## Qing Zhong

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4952440/publications.pdf

Version: 2024-02-01

39 papers 1,423 citations

18 h-index 32 g-index

46 all docs

46 docs citations

times ranked

46

2864 citing authors

| #  | Article  | IF          | CITATIONS |
|----|--|-------------|-----------|
| 1  | Prostate cancer–associated SPOP mutations confer resistance to BET inhibitors through stabilization of BRD4. Nature Medicine, 2017, 23, 1063-1071.   | 15.2        | 240       |
| 2  | TRIM24 Is an Oncogenic Transcriptional Activator in Prostate Cancer. Cancer Cell, 2016, 29, 846-858.   | 7.7         | 228       |
| 3  | Highâ€throughput proteomic analysis of <scp>FFPE</scp> tissue samples facilitates tumor stratification. Molecular Oncology, 2019, 13, 2305-2328.   | 2.1         | 100       |
| 4  | Unsupervised modeling of cell morphology dynamics for time-lapse microscopy. Nature Methods, 2012, 9, 711-713.   | 9.0         | 81        |
| 5  | Machine learning for multi-omics data integration in cancer. IScience, 2022, 25, 103798.   | 1.9         | 78        |
| 6  | Strategies to enable large-scale proteomics for reproducible research. Nature Communications, 2020, 11, 3793.  | 5.8         | 75        |
| 7  | Quantitative Proteome Landscape of the NCI-60 Cancer Cell Lines. IScience, 2019, 21, 664-680.  | 1.9         | 52        |
| 8  | Pan-cancer proteomic map of 949 human cell lines. Cancer Cell, 2022, 40, 835-849.e8.   | 7.7         | 52        |
| 9  | Multi-region proteome analysis quantifies spatial heterogeneity of prostate tissue biomarkers. Life Science Alliance, 2018, 1, e201800042.   | 1.3         | 51        |
| 10 | Cytology smears as excellent starting material for nextâ€generation sequencingâ€based molecular testing of patients with adenocarcinoma of the lung. Cancer Cytopathology, 2017, 125, 30-40. | 1.4         | 47        |
| 11 | MiR-99b-5p expression and response to tyrosine kinase inhibitor treatment in clear cell renal cell carcinoma patients. Oncotarget, 2016, 7, 78433-78447.                                     | 0.8         | 45        |
| 12 | Comparison of the Proliferation and Differentiation Potential of Human Urine-, Placenta Decidua Basalis-, and Bone Marrow-Derived Stem Cells. Stem Cells International, 2018, 2018, 1-11.    | 1.2         | 41        |
| 13 | Comparative analysis of mRNA and protein degradation in prostate tissues indicates high stability of proteins. Nature Communications, 2019, 10, 2524.  | <b>5.</b> 8 | 35        |
| 14 | Connexin 43 expression predicts poor progression-free survival in patients with non-muscle invasive urothelial bladder cancer. Journal of Clinical Pathology, 2015, 68, 819-824.             | 1.0         | 34        |
| 15 | Image-based computational quantification and visualization of genetic alterations and tumour heterogeneity. Scientific Reports, 2016, 6, 24146.  | 1.6         | 28        |
| 16 | Addressing the Challenges of Highâ€Throughput Cancer Tissue Proteomics for Clinical Application: ProCan. Proteomics, 2019, 19, e1900109.   | 1.3         | 25        |
| 17 | <scp>KPNA2</scp> is overexpressed in human and mouse endometrial cancers and promotes cellular proliferation. Journal of Pathology, 2014, 234, 239-252.                                      | 2.1         | 23        |
| 18 | Application of Nanosecond Laser Photolysis Protein Footprinting to Study EGFR Activation by EGF in Cells. Journal of Proteome Research, 2017, 16, 2282-2293.                                 | 1.8         | 21        |

