# Jiaqi Liu

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7073388/publications.pdf

Version: 2024-02-01

| 49 | 904 | 471509 17 | 526287 27 |
|---|---|---|---|
| papers | citations | h-index | g-index |

| 52 | 52 | 52 | 1391 |
|---|---|---|---|
| all docs | docs citations | times ranked | citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | Longâ€term outcomes of intraoperative radiotherapy forÂearlyâ€stage breast cancer in China: a multicenter realâ€worldÂstudy. Cancer Communications, 2022, 42, 277-280. | 9.2 | 2 |
| 2 | DrABC: deep learning accurately predicts germline pathogenic mutation status in breast cancer patients based on phenotype data. Genome Medicine, 2022, 14, 21. | 8.2 | 4 |
| 3 | Hsa_circ_0069094 accelerates cell malignancy and glycolysis through regulating the miR-591/HK2 axis in breast cancer. Cellular Signalling, 2021, 79, 109878. | 3.6 | 13 |
| 4 | Prevalence and reclassification of BRCA1 and BRCA2 variants in a large, unselected Chinese Han breast cancer cohort. Journal of Hematology and Oncology, 2021, 14, 18. | 17.0 | 12 |
| 5 | Genomic instability-derived plasma extracellular vesicle-microRNA signature as a minimally invasive predictor of risk and unfavorable prognosis in breast cancer. Journal of Nanobiotechnology, 2021, 19, 22. | 9.1 | 52 |
| 6 | Abstract PS1-31: Nomogram for predicting axillary lymph node pathological response in node-positive breast cancer patients after neoadjuvant chemotherapy. , 2021, , . | | 0 |
| 7 | Genome-wide cell-free DNA methylation analyses improve accuracy of non-invasive diagnostic imaging for early-stage breast cancer. Molecular Cancer, 2021, 20, 36. | 19.2 | 30 |
| 8 | Silencing of Nek2 suppresses the proliferation, migration and invasion and induces apoptosis of breast cancer cells by regulating ERK/MAPK signaling. Journal of Molecular Histology, 2021, 52, 809-821. | 2.2 | 5 |
| 9 | Circulating exosomal miRâ€363â€5p inhibits lymph node metastasis by downregulating PDGFB and serves as a potential noninvasive biomarker for breast cancer. Molecular Oncology, 2021, 15, 2466-2479. | 4.6 | 31 |
| 10 | Effects of Ulinastatin on Proliferation and Apoptosis of Breast Cancer Cells by Inhibiting the ERK Signaling Pathway. BioMed Research International, 2021, 2021, 1-6. | 1.9 | 7 |
| 11 | The Distinct Performances of Ultrasound, Mammograms, and MRI in Detecting Breast Cancer in Patients With Germline Pathogenic Variants in Cancer Predisposition Genes. Frontiers in Oncology, 2021, 11, 710156. | 2.8 | 2 |
| 12 | CircRNA circ-PDCD11 promotes triple-negative breast cancer progression via enhancing aerobic glycolysis. Cell Death Discovery, 2021, 7, 218. | 4.7 | 24 |
| 13 | Effect of MiR-210 on the Chemosensitivity of Breast Cancer by Regulating JAK-STAT Signaling Pathway. BioMed Research International, 2021, 2021, 1-8. | 1.9 | 12 |
| 14 | Contralateral risk-reducing local therapy in breast cancer patients with BRCA1/2 mutations: systemic review and meta-analysis. Cancer Cell International, 2021, 21, 512. | 4.1 | 4 |
| 15 | LINC00337 induces tumor development and chemoresistance to paclitaxel of breast cancer by recruiting M2 tumor-associated macrophages. Molecular Immunology, 2021, 138, 1-9. | 2.2 | 20 |
| 16 | Anti-vascular endothelial growth factor therapy in breast cancer: Molecular pathway, potential targets, and current treatment strategies. Cancer Letters, 2021, 520, 422-433. | 7.2 | 15 |
| 17 | 18:0 Lyso PC, a natural product with potential PPAR-Î³ agonistic activity, plays hypoglycemic effect with lower liver toxicity and cardiotoxicity in db/db mice. Biochemical and Biophysical Research Communications, 2021, 579, 168-174. | 2.1 | 8 |
| 18 | Overexpression of lncRNA SAMMSON Promotes Triple-Negative Breast Cancer Cell Proliferation by Interacting with p53. Critical Reviews in Eukaryotic Gene Expression, 2021, 31, 1-8. | 0.9 | 3 |

