

Yoshinori Muto

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1925541/publications.pdf>

Version: 2024-02-01

13
papers

188
citations

1478505

6
h-index

1199594

12
g-index

13
all docs

13
docs citations

13
times ranked

444
citing authors

#	ARTICLE	IF	CITATIONS
1	N-Terminally extended human ubiquitin-conjugating enzymes (E2s) mediate the ubiquitination of RING-finger proteins, ARA54 and RNF8. <i>FEBS Journal</i> , 2001, 268, 2725-2732.	0.2	88
2	Molecular Evolution of the TET Gene Family in Mammals. <i>International Journal of Molecular Sciences</i> , 2015, 16, 28472-28485.	4.1	22
3	Adaptive Evolution of Formyl Peptide Receptors in Mammals. <i>Journal of Molecular Evolution</i> , 2015, 80, 130-141.	1.8	21
4	An evolutionarily conserved leucine-rich repeat protein CLERC is a centrosomal protein required for spindle pole integrity. <i>Cell Cycle</i> , 2008, 7, 2738-2748.	2.6	13
5	Heterotrimeric G protein G β s subunit attenuates PLEKHG2, a Rho family-specific guanine nucleotide exchange factor, by direct interaction. <i>Cellular Signalling</i> , 2017, 32, 115-123.	3.6	10
6	Climacostol inhibits <i>Tetrahymena</i> motility and mitochondrial respiration. <i>Open Life Sciences</i> , 2011, 6, 99-104.	1.4	8
7	Genome-wide evidence of positive selection in <i>Bacteroides fragilis</i> . <i>Computational Biology and Chemistry</i> , 2014, 52, 43-50.	2.3	7
8	Systems Level Analysis and Identification of Pathways and Key Genes Associated with Delirium. <i>Genes</i> , 2020, 11, 1225.	2.4	7
9	CLERC and centrosomal leucine-rich repeat proteins. <i>Open Life Sciences</i> , 2010, 5, 1-10.	1.4	4
10	Network and Evolutionary Analysis of Human Epigenetic Regulators to Unravel Disease Associations. <i>Genes</i> , 2020, 11, 1457.	2.4	4
11	Evolutionary Features and Intracellular Behavior of the PRTB Protein. <i>Biochemical Genetics</i> , 2011, 49, 458-473.	1.7	3
12	Genome-wide analyses reveal genes subject to positive selection in <i>Toxoplasma gondii</i> . <i>Gene</i> , 2019, 699, 73-79.	2.2	1
13	A network-based analysis of the human TET Gene Family. <i>Biologia (Poland)</i> , 2018, 73, 415-423.	1.5	0