## **Dustin Ebert**

## List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/5582527/dustin-ebert-publications-by-year.pdf

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

| 10                | 1,989                | 5                   | 11              |
|-------------------|----------------------|---------------------|-----------------|
| papers            | citations            | h-index             | g-index         |
| 11<br>ext. papers | 3,728 ext. citations | <b>11.6</b> avg, IF | 5.87<br>L-index |

| #  | Paper  | IF   | Citations |
|----|--|------|-----------|
| 10 | PANTHER: Making genome-scale phylogenetics accessible to all. <i>Protein Science</i> , <b>2021</b> ,   | 6.3  | 12        |
| 9  | PANTHER version 16: a revised family classification, tree-based classification tool, enhancer regions and extensive API. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, D394-D403 | 20.1 | 248       |
| 8  | PEREGRINE: A genome-wide prediction of enhancer to gene relationships supported by experimental evidence. <i>PLoS ONE</i> , <b>2020</b> , 15, e0243791                               | 3.7  | 3         |
| 7  | PhyloGenes: An online phylogenetics and functional genomics resource for plant gene function inference. <i>Plant Direct</i> , <b>2020</b> , 4, e00293                                | 3.3  | 8         |
| 6  | PEREGRINE: A genome-wide prediction of enhancer to gene relationships supported by experimental evidence <b>2020</b> , 15, e0243791  |      |           |
| 5  | PEREGRINE: A genome-wide prediction of enhancer to gene relationships supported by experimental evidence <b>2020</b> , 15, e0243791  |      |           |
| 4  | PEREGRINE: A genome-wide prediction of enhancer to gene relationships supported by experimental evidence <b>2020</b> , 15, e0243791  |      |           |
| 3  | PEREGRINE: A genome-wide prediction of enhancer to gene relationships supported by experimental evidence <b>2020</b> , 15, e0243791  |      |           |
| 2  | Protocol Update for large-scale genome and gene function analysis with the PANTHER classification system (v.14.0). <i>Nature Protocols</i> , <b>2019</b> , 14, 703-721               | 18.8 | 462       |
| 1  | PANTHER version 14: more genomes, a new PANTHER GO-slim and improvements in enrichment analysis tools. <i>Nucleic Acids Research</i> , <b>2019</b> , 47, D419-D426                   | 20.1 | 1256      |