## Yue Zhao

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6589573/publications.pdf

Version: 2024-02-01

|          | 2682572        |              | 2272923        |  |
|----------|----------------|--------------|----------------|--|
| 7        | 31             | 2            | 4              |  |
| papers   | citations      | h-index      | g-index        |  |
|          |                |              |                |  |
| 7        | 7              | 7            | 29             |  |
| all docs | docs citations | times ranked | citing authors |  |
|          |                |              |                |  |

| # | Article  | IF  | CITATIONS |
|---|--|-----|-----------|
| 1 | Deep Pathway Analysis V2.0: A Pathway Analysis Framework Incorporating Multi-Dimensional Omics Data. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2021, 18, 373-385. | 3.0 | 1         |
| 2 | A framework using topological pathways for deeper analysis of transcriptome data. BMC Genomics, 2020, 21, 834.   | 2.8 | 0         |
| 3 | BioTarget: A Computational Framework Identifying Cancer Type Specific Transcriptional Targets of Immune Response Pathways. Scientific Reports, 2019, 9, 9029.                              | 3.3 | 7         |
| 4 | Recurrent Neural Network for Gene Regulation Network Construction on Time Series Expression Data. , 2019, , .  |     | 4         |
| 5 | An Extension of Deep Pathway Analysis: A Pathway Route Analysis Framework Incorporating<br>Multi-dimensional Cancer Genomics Data. Lecture Notes in Computer Science, 2018, , 113-124.     | 1.3 | 1         |
| 6 | A route-based pathway analysis framework integrating mutation information and gene expression data. Methods, 2017, 124, 3-12.  | 3.8 | 16        |
| 7 | Deep pathway analysis incorporating mutation information and gene expression data. , $2016, \ldots$  |     | 2         |