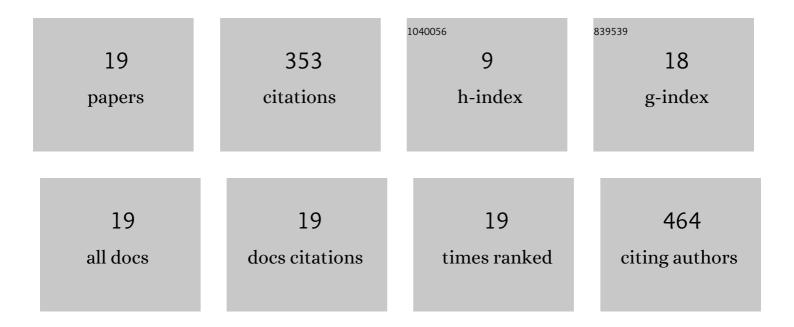
## Xujun Liang

List of Publications by Year in descending order

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| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | LRSSL: predict and interpret drug–disease associations based on data integration using sparse subspace learning. Bioinformatics, 2017, 33, 1187-1196.  | 4.1  | 106       |
| 2  | Structural basis for DNA recognition by FOXC2. Nucleic Acids Research, 2019, 47, 3752-3764.  | 14.5 | 36        |
| 3  | Identifying DCN and HSPD1 as Potential Biomarkers in Colon Cancer Using 2D-LC-MS/MS Combined with iTRAQ Technology. Journal of Cancer, 2017, 8, 479-489.   | 2.5  | 35        |
| 4  | Mechanism of forkhead transcription factors binding to a novel palindromic DNA site. Nucleic Acids<br>Research, 2021, 49, 3573-3583.   | 14.5 | 28        |
| 5  | Structural basis of binding of homodimers of the nuclear receptor NR4A2 to selective Nur-responsive DNA elements. Journal of Biological Chemistry, 2019, 294, 19795-19803.   | 3.4  | 23        |
| 6  | lncRNA RP11-624L4.1 Is Associated with Unfavorable Prognosis and Promotes Proliferation via the<br>CDK4/6-Cyclin D1-Rb-E2F1 Pathway in NPC. Molecular Therapy - Nucleic Acids, 2020, 22, 1025-1039.  | 5.1  | 20        |
| 7  | The Significance of Serum S100A9 and TNC Levels as Biomarkers in Colorectal Cancer. Journal of Cancer, 2019, 10, 5315-5323.  | 2.5  | 19        |
| 8  | Quantitative proteomic profiling of tumor-associated vascular endothelial cells in colorectal cancer. Biology Open, 2019, 8, .   | 1.2  | 14        |
| 9  | DNA-binding properties of FOXP3 transcription factor. Acta Biochimica Et Biophysica Sinica, 2017, 49, 792-799.   | 2.0  | 12        |
| 10 | Identification of VWF as a Novel Biomarker in Lung Adenocarcinoma by Comprehensive Analysis.<br>Frontiers in Oncology, 2021, 11, 639600.   | 2.8  | 10        |
| 11 | S-Adenosylmethionine Affects Cell Cycle Pathways and Suppresses Proliferation in Liver Cells. Journal of Cancer, 2019, 10, 4368-4379.  | 2.5  | 9         |
| 12 | Simultaneous Inhibition of EGFR and HER2 via Afatinib Augments the Radiosensitivity of<br>Nasopharyngeal Carcinoma Cells. Journal of Cancer, 2019, 10, 2063-2073.  | 2.5  | 8         |
| 13 | Learning important features from multi-view data to predict drug side effects. Journal of<br>Cheminformatics, 2019, 11, 79.  | 6.1  | 8         |
| 14 | Identification of prognostic genes in the pancreatic adenocarcinoma immune microenvironment by integrated bioinformatics analysis. Cancer Immunology, Immunotherapy, 2022, 71, 1757-1769.  | 4.2  | 8         |
| 15 | Coupling of Cell Surface Biotinylation and SILAC-Based Quantitative Proteomics Identified Myoferlin<br>as a Potential Therapeutic Target for Nasopharyngeal Carcinoma Metastasis. Frontiers in Cell and<br>Developmental Biology, 2021, 9, 621810. | 3.7  | 6         |
| 16 | Identification of a pyroptosis-related IncRNA risk model for predicting prognosis and immune response in colon adenocarcinoma. World Journal of Surgical Oncology, 2022, 20, 118.  | 1.9  | 4         |
| 17 | Mechanistic Insights into the Preference for Tandem Binding Sites in DNA Recognition by FOXM1.<br>Journal of Molecular Biology, 2022, 434, 167426.   | 4.2  | 3         |
| 18 | A novel machine learning model based on sparse structure learning with adaptive graph<br>regularization for predicting drug side effects. Journal of Biomedical Informatics, 2022, 132, 104131.  | 4.3  | 3         |

| #  | Article  | IF  | CITATIONS |
|----|--|-----|-----------|
| 19 | Detection and Analysis of RNAs Expression Profile for Methylated Candidate Tumor Suppressor Genes in Nasopharyngeal Carcinoma. Anti-Cancer Agents in Medicinal Chemistry, 2019, 19, 772-782. | 1.7 | 1         |