## Tianyi Xu

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9191036/publications.pdf

Version: 2024-02-01

15	860	840119 <b>11</b>	940134	
papers	citations	h-index	g-index	
16	16	16	725	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	Database Resources of the National Genomics Data Center, China National Center for Bioinformation in 2022. Nucleic Acids Research, 2022, 50, D27-D38.	6.5	364
2	Circular RNA expression profiles and features in human tissues: a study using RNA-seq data. BMC Genomics, 2017, 18, 680.	1,2	193
3	IRESfinder: Identifying RNA internal ribosome entry site in eukaryotic cell using framed k-mer features. Journal of Genetics and Genomics, 2018, 45, 403-406.	1.7	77
4	CircAST: Full-length Assembly and Quantification of Alternatively Spliced Isoforms in Circular RNAs. Genomics, Proteomics and Bioinformatics, 2019, 17, 522-534.	3.0	40
5	VISDB: a manually curated database of viral integration sites in the human genome. Nucleic Acids Research, 2020, 48, D633-D641.	6.5	39
6	Circular RNA expression profiles during the differentiation of mouse neural stem cells. BMC Systems Biology, 2018, 12, 128.	3.0	31
7	CeDR Atlas: a knowledgebase of cellular drug response. Nucleic Acids Research, 2022, 50, D1164-D1171.	6.5	22
8	Gene Expression Nebulas (GEN): a comprehensive data portal integrating transcriptomic profiles across multiple species at both bulk and single-cell levels. Nucleic Acids Research, 2022, 50, D1016-D1024.	6.5	18
9	Genome-Wide Analysis of the Expression of Circular RNA Full-Length Transcripts and Construction of the circRNA-miRNA-mRNA Network in Cervical Cancer. Frontiers in Cell and Developmental Biology, 2020, 8, 603516.	1.8	16
10	Identification of Potential Prognostic Competing Triplets in High-Grade Serous Ovarian Cancer. Frontiers in Genetics, 2020, 11, 607722.	1.1	14
11	Changes in the Microbial Community Diversity of Oil Exploitation. Genes, 2019, 10, 556.	1.0	13
12	The Expression Profiles of IncRNAs and Their Regulatory Network During Smek1/2 Knockout Mouse Neural Stem Cells Differentiation. Current Bioinformatics, 2020, 15, 77-88.	0.7	11
13	BrainBase: a curated knowledgebase for brain diseases. Nucleic Acids Research, 2022, 50, D1131-D1138.	6.5	11
14	An Integrative Transcriptomic and Methylation Approach for Identifying Differentially Expressed Circular RNAs Associated with DNA Methylation Change. Biomedicines, 2021, 9, 657.	1.4	6
15	Identification and Analysis of RNA Editing Events in Ovarian Serous Cystadenoma Using RNA-seq Data. Current Gene Therapy, 2021, 21, 258-269.	0.9	4