

Carlos Alberto Oliveira de Biagi Junior

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7516058/publications.pdf>

Version: 2024-02-01

15
papers

704
citations

1478505

6
h-index

1372567

10
g-index

17
all docs

17
docs citations

17
times ranked

1645
citing authors

#	ARTICLE	IF	CITATIONS
1	Upregulation of the novel lncRNA U731166 is associated with migration, invasion and vemurafenib resistance in melanoma. <i>Journal of Cellular and Molecular Medicine</i> , 2022, 26, 671-683.	3.6	6
2	RANKL Impairs the TLR4 Pathway by Increasing TRAF6 and RANK Interaction in Macrophages. <i>BioMed Research International</i> , 2022, 2022, 1-13.	1.9	3
3	Ultraconserved long non-coding RNA uc.112 is highly expressed in childhood T versus B-cell acute lymphoblastic leukemia. <i>Hematology, Transfusion and Cell Therapy</i> , 2021, 43, 28-34.	0.2	15
4	Transcriptomic studies on <i>Purpureocillium lilacinum</i> reveal molecular mechanisms of response to fluconazole and itraconazole. <i>Brazilian Journal of Microbiology</i> , 2021, 52, 491-501.	2.0	1
5	Multidimensional Single-Nuclei RNA-Seq Reconstruction of Adipose Tissue Reveals Adipocyte Plasticity Underlying Thermogenic Response. <i>Cells</i> , 2021, 10, 3073.	4.1	11
6	YAP1 Is a Potential Predictive Molecular Biomarker for Response to SMO Inhibitor in Medulloblastoma Cells. <i>Cancers</i> , 2021, 13, 6249.	3.7	1
7	Elevated Glucose Levels Favor SARS-CoV-2 Infection and Monocyte Response through a HIF-1 α /Glycolysis-Dependent Axis. <i>Cell Metabolism</i> , 2020, 32, 437-446.e5.	16.2	578
8	Lipopolysaccharide triggers different transcriptional signatures in taurine and indicine cattle macrophages: Reactive oxygen species and potential outcomes to the development of immune response to infections. <i>PLoS ONE</i> , 2020, 15, e0241861.	2.5	5
9	Title is missing!. , 2020, 15, e0241861.		0
10	Title is missing!. , 2020, 15, e0241861.		0
11	Title is missing!. , 2020, 15, e0241861.		0
12	Title is missing!. , 2020, 15, e0241861.		0
13	Whole transcriptome analysis reveals correlation of long noncoding RNA ZEB1-AS1 with invasive profile in melanoma. <i>Scientific Reports</i> , 2019, 9, 11350.	3.3	21
14	A simplified approach using Taqman low-density array for medulloblastoma subgrouping. <i>Acta Neuropathologica Communications</i> , 2019, 7, 33.	5.2	18
15	Mitochondrial transcription factor A (TFAM) shapes metabolic and invasion gene signatures in melanoma. <i>Scientific Reports</i> , 2018, 8, 14190.	3.3	41