| # | Article | IF | Citations |
|---|---|---|---|
| 19 | Cis -acting lnc-Cxcl2 restrains neutrophil-mediated lung inflammation by inhibiting epithelial cell CXCL2 expression in virus infection. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, . | 7.1 | 24 |
| 20 | Expression and prognostic value of , , and gene clusters in human breast cancer. Journal of International Medical Research, 2021, 49, 300060520980647. | 1.0 | 4 |
| 21 | Cost-Effectiveness Analysis of Imaging Modalities for Breast Cancer Surveillance Among BRCA1/2 Mutation Carriers: A Systematic Review. Frontiers in Oncology, 2021, 11, 763161. | 2.8 | 2 |
| 22 | CFEA: a cell-free epigenome atlas in human diseases. Nucleic Acids Research, 2020, 48, D40-D44. | 14.5 | 32 |
| 23 | TBX6 missense variants expand the mutational spectrum in a nonâ€Mendelian inheritance disease. Human Mutation, 2020, 41, 182-195. | 2.5 | 27 |
| 24 | Front Cover, Volume 41, Issue 1. Human Mutation, 2020, 41, i. | 2.5 | 0 |
| 25 | Decreased level of peripheral CD8<sup>+</sup>CD28<sup>+</sup> T cells is associated with lymph node metastasis in patients with breast cancer. Future Oncology, 2020, 16, 2611-2617. | 2.4 | 2 |
| 26 | Cost-effectiveness analysis of using the TBX6-associated congenital scoliosis risk score (TACScore) in genetic diagnosis of congenital scoliosis. Orphanet Journal of Rare Diseases, 2020, 15, 250. | 2.7 | 2 |
| 27 | Clinical Application and Feasibility of MRI-Guided Breast Biopsy of Breast Minimal Lesions in Chinese Population. Frontiers in Oncology, 2020, 10, 257. | 2.8 | 2 |
| 28 | Clinicopathological Relevance and Prognostic Value of Androgen Receptor in Mammary Pagetâ€™s Disease with Underlying Invasive Ductal Carcinoma. Oncology Research and Treatment, 2020, 43, 346-353. | 1.2 | 2 |
| 29 | Mutational landscape and genetic signatures of cellâ€free DNA in tumourâ€induced osteomalacia. Journal of Cellular and Molecular Medicine, 2020, 24, 4931-4943. | 3.6 | 4 |
| 30 | Human and mouse studies establish TBX6 in Mendelian CAKUT and as a potential driver of kidney defects associated with the 16p11.2 microdeletion syndrome. Kidney International, 2020, 98, 1020-1030. | 5.2 | 17 |
| 31 | Large-scale integrated analysis of ovarian cancer tumors and cell lines identifies an individualized gene expression signature for predicting response to platinum-based chemotherapy. Cell Death and Disease, 2019, 10, 661. | 6.3 | 18 |
| 32 | A20 as a novel target for the anti-neuroinflammatory effect of chrysin via inhibition of NF-ÎºB signaling pathway. Brain, Behavior, and Immunity, 2019, 79, 228-235. | 4.1 | 16 |
| 33 | Genetic polymorphisms of PAX1 are functionally associated with different PUMC types of adolescent idiopathic scoliosis in a northern Chinese Han population. Gene, 2019, 688, 215-220. | 2.2 | 19 |
| 34 | Impact of Locoregional Treatment on Prognosis of de novo Stage IV Breast Cancer: A Retrospective Long-Term Study of Chinese Population. Gynecologic and Obstetric Investigation, 2019, 84, 248-258. | 1.6 | 4 |
| 35 | TBX6-associated congenital scoliosis (TACS) as a clinically distinguishable subtype of congenital scoliosis: further evidence supporting the compound inheritance and TBX6 gene dosage model. Genetics in Medicine, 2019, 21, 1548-1558. | 2.4 | 60 |
| 36 | TBX6 compound inheritance leads to congenital vertebral malformations in humans and mice. Human Molecular Genetics, 2019, 28, 539-547. | 2.9 | 46 |

| # | Article | IF | Citations |
|---|---|---|---|
| 37 | Phenotypic heterogeneity of intellectual disability in patients with congenital insensitivity to pain with anhidrosis: A case report and literature review. Journal of International Medical Research, 2018, 46, 2445-2457. | 1.0 | 11 |
| 38 | Genetic polymorphisms of <scp>GPR</scp>126 are functionally associated with <scp>PUMC</scp> classifications of adolescent idiopathic scoliosis in a Northern Han population. Journal of Cellular and Molecular Medicine, 2018, 22, 1964-1971. | 3.6 | 31 |
| 39 | Comparative analysis of serum proteome in congenital scoliosis patients with <scp>TBX</scp>6 haploinsufficiency â€“ a first report pointing to lipid metabolism. Journal of Cellular and Molecular Medicine, 2018, 22, 533-545. | 3.6 | 16 |
| 40 | The coexistence of copy number variations (CNVs) and single nucleotide polymorphisms (SNPs) at a locus can result in distorted calculations of the significance in associating SNPs to disease. Human Genetics, 2018, 137, 553-567. | 3.8 | 57 |
| 41 | Progress and Application of CRISPR/Cas Technology in Biological and Biomedical Investigation. Journal of Cellular Biochemistry, 2017, 118, 3061-3071. | 2.6 | 10 |
| 42 | Genetic Polymorphism of LBX1 Is Associated With Adolescent Idiopathic Scoliosis in Northern Chinese Han Population. Spine, 2017, 42, 1125-1129. | 2.0 | 45 |
| 43 | Molecular therapeutic strategies for FGFR3 gene-related skeletal dysplasia. Journal of Molecular Medicine, 2017, 95, 1303-1313. | 3.9 | 6 |
| 44 | CRISPR/Cas9 in zebrafish: an efficient combination for human genetic diseases modeling. Human Genetics, 2017, 136, 1-12. | 3.8 | 83 |
| 45 | Breast cancer in young women of Chinese Han population: A retrospective study of patients under 25 years. Pathology Research and Practice, 2016, 212, 1015-1020. | 2.3 | 5 |
| 46 | The Association Between Body Size and Breast Cancer in Han Women in Northern and Eastern China. Oncologist, 2016, 21, 1362-1368. | 3.7 | 22 |
| 47 | Association betweenADAMTS-4gene polymorphism and lumbar disc degeneration in Chinese Han population. Journal of Orthopaedic Research, 2016, 34, 860-864. | 2.3 | 26 |
| 48 | FusionCancer: a database of cancer fusion genes derived from RNA-seq data. Diagnostic Pathology, 2015, 10, 131. | 2.0 | 61 |
| 49 | Novel NTRK1 Frameshift Mutation in Congenital Insensitivity to Pain With Anhidrosis. Journal of Child Neurology, 2015, 30, 1357-1361. | 1.4 | 6 |