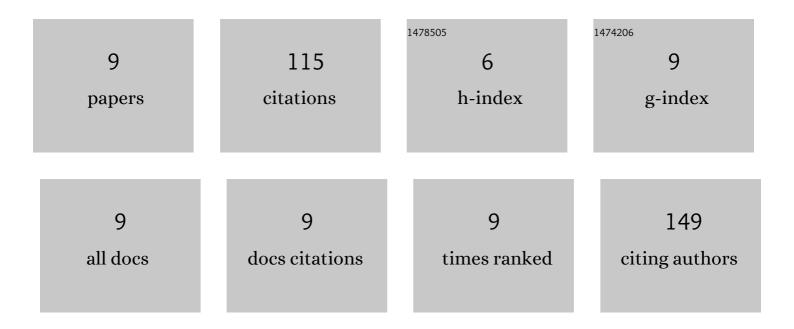
Daizhen Sun

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8747905/publications.pdf Version: 2024-02-01



| # | Article | IF | CITATIONS |
|---|--|-------------------|-------------|
| 1 | Association of yield-related traits in founder genotypes and derivatives of common wheat (Triticum) Tj ETQq1 1 C |).784314 r 3.6 | gBT /Overlo |
| 2 | Identification of TaPPH-7A haplotypes and development of a molecular marker associated with important agronomic traits in common wheat. BMC Plant Biology, 2019, 19, 296. | 3.6 | 20 |
| 3 | QTL analysis of wheat kernel traits, and genetic effects of qKW-6A on kernel width. Euphytica, 2019, 215, 1. | 1.2 | 18 |
| 4 | QTL mapping and candidate gene analysis of seed vigor-related traits during artificial aging in wheat (Triticum aestivum). Scientific Reports, 2020, 10, 22060. | 3.3 | 18 |
| 5 | QTL mapping for flag leaf-related traits and genetic effect of QFLW-6A on flag leaf width using two related introgression line populations in wheat. PLoS ONE, 2020, 15, e0229912. | 2.5 | 11 |
| 6 | In Silico and Transcription Analysis of Trehalose-6-phosphate Phosphatase Gene Family of Wheat: Trehalose Synthesis Genes Contribute to Salinity, Drought Stress and Leaf Senescence. Genes, 2021, 12, 1652. | 2.4 | 7 |
| 7 | Genome-wide identification and characterization of long non-coding RNAs related to grain yield in foxtail millet [Setaria italica (L.) P. Beauv.]. BMC Genomics, 2020, 21, 853. | 2.8 | 6 |
| 8 | Mining the stable quantitative trait loci for agronomic traits in wheat (Triticum aestivum L.) based on an introgression line population. BMC Plant Biology, 2020, 20, 275. | 3.6 | 6 |
| 9 | Quantitative trait loci mapping and candidate gene analysis of stoma-related traits in wheat (<i>Triticum aestivum</i> L.) glumes. PeerJ, 2022, 10, e13262. | 2.0 | 1 |