

# Chinac Thammamongtham

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/399369/publications.pdf>

Version: 2024-02-01

10  
papers

190  
citations

1874746

5  
h-index

1637695

9  
g-index

10  
all docs

10  
docs citations

10  
times ranked

338  
citing authors

#	ARTICLE	IF	CITATIONS
1	Ensemble-AHTPpred: A Robust Ensemble Machine Learning Model Integrated With a New Composite Feature for Identifying Antihypertensive Peptides. <i>Frontiers in Genetics</i> , 2022, 13, 883766.	1.1	4
2	Ensemble-AMPPred: Robust AMP Prediction and Recognition Using the Ensemble Learning Method with a New Hybrid Feature for Differentiating AMPs. <i>Genes</i> , 2021, 12, 137.	1.0	20
3	Ensemble of Multiple Classifiers for Multilabel Classification of Plant Protein Subcellular Localization. <i>Life</i> , 2021, 11, 293.	1.1	13
4	Metabolic Regulation of Sugar Assimilation for Lipid Production in <i>Aspergillus oryzae</i> BCC7051 through Comparative Transcriptome Perspective. <i>Biology</i> , 2021, 10, 885.	1.3	5
5	Systematic genome analysis of a novel arachidonic acid-producing strain uncovered unique metabolic traits in the production of acetyl-CoA-derived products in <i>Mortierella</i> fungi. <i>Gene</i> , 2020, 741, 144559.	1.0	3
6	PSO-LocBact: A Consensus Method for Optimizing Multiple Classifier Results for Predicting the Subcellular Localization of Bacterial Proteins. <i>BioMed Research International</i> , 2019, 2019, 1-11.	0.9	5
7	Genome Characterization of Oleaginous <i>Aspergillus oryzae</i> BCC7051: A Potential Fungal-Based Platform for Lipid Production. <i>Current Microbiology</i> , 2018, 75, 57-70.	1.0	30
8	Networking Omic Data to Envisage Systems Biological Regulation. <i>Advances in Biochemical Engineering/Biotechnology</i> , 2016, 160, 121-141.	0.6	0
9	Integrative computational approach for genome-based study of microbial lipid-degrading enzymes. <i>World Journal of Microbiology and Biotechnology</i> , 2016, 32, 122.	1.7	4
10	Alternative routes of acetyl-CoA synthesis identified by comparative genomic analysis: involvement in the lipid production of oleaginous yeast and fungi. <i>Microbiology (United Kingdom)</i> , 2012, 158, 217-228.	0.7	106