Ayako Yachie

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8154049/publications.pdf

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		1040056	1058476	
13	724	9	14	
papers	citations	h-index	g-index	
14	14	14	1343	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	Lenvatinib plus anti-PD-1 antibody combination treatment activates CD8+ T cells through reduction of tumor-associated macrophage and activation of the interferon pathway. PLoS ONE, 2019, 14, e0212513.	2.5	294
2	Paradoxical ATP Elevation in Ischemic Penumbra Revealed by Quantitative Imaging Mass Spectrometry. Antioxidants and Redox Signaling, 2010, 13, 1157-1167.	5.4	102
3	CD24 tracks divergent pluripotent states in mouse and human cells. Nature Communications, 2015, 6, 7329.	12.8	76
4	Modeling signalingâ€dependent pluripotency with Boolean logic to predict cell fate transitions. Molecular Systems Biology, 2018, 14, e7952.	7.2	49
5	In silico modeling and metabolome analysis of long-stored erythrocytes to improve blood storage methods. Journal of Biotechnology, 2009, 144, 212-223.	3.8	46
6	Dynamic Simulation and Metabolome Analysis of Long-Term Erythrocyte Storage in Adenine–Guanosine Solution. PLoS ONE, 2013, 8, e71060.	2.5	34
7	Hypotaurine is an Energy-Saving Hepatoprotective Compound against Ischemia-Reperfusion Injury of the Rat Liver. Journal of Clinical Biochemistry and Nutrition, 2010, 46, 126-134.	1.4	34
8	Artificial intelligence-based computational framework for drug-target prioritization and inference of novel repositionable drugs for Alzheimer's disease. Alzheimer's Research and Therapy, 2021, 13, 92.	6.2	29
9	A Metabolic Model of Human Erythrocytes: Practical Application of the E-Cell Simulation Environment. Journal of Biomedicine and Biotechnology, 2010, 2010, 1-14.	3.0	24
10	IQCELL: A platform for predicting the effect of gene perturbations on developmental trajectories using single-cell RNA-seq data. PLoS Computational Biology, 2022, 18, e1009907.	3.2	13
11	Disease-related cellular protein networks differentially affected under different EGFR mutations in lung adenocarcinoma. Scientific Reports, 2020, 10, 10881.	3.3	9
12	LimeMap: a comprehensive map of lipid mediator metabolic pathways. Npj Systems Biology and Applications, 2021, 7, 6.	3.0	8
13	Effects of maoto (ma-huang-tang) on host lipid mediator and transcriptome signature in influenza virus infection. Scientific Reports, 2021, 11, 4232.	3.3	5