

Andreia J Amaral

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7640159/publications.pdf>

Version: 2024-02-01

17
papers

1,033
citations

933264

10
h-index

1058333

14
g-index

17
all docs

17
docs citations

17
times ranked

1665
citing authors

#	ARTICLE	IF	CITATIONS
1	IsomiR_Window: a system for analyzing small-RNA-seq data in an integrative and user-friendly manner. BMC Bioinformatics, 2021, 22, 37.	1.2	3
2	Genomic Tools for the Conservation and Genetic Improvement of a Highly Fragmented Breedâ€”The Ramo Grande Cattle from the Azores. Animals, 2020, 10, 1089.	1.0	5
3	Combining genome-wide association analyses and gene interaction networks to reveal new genes associated with carcass traits, meat quality and fatty acid profiles in pigs. Livestock Science, 2019, 220, 180-189.	0.6	8
4	Role of mitofusin-2 in NAFLD and targeting by miRNAs. Journal of Hepatology, 2018, 68, S356.	1.8	0
5	miRNA profiling of human naive CD4 T cells links miR-34c5p to cell activation and HIV replication. EMBO Journal, 2017, 36, 346-360.	3.5	32
6	Activation of miR-34a contributes to liver and muscle insulin resistance in experimental and human non-alcoholic fatty liver disease. Journal of Hepatology, 2017, 66, S163.	1.8	1
7	Autoimmunity and allergy control in adults submitted to complete thymectomy early in infancy. PLoS ONE, 2017, 12, e0180385.	1.1	14
8	Stimulation of RAC1/PAK1 signalling upregulates DNA damage repair genes via the BCL6/STAT5-switch. Annals of Oncology, 2017, 28, v19-v20.	0.6	0
9	Prospects and challenges for the conservation of farm animal genomic resources, 2015-2025. Frontiers in Genetics, 2015, 6, 314.	1.1	64
10	P0944 : Modulation of muscle microRNA expression profiles in patients with NAFLD and in C2C12 cells incubated with palmitic acid. Journal of Hepatology, 2015, 62, S699.	1.8	0
11	Quality assessment and control of tissue specific RNA-seq libraries of Drosophila transgenic RNAi models. Frontiers in Genetics, 2014, 5, 43.	1.1	14
12	A long journey to effective obesity treatments: is there light at the end of the tunnel?. Experimental Biology and Medicine, 2013, 238, 491-501.	1.1	10
13	Expression Profile of microRNAs Regulating Proliferation and Differentiation in Mouse Adult Cardiac Stem Cells. PLoS ONE, 2013, 8, e63041.	1.1	17
14	Genome-Wide Footprints of Pig Domestication and Selection Revealed through Massive Parallel Sequencing of Pooled DNA. PLoS ONE, 2011, 6, e14782.	1.1	135
15	Design of a High Density SNP Genotyping Assay in the Pig Using SNPs Identified and Characterized by Next Generation Sequencing Technology. PLoS ONE, 2009, 4, e6524.	1.1	568
16	Application of massive parallel sequencing to whole genome SNP discovery in the porcine genome. BMC Genomics, 2009, 10, 374.	1.2	44
17	Linkage Disequilibrium Decay and Haplotype Block Structure in the Pig. Genetics, 2008, 179, 569-579.	1.2	118