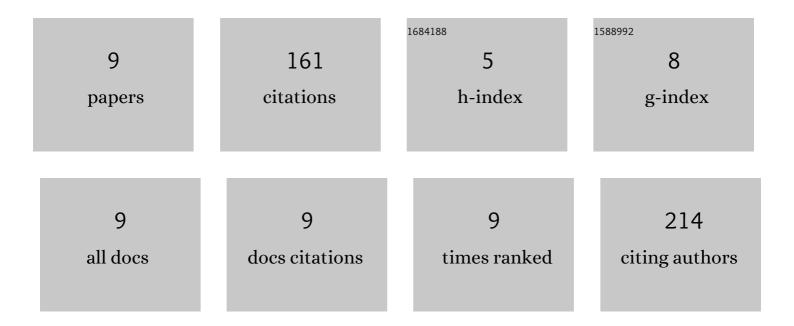
Xiaoyuan Tao

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8708962/publications.pdf

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Χιλοχιμλη Τλο

| # | Article | IF | CITATIONS |
|---|--|-----|-----------|
| 1 | Profiling of H3K4me3 Modification in Plants using Cleavage under Targets and Tagmentation. Journal of Visualized Experiments, 2022, , . | 0.3 | 0 |
| 2 | Functional analysis of the cotton CLE polypeptide signaling gene family in plant growth and development. Scientific Reports, 2021, 11, 5060. | 3.3 | 7 |
| 3 | Neofunctionalization of a polyploidization-activated cotton long intergenic non-coding RNA <i>DAN1</i> during drought stress regulation. Plant Physiology, 2021, 186, 2152-2168. | 4.8 | 27 |
| 4 | Absence of CG methylation alters the long noncoding transcriptome landscape in multiple species. FEBS Letters, 2021, 595, 1734-1747. | 2.8 | 4 |
| 5 | Functional examination of IncRNAs in allotetraploid Gossypium hirsutum. BMC Genomics, 2021, 22, 443. | 2.8 | 4 |
| 6 | Efficient chromatin profiling of H3K4me3 modification in cotton using CUT&Tag. Plant Methods, 2020, 16, 120. | 4.3 | 31 |
| 7 | Role of phasiRNAs from two distinct phasing frames of GhMYB2 loci in cis- gene regulation in the cotton genome. BMC Plant Biology, 2020, 20, 219. | 3.6 | 5 |
| 8 | Pseudomonas species isolated via high-throughput screening significantly protect cotton plants against verticillium wilt. AMB Express, 2020, 10, 193. | 3.0 | 24 |
| 9 | LncRNAs in polyploid cotton interspecific hybrids are derived from transposon neofunctionalization. Genome Biology, 2018, 19, 195. | 8.8 | 59 |