

Meng Li

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9882977/publications.pdf>

Version: 2024-02-01

10
papers

1,591
citations

1478505

6
h-index

1474206

9
g-index

10
all docs

10
docs citations

10
times ranked

3078
citing authors

#	ARTICLE	IF	CITATIONS
1	Exosome and Exosomal MicroRNA: Trafficking, Sorting, and Function. Genomics, Proteomics and Bioinformatics, 2015, 13, 17-24.	6.9	1,466
2	Small extracellular vesicles derived from hypoxic mesenchymal stem cells promote vascularized bone regeneration through the miR-210-3p/EFNA3/PI3K pathway. Acta Biomaterialia, 2022, 150, 413-426.	8.3	38
3	Tumor-derived circulating exosomal miR-342-5p and miR-574-5p as promising diagnostic biomarkers for early-stage Lung Adenocarcinoma. International Journal of Medical Sciences, 2020, 17, 1428-1438.	2.5	33
4	Secretome of Activated Fibroblasts Induced by Exosomes for the Discovery of Biomarkers in Nonâ€Small Cell Lung Cancer. Small, 2021, 17, e2004750.	10.0	18
5	Dietary Intervention With Î±-Amylase Inhibitor in White Kidney Beans Added Yogurt Modulated Gut Microbiota to Adjust Blood Glucose in Mice. Frontiers in Nutrition, 2021, 8, 664976.	3.7	12
6	Heterogeneous Nuclear Ribonucleoprotein A1 Loads Batched Tumor-Promoting MicroRNAs Into Small Extracellular Vesicles With the Assist of Caveolin-1 in A549 Cells. Frontiers in Cell and Developmental Biology, 2021, 9, 687912.	3.7	9
7	The long noncoding RNA HOTAIRM1 controlled by AML1 enhances glucocorticoid resistance by activating RHOA/ROCK1 pathway through suppressing ARHGAP18. Cell Death and Disease, 2021, 12, 702.	6.3	8
8	Disequilibrium in chicken gut microflora with avian colibacillosis is related to microenvironment damaged by antibiotics. Science of the Total Environment, 2021, 762, 143058.	8.0	6
9	RBUD: A New Functional Potential Analysis Approach for Whole Microbial Genome Shotgun Sequencing. Microorganisms, 2020, 8, 1563.	3.6	1
10	Identification of Potential Key lncRNAs in the Context of Mouse Myeloid Differentiation by Systematic Transcriptomics Analysis. Genes, 2021, 12, 630.	2.4	0