Silvia Liu

List of Publications by Year in descending order

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414034 393982 1,274 48 19 32 citations h-index g-index papers 49 49 49 1953 citing authors all docs docs citations times ranked

#	Article	IF	CITATIONS
1	Engineering osteoarthritic cartilage model through differentiating senescent human mesenchymal stem cells for testing disease-modifying drugs. Science China Life Sciences, 2022, 65, 309-327.	2.3	9
2	Oncogenic Activity of Solute Carrier Family 45 Member 2 and Alphaâ€Methylacylâ€Coenzyme A Racemase Gene Fusion Is Mediated by Mitogenâ€Activated Protein Kinase. Hepatology Communications, 2022, 6, 209-222.	2.0	3
3	Transcriptome and Exome Analyses of Hepatocellular Carcinoma Reveal Patterns to Predict Cancer Recurrence in Liver Transplant Patients. Hepatology Communications, 2022, 6, 710-727.	2.0	9
4	Four-dimensional nuclear speckle phase separation dynamics regulate proteostasis. Science Advances, 2022, 8, eabl4150.	4.7	16
5	Changes in beta-catenin expression and activation during progression of primary sclerosing cholangitis predict disease recurrence. Scientific Reports, 2022, 12, 206.	1.6	2
6	Transcriptomic profiling of a multiethnic pediatric NAFLD cohort reveals genes and pathways associated with disease. Hepatology Communications, 2022, 6, 1598-1610.	2.0	6
7	Differential responses to immune checkpoint inhibitor dictated by pre-existing differential immune profiles in squamous cell carcinomas caused by same initial oncogenic drivers. Journal of Experimental and Clinical Cancer Research, 2022, 41, 123.	3.5	10
8	ADAR1 RNA editing regulates endothelial cell functions via the MDA-5 RNA sensing signaling pathway. Life Science Alliance, 2022, 5, e202101191.	1.3	7
9	Human Mesenchymal Stem Cellâ€Derived Miniature Joint System for Disease Modeling and Drug Testing. Advanced Science, 2022, 9, e2105909.	5.6	22
10	NOTCH-YAP1/TEAD-DNMT1 Axis Drives Hepatocyte Reprogramming Into Intrahepatic Cholangiocarcinoma. Gastroenterology, 2022, 163, 449-465.	0.6	23
11	In the Absence of YAP, TAZ Contributes to Hepatocyte Adaptation in Chronic Cholestasis in Females. FASEB Journal, 2022, 36, .	0.2	О
12	Functional analysis of the <i>Vsx2</i> super-enhancer uncovers distinct <i>cis</i> -regulatory circuits controlling <i>Vsx2</i> expression during retinogenesis. Development (Cambridge), 2022, 149, .	1.2	2
13	Gene Regulatory Network Analysis and Engineering Directs Development and Vascularization of Multilineage Human Liver Organoids. Cell Systems, 2021, 12, 41-55.e11.	2.9	59
14	Gestational Age-Specific Complete Blood Count Signatures in Necrotizing Enterocolitis. Frontiers in Pediatrics, 2021, 9, 604899.	0.9	8
15	Targeted transcriptome analysis using synthetic long read sequencing uncovers isoform reprograming in the progression of colon cancer. Communications Biology, 2021, 4, 506.	2.0	10
16	Dual \hat{I}^2 -Catenin and \hat{I}^3 -Catenin Loss in Hepatocytes Impacts Their Polarity through Altered Transforming Growth Factor- \hat{I}^2 and Hepatocyte Nuclear Factor 4 \hat{I} ± Signaling. American Journal of Pathology, 2021, 191, 885-901.	1.9	3
17	Nuclear factor erythroid 2–related factor 2 and βâ€Catenin Coactivation in Hepatocellular Cancer: Biological and Therapeutic Implications. Hepatology, 2021, 74, 741-759.	3.6	32
18	Detection of fusion gene transcripts in the blood samples of prostate cancer patients. Scientific Reports, 2021, 11, 16995.	1.6	7

