

Fuduan Peng

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9129130/publications.pdf>

Version: 2024-02-01

12
papers

390
citations

1040056

9
h-index

1199594

12
g-index

13
all docs

13
docs citations

13
times ranked

799
citing authors

#	ARTICLE	IF	CITATIONS
1	Differential expression analysis at the individual level reveals a lncRNA prognostic signature for lung adenocarcinoma. <i>Molecular Cancer</i> , 2017, 16, 98.	19.2	101
2	Autophagy-related prognostic signature for breast cancer. <i>Molecular Carcinogenesis</i> , 2016, 55, 292-299.	2.7	68
3	Meta-analysis of genome-wide association studies identifies 8 novel loci involved in shape variation of human head hair. <i>Human Molecular Genetics</i> , 2018, 27, 559-575.	2.9	51
4	Systematic feature selection improves accuracy of methylation-based forensic age estimation in Han Chinese males. <i>Forensic Science International: Genetics</i> , 2018, 35, 38-45.	3.1	46
5	The BRCA1/2-directed miRNA signature predicts a good prognosis in ovarian cancer patients with wild-type BRCA1/2. <i>Oncotarget</i> , 2015, 6, 2397-2406.	1.8	36
6	EMT-Regulome: a database for EMT-related regulatory interactions, motifs and network. <i>Cell Death and Disease</i> , 2017, 8, e2872-e2872.	6.3	21
7	Genome-wide association studies and CRISPR/Cas9-mediated gene editing identify regulatory variants influencing eyebrow thickness in humans. <i>PLoS Genetics</i> , 2018, 14, e1007640.	3.5	20
8	Genome-Wide Association Studies Identify Multiple Genetic Loci Influencing Eyebrow Color Variation in Europeans. <i>Journal of Investigative Dermatology</i> , 2019, 139, 1601-1605.	0.7	17
9	Validation of methylation-based forensic age estimation in time-series bloodstains on FTA cards and gauze at room temperature conditions. <i>Forensic Science International: Genetics</i> , 2019, 40, 168-174.	3.1	11
10	A landscape of synthetic viable interactions in cancer. <i>Briefings in Bioinformatics</i> , 2018, 19, bbw142.	6.5	9
11	A genome-wide association study identifies <i>FSHR</i> rs2300441 associated with follicle-stimulating hormone levels. <i>Clinical Genetics</i> , 2020, 97, 869-877.	2.0	8
12	Population-scale genetic control of alternative polyadenylation and its association with human diseases. <i>Quantitative Biology</i> , 2022, 10, 44-54.	0.5	2