

# Tsuyoshi Esaki

## List of Publications by Year in descending order

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Version: 2024-02-01

18  
papers

621  
citations

687363

13  
h-index

940533

16  
g-index

20  
all docs

20  
docs citations

20  
times ranked

922  
citing authors

#	ARTICLE	IF	CITATIONS
1	Tracking metabolites at single-cell resolution reveals metabolic dynamics during plant mitosis. <i>Plant Physiology</i> , 2022, , .	4.8	3
2	Development of an <i>In Silico</i> Prediction Model for P-glycoprotein Efflux Potential in Brain Capillary Endothelial Cells toward the Prediction of Brain Penetration. <i>Journal of Medicinal Chemistry</i> , 2021, 64, 2725-2738.	6.4	22
3	Appropriate Evaluation Measurements for Regression Models. <i>Chem-Bio Informatics Journal</i> , 2021, 21, 59-69.	0.3	1
4	Novel anti-flavivirus drugs targeting the nucleolar distribution of core protein. <i>Virology</i> , 2020, 541, 41-51.	2.4	15
5	Open Innovation Platform using Cloud-based Applications and Collaborative Space: A Case Study of Solubility Prediction Model Development. <i>Chem-Bio Informatics Journal</i> , 2020, 20, 5-18.	0.3	0
6	Estimation of relationships between chemical substructures and antibiotic resistance-related gene expression in bacteria: Adapting a canonical correlation analysis for small sample data of gathered features using consensus clustering. <i>Chem-Bio Informatics Journal</i> , 2020, 20, 58-61.	0.3	0
7	Computational Model To Predict the Fraction of Unbound Drug in the Brain. <i>Journal of Chemical Information and Modeling</i> , 2019, 59, 3251-3261.	5.4	16
8	Constructing an In Silico Three-Class Predictor of Human Intestinal Absorption With Caco-2 Permeability and Dried-DMSO Solubility. <i>Journal of Pharmaceutical Sciences</i> , 2019, 108, 3630-3639.	3.3	17
9	Development of Simplified in Vitro P-Glycoprotein Substrate Assay and in Silico Prediction Models To Evaluate Transport Potential of P-Glycoprotein. <i>Molecular Pharmaceutics</i> , 2019, 16, 1851-1863.	4.6	41
10	Development of an in silico prediction system of human renal excretion and clearance from chemical structure information incorporating fraction unbound in plasma as a descriptor. <i>Scientific Reports</i> , 2019, 9, 18782.	3.3	27
11	Data Curation can Improve the Prediction Accuracy of Metabolic Intrinsic Clearance. <i>Molecular Informatics</i> , 2019, 38, e1800086.	2.5	24
12	Predicting Fraction Unbound in Human Plasma from Chemical Structure: Improved Accuracy in the Low Value Ranges. <i>Molecular Pharmaceutics</i> , 2018, 15, 5302-5311.	4.6	95
13	Dual Roles of Glutathione in Ecdysone Biosynthesis and Antioxidant Function During Larval Development in <i>Drosophila</i> . <i>Genetics</i> , 2017, 207, 1519-1532.	2.9	27
14	Direct evidence of specific localization of sesquiterpenes and marchantin A in oil body cells of <i>Marchantia polymorpha</i> L. <i>Phytochemistry</i> , 2016, 130, 77-84.	2.9	52
15	Comprehensive chemical secretory measurement of single cells trapped in a micro-droplet array with mass spectrometry. <i>RSC Advances</i> , 2015, 5, 16968-16971.	3.6	22
16	Fluorescence Probing Live Single-cell Mass Spectrometry for Direct Analysis of Organelle Metabolism. <i>Analytical Sciences</i> , 2015, 31, 1211-1213.	1.6	28
17	Live Single-Cell Plant Hormone Analysis by Video-Mass Spectrometry. <i>Plant and Cell Physiology</i> , 2015, 56, 1287-1296.	3.1	39
18	Direct metabolomics for plant cells by live single-cell mass spectrometry. <i>Nature Protocols</i> , 2015, 10, 1445-1456.	12.0	192