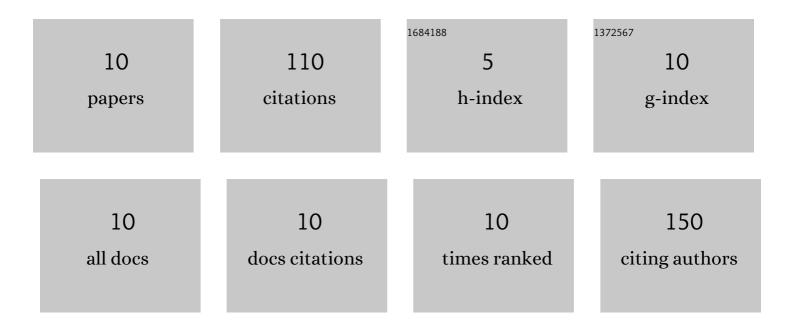
Xuehai Hu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/828430/publications.pdf Version: 2024-02-01



Хиенлі Ни

| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 1 | A statistical framework for predicting critical regions of p53-dependent enhancers. Briefings in Bioinformatics, 2021, 22, . | 6.5 | 4 |
| 2 | Boosting predictabilities of agronomic traits in rice using bivariate genomic selection. Briefings in Bioinformatics, 2021, 22, . | 6.5 | 6 |
| 3 | PlantDeepSEA, a deep learning-based web service to predict the regulatory effects of genomic variants in plants. Nucleic Acids Research, 2021, 49, W523-W529. | 14.5 | 20 |
| 4 | TSPTFBS: a Docker image for trans-species prediction of transcription factor binding sites in plants. Bioinformatics, 2021, 37, 260-262. | 4.1 | 10 |
| 5 | Improved Prediction of Regulatory Element Using Hybrid Abelian Complexity Features with DNA Sequences. International Journal of Molecular Sciences, 2019, 20, 1704. | 4.1 | 5 |
| 6 | A directed learning strategy integrating multiple omic data improves genomic prediction. Plant Biotechnology Journal, 2019, 17, 2011-2020. | 8.3 | 29 |
| 7 | Genome-Wide Prediction of DNA Methylation Using DNA Composition and Sequence Complexity in Human. International Journal of Molecular Sciences, 2017, 18, 420. | 4.1 | 8 |
| 8 | Accurate prediction of nuclear receptors with conjoint triad feature. BMC Bioinformatics, 2015, 16, 402. | 2.6 | 25 |
| 9 | Arbitrarily Long Arithmetic Progressions for Continued Fractions of Laurent Series. Acta Mathematica Scientia, 2013, 33, 943-949. | 1.0 | 2 |
| 10 | Evaluation of the occurrence possibility of SNP in Brassica napus with sliding window features by using RBF networks. Wuhan University Journal of Natural Sciences, 2011, 16, 73-78. | 0.4 | 1 |