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19	WNT7B Regulates Cholangiocyte Proliferation and Function During Murine Cholestasis. Hepatology Communications, 2021, 5, 2019-2034.	2.0	9
20	The hepatocyte growth factor/c-met pathway is a key determinant of the fibrotic kidney local microenvironment. IScience, 2021, 24, 103112.	1.9	5
21	Pten-NOLC1 fusion promotes cancers involving MET and EGFR signalings. Oncogene, 2021, 40, 1064-1076.	2.6	9
22	Serum integrative omics reveals the landscape of human diabetic kidney disease. Molecular Metabolism, 2021, 54, 101367.	3.0	20
23	The Thyromimetic Sobetirome (GC-1) Alters Bile Acid Metabolism in a Mouse Model of Hepatic Cholestasis. American Journal of Pathology, 2020, 190, 1006-1017.	1.9	3
24	Inhibition of Estrogen Sulfotransferase (SULT1E1/EST) Ameliorates Ischemic Acute Kidney Injury in Mice. Journal of the American Society of Nephrology: JASN, 2020, 31, 1496-1508.	3.0	12
25	Recent Advances in Computer-Assisted Algorithms for Cell Subtype Identification of Cytometry Data. Frontiers in Cell and Developmental Biology, 2020, 8, 234.	1.8	19
26	Blocking integrin $\hat{l}\pm4\hat{l}^2$ 7-mediated CD4 T cell recruitment to the intestine and liver protects mice from western diet-induced non-alcoholic steatohepatitis. Journal of Hepatology, 2020, 73, 1013-1022.	1.8	47
27	Impaired Bile Secretion Promotes Hepatobiliary Injury in Sickle Cell Disease. Hepatology, 2020, 72, 2165-2181.	3.6	12
28	Transcriptomic and functional studies reveal undermined chemotactic and angiostimulatory properties of aged microglia during stroke recovery. Journal of Cerebral Blood Flow and Metabolism, 2020, 40, S81-S97.	2.4	29
29	12-h clock regulation of genetic information flow by XBP1s. PLoS Biology, 2020, 18, e3000580.	2.6	46
30	Concomitant NFE2L2 and CTNNB1 mutations in a subset of HCC patients: Synergy between Nrf2 and Wnt pathway in hepatocarcinogenesis. FASEB Journal, 2020, 34, 1-1.	0.2	0
31	Diabetic kidney diseases revisited: A new perspective for a new era. Molecular Metabolism, 2019, 30, 250-263.	3.0	122
32	Inhibiting Glutamine-Dependent mTORC1 Activation Ameliorates Liver Cancers Driven by \hat{l}^2 -Catenin Mutations. Cell Metabolism, 2019, 29, 1135-1150.e6.	7.2	92
33	MetaOmics: analysis pipeline and browser-based software suite for transcriptomic meta-analysis. Bioinformatics, 2019, 35, 1597-1599.	1.8	37
34	Detection of fusion transcripts in the serum samples of patients with hepatocellular carcinoma. Oncotarget, 2019, 10, 3352-3360.	0.8	20
35	NFE2L2 synergizes with betaâ€catenin gene mutations to induce HCC in patients and mice. FASEB Journal, 2019, 33, 126.12.	0.2	1
36	FGF19 and Met coâ€activation in murine liver induces HCC: Biological and clinical relevance. FASEB Journal, 2019, 33, 496.36.	0.2	0

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37	Detection of fusion transcripts in the serum samples of patients with hepatocellular carcinoma. Oncotarget, 2019, 10, 3352-3360.	0.8	10
38	Combined Systemic Disruption of MET and Epidermal Growth Factor Receptor Signaling Causes Liver Failure in Normal Mice. American Journal of Pathology, 2018, 188, 2223-2235.	1.9	20
39	MAN2A1–FER Fusion Gene Is Expressed by Human Liver and Other Tumor Types and Has Oncogenic Activity in Mice. Gastroenterology, 2017, 153, 1120-1132.e15.	0.6	44
40	Targeting genomic rearrangements in tumor cells through Cas9-mediated insertion of a suicide gene. Nature Biotechnology, 2017, 35, 543-550.	9.4	91
41	Meta-analytic framework for liquid association. Bioinformatics, 2017, 33, 2140-2147.	1.8	9
42	Combined systemic elimination of MET and epidermal growth factor receptor signaling completely abolishes liver regeneration and leads to liver decompensation. Hepatology, 2016, 64, 1711-1724.	3.6	89
43	Comprehensive evaluation of fusion transcript detection algorithms and a meta-caller to combine top performing methods in paired-end RNA-seq data. Nucleic Acids Research, 2016, 44, e47-e47.	6.5	141
44	Meta-Analytic Framework for Sparse <i>K</i> -Means to Identify Disease Subtypes in Multiple Transcriptomic Studies. Journal of the American Statistical Association, 2016, 111, 27-42.	1.8	22
45	Genomic Copy Number Variations in the Genomes of Leukocytes Predict Prostate Cancer Clinical Outcomes. PLoS ONE, 2015, 10, e0135982.	1.1	7
46	Discovery and Classification of Fusion Transcripts in Prostate Cancer and Normal Prostate Tissue. American Journal of Pathology, 2015, 185, 1834-1845.	1.9	26
47	Oncogenic Activity of miR-650 in Prostate Cancer Is Mediated by Suppression of CSR1 Expression. American Journal of Pathology, 2015, 185, 1991-1999.	1.9	41
48	Novel Fusion Transcripts Associate with Progressive Prostate Cancer. American Journal of Pathology, 2014, 184, 2840-2849.	1.9	62