Kishor Dhaygude

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9335777/publications.pdf

Version: 2024-02-01

	1163117	1125743	
377	8	13	
citations	h-index	g-index	
1.0	1.0	0.40	
18	18	848	
docs citations	times ranked	citing authors	
	citations 18	377 8 citations h-index 18 18	

#	Article	IF	CITATIONS
1	Not Only for Egg Yolkâ€"Functional and Evolutionary Insights from Expression, Selection, and Structural Analyses of Formica Ant Vitellogenins. Molecular Biology and Evolution, 2014, 31, 2181-2193.	8.9	78
2	Identification of Optimum Sequencing Depth Especially for De Novo Genome Assembly of Small Genomes Using Next Generation Sequencing Data. PLoS ONE, 2013, 8, e60204.	2.5	73
3	Ancient Duplications Have Led to Functional Divergence of Vitellogenin-Like Genes Potentially Involved in Inflammation and Oxidative Stress in Honey Bees. Genome Biology and Evolution, 2016, 8, 495-506.	2.5	60
4	A Metatranscriptomic Approach to the Identification of Microbiota Associated with the Ant Formica exsecta. PLoS ONE, 2013, 8, e79777.	2.5	52
5	Donor Simvastatin Treatment in Heart Transplantation. Circulation, 2019, 140, 627-640.	1.6	24
6	Evaluating responses to temperature during pre-metamorphosis and carry-over effects at post-metamorphosis in the wood tiger moth ($\langle i \rangle$ Arctia plantaginis $\langle i \rangle$). Philosophical Transactions of the Royal Society B: Biological Sciences, 2019, 374, 20190295.	4.0	21
7	The first draft genomes of the ant Formica exsecta, and its Wolbachia endosymbiont reveal extensive gene transfer from endosymbiont to host. BMC Genomics, 2019, 20, 301.	2.8	18
8	Instability of natural selection at candidate barrier loci underlying speciation in wood ants. Molecular Ecology, 2020, 29, 3988-3999.	3.9	13
9	Genome organization and molecular characterization of the three <i>Formica exsecta</i> virusesâ€"FeV1, FeV2 and FeV4. PeerJ, 2019, 6, e6216.	2.0	13
10	De novo transcriptome assembly and its annotation for the aposematic wood tiger moth (Parasemia) Tj ETQq0 C	0_rgBT /C	Overlock 10 Tf
11	Transcriptome sequencing reveals high isoform diversity in the ant <i>Formica exsecta</i> . PeerJ, 2017, 5, e3998.	2.0	7
12	Plasma proteome of brain-dead organ donors predicts heart transplant outcome. Journal of Heart and Lung Transplantation, 2022, 41, 311-324.	0.6	7
13	Inhibition of Vascular Endothelial Growth Factor Receptors 1 and 2 Attenuates Natural Killer Cell and Innate Immune Responses in an Experimental Model for Obliterative Bronchiolitis. American Journal of Pathology, 2022, 192, 254-269.	3.8	3
14	Transcriptome Profiles of the Circulating Extracellular Vesicles in Acute Lung Allograft Rejection. Journal of Heart and Lung Transplantation, 2019, 38, S146.	0.6	0
15	EFFECT OF DONOR SIMVASTATIN TREATMENT ON METABOLIC PROFILING IN RECIPIENTS DURING ISCHEMIA-REPERFUSION INJURY. Transplantation, 2020, 104, S174-S174.	1.0	0
16	DONOR SIMVASTATIN TREATMENT INFLUENCES TRANSCRIPTOMIC PROFILES OF EXTRACELLULAR VESICLES IN RECIPIENTS AFTER HEART TRANSPLANTATION. Transplantation, 2020, 104, S170-S170.	1.0	0
17	BRAIN DEATH ALTERS ECTONUCLEOTIDASE ACTIVITIES AND PURINE NUCLEOTIDE LEVELS IN HUMAN PLASMA. Transplantation, 2020, 104, S101-S101.	1.0	0