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|----|--|-----|-----------|
| 19 | A curated collection of tissue microarray images and clinical outcome data of prostate cancer patients. Scientific Data, 2017, 4, 170014.  | 2.4 | 21        |
| 20 | Value of postmortem studies in deceased neonatal and pediatric intensive care unit patients. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2017, 470, 217-223.                                   | 1.4 | 18        |
| 21 | Convergent network effects along the axis of gene expression during prostate cancer progression. Genome Biology, 2020, 21, 302.  | 3.8 | 17        |
| 22 | Detection of <i>CCNE1/URI</i> (19q12) amplification by <i>in situ</i> hybridisation is common in high grade and type II endometrial cancer. Oncotarget, 2017, 8, 14794-14805.  | 0.8 | 16        |
| 23 | Positive fibroblast growth factor receptor 3 immunoreactivity is associated with low-grade non-invasive urothelial bladder cancer. Oncology Letters, 2015, 10, 2753-2760.  | 0.8 | 13        |
| 24 | Targeted next-generation-sequencing for reliable detection of targetable rearrangements in lung adenocarcinoma—a single center retrospective study. Pathology Research and Practice, 2018, 214, 572-578.                                 | 1.0 | 13        |
| 25 | Multi-laboratory proficiency testing of clinical cancer genomic profiling by next-generation sequencing. Pathology Research and Practice, 2018, 214, 957-963.  | 1.0 | 11        |
| 26 | Oxygen supply maps for hypoxic microenvironment visualization in prostate cancer. Journal of Pathology Informatics, 2016, 7, 3.  | 0.8 | 10        |
| 27 | A novel germline mutation of PDGFR- $\hat{l}^2$ might be associated with clinical response of colorectal cancer to regorafenib. Annals of Oncology, 2015, 26, 246-248.   | 0.6 | 8         |
| 28 | A computational framework for disease grading using protein signatures. , 2016, , .  |     | 7         |
| 29 | Modelling of a genetically diverse evolution of Systemic Mastocytosis with Chronic Myelomonocytic Leukemia (SM-CMML) by Next Generation Sequencing. Experimental Hematology and Oncology, 2014, 3, 18.                                   | 2.0 | 5         |
| 30 | Prevalence of hypertension and diabetes after exposure to extracorporeal shock-wave lithotripsy in patients with renal calculi: a retrospective non-randomized data analysis. International Urology and Nephrology, 2018, 50, 1227-1233. | 0.6 | 4         |
| 31 | PIONEER: Pipeline for Generating Highâ€Quality Spectral Libraries for DIAâ€MS Data. Current Protocols, 2021, 1, e69.   | 1.3 | 4         |
| 32 | Improved identification and quantification of peptides in mass spectrometry data via chemical and random additive noise elimination (CRANE). Bioinformatics, 2021, 37, 4719-4726.  | 1.8 | 4         |
| 33 | Heterogeneity characterization of immunohistochemistry stained tissue using convolutional autoencoder., 2017,,.  |     | 3         |
| 34 | Deciphering protein signatures using color, morphological, and topological analysis of immunohistochemically stained human tissues. Proceedings of SPIE, $2016, \ldots$  | 0.8 | 2         |
| 35 | Disease grading of heterogeneous tissue using convolutional autoencoder. , 2017, , .   |     | 1         |
| 36 | Abstract 5565: Multi-omic profiling of prostate cancer evolution in 39 patients., 2017,,.  |     | 1         |

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|----|--|-----|-----------|
| 37 | CD10 expression in 325 testicular germ cell tumours. Journal of Clinical Pathology, 2015, 68, 400-403.   | 1.0 | O         |
| 38 | MP62-11 EXTRACORPOREAL SHOCK-WAVE LITHOTRIPSY (ESWL) FOR RENAL STONES IS ASSOCIATED WITH DECREASED KIDNEY FUNCTION AFTER LONG TERM FOLLOW-UP. Journal of Urology, 2017, 197, . | 0.2 | 0         |
| 39 | Computational Pathology. , 2017, , 263-279.  |     | 0         |