

Jorge Andrade

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

25
papers

1,694
citations

14
h-index

26
g-index

26
ext. papers

2,427
ext. citations

12.3
avg, IF

3.8
L-index

#	Paper	IF	Citations
25	Transcriptomic Changes of Murine Visceral Fat Exposed to Intermittent Hypoxia at Single Cell Resolution. <i>International Journal of Molecular Sciences</i> , 2020 , 22,	6.3	1
24	Circulating plasma exosomes in obstructive sleep apnoea and reverse dipping blood pressure. <i>European Respiratory Journal</i> , 2020 , 55,	13.6	9
23	Circulating Exosomal miRNAs Signal Circadian Misalignment to Peripheral Metabolic Tissues. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	11
22	Pathogenesis of Human Papillomaviruses Requires the ATR/p62 Autophagy-Related Pathway. <i>MBio</i> , 2020 , 11,	7.8	4
21	Proteomics reveals NNMT as a master metabolic regulator of cancer-associated fibroblasts. <i>Nature</i> , 2019 , 569, 723-728	50.4	155
20	Topoisomerase IIβ binding protein 1 activates expression of E2F1 and p73 in HPV-positive cells for genome amplification upon epithelial differentiation. <i>Oncogene</i> , 2019 , 38, 3274-3287	9.2	5
19	Late gestational intermittent hypoxia induces metabolic and epigenetic changes in male adult offspring mice. <i>Journal of Physiology</i> , 2017 , 595, 2551-2568	3.9	35
18	Visceral White Adipose Tissue after Chronic Intermittent and Sustained Hypoxia in Mice. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2017 , 56, 477-487	5.7	49
17	miRNA profiling of human naive CD4 T cells links miR-34c-5p to cell activation and HIV replication. <i>EMBO Journal</i> , 2017 , 36, 346-360	13	19
16	Tumor Cell Malignant Properties Are Enhanced by Circulating Exosomes in Sleep Apnea. <i>Chest</i> , 2016 , 150, 1030-1041	5.3	40
15	Genomics of Ovarian Cancer Progression Reveals Diverse Metastatic Trajectories Including Intraepithelial Metastasis to the Fallopian Tube. <i>Cancer Discovery</i> , 2016 , 6, 1342-1351	24.4	116
14	Epigenomic Diversity in a Global Collection of Arabidopsis thaliana Accessions. <i>Cell</i> , 2016 , 166, 492-505	56.2	353
13	1,135 Genomes Reveal the Global Pattern of Polymorphism in Arabidopsis thaliana. <i>Cell</i> , 2016 , 166, 481-491	56.2	620
12	Post-Transcriptional Regulation of KLF4 by High-Risk Human Papillomaviruses Is Necessary for the Differentiation-Dependent Viral Life Cycle. <i>PLoS Pathogens</i> , 2016 , 12, e1005747	7.6	24
11	Extracellular microvesicle microRNAs in children with sickle cell anaemia with divergent clinical phenotypes. <i>British Journal of Haematology</i> , 2016 , 174, 786-98	4.5	31
10	Effect on Intermittent Hypoxia on Plasma Exosomal Micro RNA Signature and Endothelial Function in Healthy Adults. <i>Sleep</i> , 2016 , 39, 2077-2090	1.1	52
9	Circulating Plasma Extracellular Microvesicle MicroRNA Cargo and Endothelial Dysfunction in Children with Obstructive Sleep Apnea. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2016 , 194, 1116-1126	10.2	74

8	ExScalibur: A High-Performance Cloud-Enabled Suite for Whole Exome Germline and Somatic Mutation Identification. <i>PLoS ONE</i> , 2015 , 10, e0135800	3-7	11
7	Linkage analysis of autopsy-confirmed familial Alzheimer disease supports an Alzheimer disease locus in 8q24. <i>Dementia and Geriatric Cognitive Disorders</i> , 2011 , 31, 109-18	2-6	4
6	Linkage to 20p13 including the ANGPT4 gene in families with mixed Alzheimer's disease and vascular dementia. <i>Journal of Human Genetics</i> , 2010 , 55, 649-55	4-3	8
5	Expanded high-resolution genetic study of 109 Swedish families with Alzheimer's disease. <i>European Journal of Human Genetics</i> , 2008 , 16, 202-8	5-3	17
4	The epitope space of the human proteome. <i>Protein Science</i> , 2008 , 17, 606-13	6-3	32
3	The use of grid computing to drive data-intensive genetic research. <i>European Journal of Human Genetics</i> , 2007 , 15, 694-702	5-3	13
2	Applications of Grid Computing in Genetics and Proteomics 2006 , 791-798		2
1	Using Grid technology for computationally intensive applied bioinformatics analyses. <i>In Silico Biology</i> , 2006 , 6, 495-504	2	5