

# Xuehai Hu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/828430/publications.pdf>

Version: 2024-02-01

10  
papers

110  
citations

1684188

5  
h-index

1372567

10  
g-index

10  
all docs

10  
docs citations

10  
times ranked

150  
citing authors

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