

# Diem-Trang T Tran

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4415993/publications.pdf>

Version: 2024-02-01

12  
papers

37  
citations

2258059

3  
h-index

1872680

6  
g-index

13  
all docs

13  
docs citations

13  
times ranked

81  
citing authors

| #  | ARTICLE  | IF  | CITATIONS |
|----|--|-----|-----------|
| 1  | cdev: a ground-truth based measure to evaluate RNA-seq normalization performance. PeerJ, 2021, 9, e12233.  | 2.0 | 2         |
| 2  | Cardioinformatics: the nexus of bioinformatics and precision cardiology. Briefings in Bioinformatics, 2020, 21, 2031-2051.   | 6.5 | 15        |
| 3  | A graph-based algorithm for RNA-seq data normalization. PLoS ONE, 2020, 15, e0227760.  | 2.5 | 3         |
| 4  | A graph-based algorithm for RNA-seq data normalization. , 2020, 15, e0227760.  |     | 0         |
| 5  | A graph-based algorithm for RNA-seq data normalization. , 2020, 15, e0227760.  |     | 0         |
| 6  | A graph-based algorithm for RNA-seq data normalization. , 2020, 15, e0227760.  |     | 0         |
| 7  | A graph-based algorithm for RNA-seq data normalization. , 2020, 15, e0227760.  |     | 0         |
| 8  | A graph-based algorithm for RNA-seq data normalization. , 2020, 15, e0227760.  |     | 0         |
| 9  | A graph-based algorithm for RNA-seq data normalization. , 2020, 15, e0227760.  |     | 0         |
| 10 | HeartBioPortal. Circulation Genomic and Precision Medicine, 2019, 12, e002426.   | 3.6 | 8         |
| 11 | anexVis: visual analytics framework for analysis of RNA expression. Bioinformatics, 2018, 34, 2510-2512.   | 4.1 | 1         |
| 12 | Discover binding pathways using the sliding binding-box docking approach: application to binding pathways of oseltamivir to avian influenza H5N1 neuraminidase. Journal of Computer-Aided Molecular Design, 2013, 27, 689-695. | 2.9 | 8         |