer Assisted Tomography, 2019, 43, 350-357.	0.9	6
36	Exosomes Are Comparable to Source Adipose Stem Cells in Fat Graft Retention with Up-Regulating Early Inflammation and Angiogenesis. Plastic and Reconstructive Surgery, 2019, 144, 816e-827e.	1.4	60

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37	Combinatorial epigenetic regulation of non-coding RNAs has profound effects on oncogenic pathways in breast cancer subtypes. Briefings in Bioinformatics, 2018, 19, bbw099.	6.5	15
38	The LINC01138 drives malignancies via activating arginine methyltransferase 5 in hepatocellular carcinoma. Nature Communications, 2018 , 9 , 1572 .	12.8	157
39	A LIN28B Tumor-Specific Transcript in Cancer. Cell Reports, 2018, 22, 2016-2025.	6.4	22
40	Scalable and cleavable polysaccharide nanocarriers for the delivery of chemotherapy drugs. Acta Biomaterialia, 2018, 72, 206-216.	8.3	21
41	Genome-wide analysis reveals that exon methylation facilitates its selective usage in the human transcriptome. Briefings in Bioinformatics, 2018, 19, 754-764.	6.5	52
42	Long noncoding RNA TSLNC8 is a tumor suppressor that inactivates the interleukinâ€6/STAT3 signaling pathway. Hepatology, 2018, 67, 171-187.	7.3	183
43	exoRBase: a database of circRNA, IncRNA and mRNA in human blood exosomes. Nucleic Acids Research, 2018, 46, D106-D112.	14.5	415
44	MicroRNA-129-5p Regulates Glycolysis and Cell Proliferation by Targeting the Glucose Transporter SLC2A3 in Gastric Cancer Cells. Frontiers in Pharmacology, 2018, 9, 502.	3 . 5	59
45	Overexpression of Prox1 Induces the Differentiation of Human Adipose-Derived Stem Cells into Lymphatic Endothelial-Like Cells <i>In Vitro</i> . Cellular Reprogramming, 2017, 19, 54-63.	0.9	18
46	Transcriptomic analyses of <scp>RNA</scp> â€binding proteins reveal <i><scp>elF</scp>3c</i> promotes cell proliferation in hepatocellular carcinoma. Cancer Science, 2017, 108, 877-885.	3.9	38
47	MetaLnc9 Facilitates Lung Cancer Metastasis via a PGK1-Activated AKT/mTOR Pathway. Cancer Research, 2017, 77, 5782-5794.	0.9	139
48	Profiling and Co-expression Network Analysis of Learned Helplessness Regulated mRNAs and IncRNAs in the Mouse Hippocampus. Frontiers in Molecular Neuroscience, 2017, 10, 454.	2.9	18
49	Magnetic resonance lymphangiography for the assessment of the lymphatic system in a lymphatic malformation patient undergoing sclerotherapy. Journal of Dermatology, 2016, 43, 981-983.	1.2	2
50	Genome-wide DNA methylome analysis reveals epigenetically dysregulated non-coding RNAs in human breast cancer. Scientific Reports, 2015, 5, 8790.	3.3	54
51	Functional dissection of virus–human crosstalk mediated by miRNAs based on the VmiReg database. Molecular BioSystems, 2015, 11, 1319-1328.	2.9	12
52	Comparative epigenetic analyses reveal distinct patterns of oncogenic pathways activation in breast cancer subtypes. Human Molecular Genetics, 2014, 23, 5378-5393.	2.9	31
53	Dextran-based fluorescent nanoprobes for sentinel lymph node mapping. Biomaterials, 2014, 35, 8227-8235.	11.4	34
54	MR lymphangiography for the assessment of the lymphatic system in a primary penoscrotal lymphedema patient undergoing surgical management. Journal of Plastic, Reconstructive and Aesthetic Surgery, 2014, 67, e173-e175.	1.0	4

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55	Integrating analysis reveals microRNA-mediated pathway crosstalk among Crohn's disease, ulcerative colitis and colorectal cancer. Molecular BioSystems, 2014, 10, 2317.	2.9	27
56	Closure of large defects after microcystic lymphatic malformations using lateral intercostal artery perforator flap. Journal of Plastic, Reconstructive and Aesthetic Surgery, 2014, 67, 1230-1236.	1.0	8
57	Emergency management of traumatic total scalp avulsion with microsurgical replantation. Ulusal Travma Ve Acil Cerrahi Dergisi, 2014, 20, 66-70.	0.3	13
58	Comprehensive analysis of the functional microRNA–mRNA regulatory network identifies miRNA signatures associated with glioma malignant progression. Nucleic Acids Research, 2013, 41, e203-e203.	14.5	112
59	Characterizing Genes with Distinct Methylation Patterns in the Context of Protein-Protein Interaction Network: Application to Human Brain Tissues. PLoS ONE, 2013, 8, e65871.	2.5	8
60	Editorial: DNA Methylation Dynamics and Human Diseases. Frontiers in Cell and Developmental Biology, 0, 10, .	3.7	0