

# Kokulapalan Wimalanathan

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1346760/publications.pdf>

Version: 2024-02-01

11  
papers

582  
citations

1307594

7  
h-index

1588992

8  
g-index

14  
all docs

14  
docs citations

14  
times ranked

1321  
citing authors

| #  | ARTICLE  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Standardized genome-wide function prediction enables comparative functional genomics: a new application area for Gene Ontologies in plants. <i>GigaScience</i> , 2022, 11, . | 6.4  | 2         |
| 2  | Gene Ontology Meta Annotator for Plants (GOMAP). <i>Plant Methods</i> , 2021, 17, 54.  | 4.3  | 21        |
| 3  | A high-resolution gene expression atlas links dedicated meristem genes to key architectural traits. <i>Genome Research</i> , 2019, 29, 1962-1973.                            | 5.5  | 35        |
| 4  | Compare Expression Profiles for Pre-defined Gene Groups with C-REx. <i>Journal of Open Source Software</i> , 2019, 4, 1255.  | 4.6  | 0         |
| 5  | Cover Image: Volume 20, Issue 5. <i>Evolution &amp; Development</i> , 2018, 20, i.   | 2.0  | 0         |
| 6  | Maize GO Annotationâ€™Methods, Evaluation, and Review (maizeâ€™GAMER). <i>Plant Direct</i> , 2018, 2, e00052.  | 1.9  | 97        |
| 7  | The maize W22 genome provides a foundation for functional genomics and transposon biology. <i>Nature Genetics</i> , 2018, 50, 1282-1288.                                     | 21.4 | 183       |
| 8  | Gene network variation and alternative paths to convergent evolution in turtles. <i>Evolution &amp; Development</i> , 2018, 20, 172-185.                                     | 2.0  | 7         |
| 9  | MaizeGDB update: new tools, data and interface for the maize model organism database. <i>Nucleic Acids Research</i> , 2016, 44, D1195-D1201.                                 | 14.5 | 170       |
| 10 | YouGenMap: a web platform for dynamic multi-comparative mapping and visualization of genetic maps. <i>Frontiers in Genetics</i> , 2014, 5, 183.                              | 2.3  | 0         |
| 11 | An annotated genetic map of loblolly pine based on microsatellite and cDNA markers. <i>BMC Genetics</i> , 2011, 12, 17.  | 2.7  | 60        |