Yukiko Matsuoka

List of Publications by Year in descending order

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687363 888059 3,307 16 13 17 citations h-index g-index papers 17 17 17 5252 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	A comprehensive pathway map of epidermal growth factor receptor signaling. Molecular Systems Biology, 2005, 1, 2005.0010.	7.2	902
2	The Systems Biology Graphical Notation. Nature Biotechnology, 2009, 27, 735-741.	17.5	828
3	CellDesigner 3.5: A Versatile Modeling Tool for Biochemical Networks. Proceedings of the IEEE, 2008, 96, 1254-1265.	21.3	380
4	Integrating Pathways of Parkinson's Disease in a Molecular Interaction Map. Molecular Neurobiology, 2014, 49, 88-102.	4.0	231
5	Software for systems biology: from tools to integrated platforms. Nature Reviews Genetics, 2011, 12, 821-832.	16.3	228
6	A comprehensive map of the mTOR signaling network. Molecular Systems Biology, 2010, 6, 453.	7.2	201
7	systemsDock: a web server for network pharmacology-based prediction and analysis. Nucleic Acids Research, 2016, 44, W507-W513.	14.5	135
8	AlzPathway: a comprehensive map of signaling pathways of Alzheimer's disease. BMC Systems Biology, 2012, 6, 52.	3.0	114
9	A comprehensive map of the influenza A virus replication cycle. BMC Systems Biology, 2013, 7, 97.	3.0	97
10	A comprehensive molecular interaction map of the budding yeast cell cycle. Molecular Systems Biology, 2010, 6, 415.	7.2	62
11	Modeling and Simulation Using CellDesigner. Methods in Molecular Biology, 2014, 1164, 121-145.	0.9	60
12	Network analyses based on comprehensive molecular interaction maps reveal robust control structures in yeast stress response pathways. Npj Systems Biology and Applications, 2016, 2, 15018.	3.0	27
13	Deconstructing the traditional Japanese medicine "Kampo― compounds, metabolites and pharmacological profile of maoto, a remedy for flu-like symptoms. Npj Systems Biology and Applications, 2017, 3, 32.	3.0	25
14	LimeMap: a comprehensive map of lipid mediator metabolic pathways. Npj Systems Biology and Applications, 2021, 7, 6.	3.0	8
15	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
16	Multi-dimensional computational pipeline for large-scale deep screening of compound effect assessment: an in silico case study on ageing-related compounds. Npj Systems Biology and Applications, 2019, 5, 42.	3.0	3