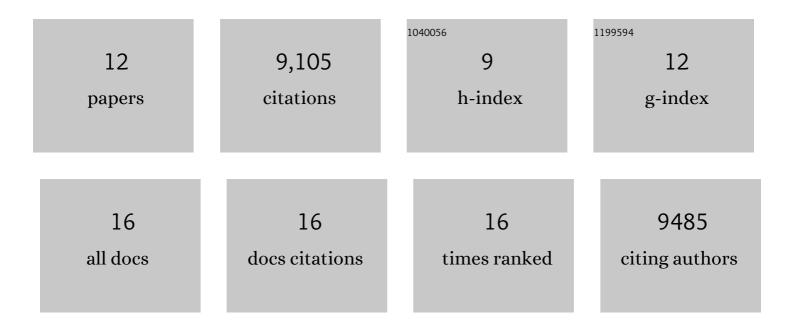
Terry M Casstevens

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1245633/publications.pdf Version: 2024-02-01



| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | TASSEL: software for association mapping of complex traits in diverse samples. Bioinformatics, 2007, 23, 2633-2635. | 4.1 | 6,407 |
| 2 | TASSEL-GBS: A High Capacity Genotyping by Sequencing Analysis Pipeline. PLoS ONE, 2014, 9, e90346. | 2.5 | 1,511 |
| 3 | Comprehensive genotyping of the USA national maize inbred seed bank. Genome Biology, 2013, 14, R55. | 8.8 | 458 |
| 4 | Gramene database in 2010: updates and extensions. Nucleic Acids Research, 2011, 39, D1085-D1094. | 14.5 | 182 |
| 5 | Gramene: a bird's eye view of cereal genomes. Nucleic Acids Research, 2006, 34, D717-D723. | 14.5 | 177 |
| 6 | Gramene: a growing plant comparative genomics resource. Nucleic Acids Research, 2007, 36, D947-D953. | 14.5 | 151 |
| 7 | Software engineering the mixed model for genome-wide association studies on large samples. Briefings in Bioinformatics, 2009, 10, 664-675. | 6.5 | 96 |
| 8 | A sorghum practical haplotype graph facilitates genomeâ€wide imputation and costâ€effective genomic prediction. Plant Genome, 2020, 13, e20009. | 2.8 | 54 |
| 9 | Genetic Architecture of Domestication-Related Traits in Maize. Genetics, 2016, 204, 99-113. | 2.9 | 39 |
| 10 | GDPC: connecting researchers with multiple integrated data sources. Bioinformatics, 2004, 20, 2839-2840. | 4.1 | 9 |
| 11 | The Generation Challenge Programme Platform: Semantic Standards and Workbench for Crop Science. International Journal of Plant Genomics, 2008, 2008, 1-6. | 2.2 | 5 |
| 12 | A multiâ€ŧrait multiâ€locus stepwise approach for conducting GWAS on correlated traits. Plant Genome, 2022, 15, e20200. | 2.8 | 4 |