Sibo Zhu

List of Publications by Year in descending order

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| 1 2 | Calcitriol ameliorates renal injury with high-salt diet-induced hypertension by upregulating GLIS2 expression and AMPK/mTOR-regulated autophagy. Gene, 2022, 820, 146239. Environmental risk factors and genetic markers of Kaposi's sarcomaâ€associated herpesvirus infection among Uygur population in Xinjiang, China. Journal of Medical Virology, 2022, , . Calcitriol ameliorates damage in high-salt diet-induced hypertension: Evidence of communication with the gut–kidney axis. Experimental Biology and Medicine, 2022, 247, 624-640. | 2.2 5.0 | 7 |
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| | among Uygur population in Xinjiang, China. Journal of Medical Virology, 2022, , . Calcitriol ameliorates damage in high-salt diet-induced hypertension: Evidence of communication with | 5.0 | 1 |
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| 3 | | 2.4 | 3 |
| 4 | Single-cell transcriptomic analysis reveals the critical molecular pattern of UV-induced cutaneous squamous cell carcinoma. Cell Death and Disease, 2022, 13, 23. | 6.3 | 19 |
| 5 | Lifestyle, multiâ€omics features, and preclinical dementia among Chinese: The Taizhou Imaging Study. Alzheimer's and Dementia, 2021, 17, 18-28. | 0.8 | 20 |
| 6 | Comprehensive analysis of metastatic gastric cancer tumour cells using single-cell RNA-seq. Scientific Reports, 2021, 11, 1141. | 3.3 | 30 |
| 7 | Inhibition of phospholipases suppresses progression of psoriasis through modulation of inflammation. Experimental Biology and Medicine, 2021, 246, 1253-1262. | 2.4 | 7 |
| 8 | Loss of retinoic acid receptor-related receptor alpha (Rorα) promotes the progression of UV-induced cSCC. Cell Death and Disease, 2021, 12, 247. | 6.3 | 8 |
| 9 | The gut microbiome in subclinical atherosclerosis: a population-based multiphenotype analysis. Rheumatology, 2021, 61, 258-269. | 1.9 | 13 |
| 10 | Functional consequences of a rare missense BARD1 c.403G>A germline mutation identified in a triple-negative breast cancer patient. Breast Cancer Research, 2021, 23, 53. | 5.0 | 4 |
| 11 | TRPM2 promotes pancreatic cancer by PKC/MAPK pathway. Cell Death and Disease, 2021, 12, 585. | 6.3 | 30 |
| 12 | A global metagenomic map of urban microbiomes and antimicrobial resistance. Cell, 2021, 184, 3376-3393.e17. | 28.9 | 164 |
| 13 | Association between oral microflora and gastrointestinal tumors (Review). Oncology Reports, 2021, 46, . | 2.6 | 6 |
| 14 | Relating the transcriptome and microbiome by paired terminal ileal Crohn disease. IScience, 2021, 24, 102516. | 4.1 | 3 |
| 15 | Prediction of response after cardiac resynchronization therapy with machine learning. International Journal of Cardiology, 2021, 344, 120-126. | 1.7 | 10 |
| 16 | Single-cell analysis reveals the intra-tumor heterogeneity and identifies MLXIPL as a biomarker in the cellular trajectory of hepatocellular carcinoma. Cell Death Discovery, 2021, 7, 14. | 4.7 | 23 |
| 17 | Negative correlation between endoglin levels and coronary atherosclerosis. Lipids in Health and Disease, 2021, 20, 127. | 3.0 | 5 |
| 18 | Risk factors and socio-economic burden in pancreatic ductal adenocarcinoma operation: a machine learning based analysis. BMC Cancer, 2020, 20, 1161. | 2.6 | 12 |

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| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 19 | Mechanisms and Future of Non-Small Cell Lung Cancer Metastasis. Frontiers in Oncology, 2020, 10, 585284. | 2.8 | 42 |
| 20 | Critical steps to tumor metastasis: alterations of tumor microenvironment and extracellular matrix in the formation of pre-metastatic and metastatic niche. Cell and Bioscience, 2020, 10, 89. | 4.8 | 36 |
| 21 | Changes in plasma bile acids are associated with gallbladder stones and polyps. BMC Gastroenterology, 2020, 20, 363. | 2.0 | 11 |
| 22 | Transcriptomic analysis of the mechanisms of alleviating renal interstitial fibrosis using the traditional Chinese medicine Kangxianling in a rat model. Scientific Reports, 2020, 10, 10682. | 3.3 | 14 |
| 23 | The progress of gut microbiome research related to brain disorders. Journal of Neuroinflammation, 2020, 17, 25. | 7.2 | 252 |
| 24 | Transcriptome analysis identifies key regulators and networks in Acute myeloid leukemia. Hematology, 2019, 24, 487-491. | 1.5 | 7 |
| 25 | Epithelial-type systemic breast carcinoma cells with a restricted mesenchymal transition are a major source of metastasis. Science Advances, 2019, 5, eaav4275. | 10.3 | 139 |
| 26 | AMPK alleviates high uric acid-induced Na+-K+-ATPase signaling impairment and cell injury in renal tubules. Experimental and Molecular Medicine, 2019, 51, 1-14. | 7.7 | 24 |
| 27 | Comparison and development of machine learning tools in the prediction of chronic kidney disease progression. Journal of Translational Medicine, 2019, 17, 119. | 4.4 | 134 |
| 28 | Specific zinc fingerâ€induced methylation of PDâ€L1 promoter inhibits its expression. FEBS Open Bio, 2019, 9, 1063-1070. | 2.3 | 17 |
| 29 | Checkpoint inhibitors for stage I to III non-small cell lung cancer treated with surgery or radiotherapy with curative intent: a generic protocol. The Cochrane Library, 2019, , . | 2.8 | 2 |
| 30 | Transcriptome Sequencing to Detect the Potential Role of Long Noncoding RNAs in Salt-Sensitive Hypertensive Rats. BioMed Research International, 2019, 2019, 1-9. | 1.9 | 7 |
| 31 | RNA-seq analyses the effect of high-salt diet in hypertension. Gene, 2018, 677, 245-250. | 2.2 | 22 |
| 32 | A Comprehensive Analysis of Gene Expression of Xenobiotic and Endogenous Metabolizing Enzymes and Transporters in Rat Multiple Organs. Current Pharmaceutical Biotechnology, 2018, 19, 240-249. | 1.6 | 0 |
| 33 | Advances in single-cell RNA sequencing and its applications in cancer research. Oncotarget, 2017, 8, 53763-53779. | 1.8 | 76 |
| 34 | Expression profiling and functional annotation of noncoding genes across 11 distinct organs in rat development. Scientific Reports, 2016, 6, 38575. | 3.3 | 4 |
| 35 | Identification of Tissue-Specific Protein-Coding and Noncoding Transcripts across 14 Human Tissues Using RNA-seq. Scientific Reports, 2016, 6, 28400. | 3.3 | 57 |
| 36 | A Newly Identified Natural Splice Variant ASN Enhances Hepatitis B Virus Amplification. Viral Immunology, 2016, 29, 27-32. | 1.3 | 0 |