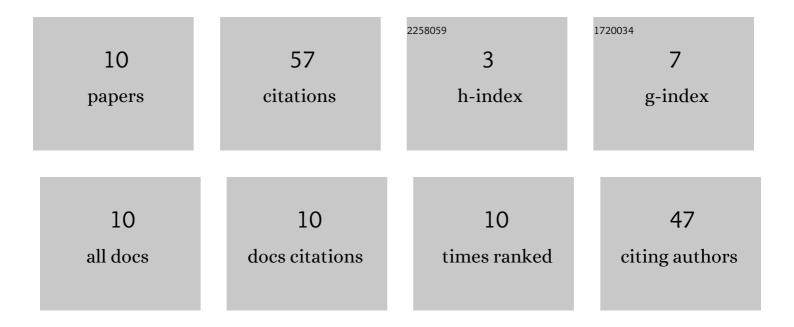
## Xu Zhang

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7709204/publications.pdf Version: 2024-02-01



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#	Article	IF	CITATIONS
1	A pan-cancer analysis based on weighted gene co-expression network analysis identifies the biomarker utility of lamin B1 in human tumors. Cancer Biomarkers, 2022, 34, 23-39.	1.7	3
2	AMDBNorm: an approach based on distribution adjustment to eliminate batch effects of gene expression data. Briefings in Bioinformatics, 2022, 23, .	6.5	1
3	Identifying of biomarkers associated with gastric cancer based on 11 topological analysis methods of CytoHubba. Scientific Reports, 2021, 11, 1331.	3.3	36
4	Construction and Comprehensive Analysis of a Special Competitive Endogenous RNAs Network to Reveal Potential Prognostic Biomarkers for Endometrial Carcinoma. Current Bioinformatics, 2021, 16, 1341-1354.	1.5	1
5	Clustering of genes from microarray data using hierarchical projective adaptive resonance theory: a case study of tuberculosis. Briefings in Functional Genomics, 2021, , .	2.7	0
6	Construction of competitive endogenous RNA network related to circular RNA and prognostic nomogram model in lung adenocarcinoma. Mathematical Biosciences and Engineering, 2021, 18, 9806-9821.	1.9	0
7	Analysis of Cyclin-Dependent Kinase 1 as an Independent Prognostic Factor for Gastric Cancer Based on Statistical Methods. Frontiers in Cell and Developmental Biology, 2020, 8, 620164.	3.7	9
8	Identifying candidate diagnostic markers for early stage of non-small cell lung cancer. PLoS ONE, 2019, 14, e0225080.	2.5	1
9	Identifying candidate diagnostic markers for tuberculosis: A critical role of co-expression and pathway analysis. Mathematical Biosciences and Engineering, 2019, 16, 541-552.	1.9	3
10	Influences of migrations from local competitive pressures on populations between patches. Journal of Applied Mathematics and Computing, 2011, 37, 313-330.	2.5	3