Kishor Dhaygude

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	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
2	Plasma proteome of brain-dead organ donors predicts heart transplant outcome. Journal of Heart and Lung Transplantation, 2022, 41, 311-324.	0.6	7
3	Instability of natural selection at candidate barrier loci underlying speciation in wood ants. Molecular Ecology, 2020, 29, 3988-3999.	3.9	13
4	EFFECT OF DONOR SIMVASTATIN TREATMENT ON METABOLIC PROFILING IN RECIPIENTS DURING ISCHEMIA-REPERFUSION INJURY. Transplantation, 2020, 104, S174-S174.	1.0	0
5	DONOR SIMVASTATIN TREATMENT INFLUENCES TRANSCRIPTOMIC PROFILES OF EXTRACELLULAR VESICLES IN RECIPIENTS AFTER HEART TRANSPLANTATION. Transplantation, 2020, 104, S170-S170.	1.0	0
6	BRAIN DEATH ALTERS ECTONUCLEOTIDASE ACTIVITIES AND PURINE NUCLEOTIDE LEVELS IN HUMAN PLASMA. Transplantation, 2020, 104, S101-S101.	1.0	0
7	Donor Simvastatin Treatment in Heart Transplantation. Circulation, 2019, 140, 627-640.	1.6	24
8	Evaluating responses to temperature during pre-metamorphosis and carry-over effects at post-metamorphosis in the wood tiger moth (<i>Arctia plantaginis</i>). Philosophical Transactions of the Royal Society B: Biological Sciences, 2019, 374, 20190295.	4.0	21
9	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
10	Transcriptome Profiles of the Circulating Extracellular Vesicles in Acute Lung Allograft Rejection. Journal of Heart and Lung Transplantation, 2019, 38, S146.	0.6	0
11	Genome organization and molecular characterization of the three <i>Formica exsecta</i> viruses—FeV1, FeV2 and FeV4. PeerJ, 2019, 6, e6216.	2.0	13
12	De novo transcriptome assembly and its annotation for the aposematic wood tiger moth (Parasemia) Tj ETQq0 C) 0_rgBT /C)verlock 10 T
13	Transcriptome sequencing reveals high isoform diversity in the ant <i>Formica exsecta</i> . PeerJ, 2017, 5, e3998.	2.0	7

14	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
15	Not Only for Egg Yolk—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
16	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
17	A Metatranscriptomic Approach to the Identification of Microbiota Associated with the Ant Formica exsecta. PLoS ONE, 2013, 8, e79777.	2.5	52