

Yuanyuan Wei

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

2,783
citations

19
h-index

27
g-index

27
ext. papers

3,276
ext. citations

11.3
avg, IF

4.83
L-index

#	Paper	IF	Citations
25	piRNA-guided intron removal from pre-mRNAs regulates density-dependent reproductive strategy.. <i>Cell Reports</i> , 2022 , 39, 110593	10.6	0
24	A host lipase prevents lipopolysaccharide-induced foam cell formation. <i>IScience</i> , 2021 , 24, 103004	6.1	0
23	A Novel Regulatory Player in the Innate Immune System: Long Non-Coding RNAs. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	2
22	HIF-1[[Hypoxia-Inducible Factor-1]]Promotes Macrophage Necroptosis by Regulating miR-210 and miR-383. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2020 , 40, 583-596	9.4	31
21	Modulators of MicroRNA Function in the Immune System. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	16
20	miR-103 promotes endothelial maladaptation by targeting lncWDR59. <i>Nature Communications</i> , 2018 , 9, 2645	17.4	40
19	Macrophage MicroRNAs as Therapeutic Targets for Atherosclerosis, Metabolic Syndrome, and Cancer. <i>International Journal of Molecular Sciences</i> , 2018 , 19,	6.3	18
18	Dicer in Macrophages Prevents Atherosclerosis by Promoting Mitochondrial Oxidative Metabolism. <i>Circulation</i> , 2018 , 138, 2007-2020	16.7	54
17	Hyperlipidemia-Induced MicroRNA-155-5p Improves ECell Function by Targeting. <i>Diabetes</i> , 2017 , 66, 3072-3084	0.9	25
16	Dicer generates a regulatory microRNA network in smooth muscle cells that limits neointima formation during vascular repair. <i>Cellular and Molecular Life Sciences</i> , 2017 , 74, 359-372	10.3	13
15	Endothelial Dicer promotes atherosclerosis and vascular inflammation by miRNA-103-mediated suppression of KLF4. <i>Nature Communications</i> , 2016 , 7, 10521	17.4	81
14	MicroRNA-276 promotes egg-hatching synchrony by up-regulating brm in locusts. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 584-9	11.5	57
13	MicroRNA regulation of macrophages in human pathologies. <i>Cellular and Molecular Life Sciences</i> , 2016 , 73, 3473-95	10.3	43
12	Regulation of Csf1r and Bcl6 in macrophages mediates the stage-specific effects of microRNA-155 on atherosclerosis. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2015 , 35, 796-803	9.4	63
11	Evidence for the expression of abundant microRNAs in the locust genome. <i>Scientific Reports</i> , 2015 , 5, 13608	4.9	20
10	MicroRNA-126-5p promotes endothelial proliferation and limits atherosclerosis by suppressing Dlk1. <i>Nature Medicine</i> , 2014 , 20, 368-76	50.5	427
9	Cardiac fibroblast-derived microRNA passenger strand-enriched exosomes mediate cardiomyocyte hypertrophy. <i>Journal of Clinical Investigation</i> , 2014 , 124, 2136-46	15.9	617

8	MicroRNA-133 inhibits behavioral aggregation by controlling dopamine synthesis in locusts. <i>PLoS Genetics</i> , 2014 , 10, e1004206	6	71
7	MicroRNA-126, -145, and -155: a therapeutic triad in atherosclerosis?. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2013 , 33, 449-54	9.4	166
6	The microRNA-342-5p fosters inflammatory macrophage activation through an Akt1- and microRNA-155-dependent pathway during atherosclerosis. <i>Circulation</i> , 2013 , 127, 1609-19	16.7	163
5	Pathogenic arterial remodeling: the good and bad of microRNAs. <i>American Journal of Physiology - Heart and Circulatory Physiology</i> , 2013 , 304, H1050-9	5.2	91
4	The role of microRNAs in arterial remodelling. <i>Thrombosis and Haemostasis</i> , 2012 , 107, 611-8	7	84
3	MicroRNA-155 promotes atherosclerosis by repressing Bcl6 in macrophages. <i>Journal of Clinical Investigation</i> , 2012 , 122, 4190-202	15.9	359
2	De novo analysis of transcriptome dynamics in the migratory locust during the development of phase traits. <i>PLoS ONE</i> , 2010 , 5, e15633	3.7	188
1	Characterization and comparative profiling of the small RNA transcriptomes in two phases of locust. <i>Genome Biology</i> , 2009 , 10, R6	18.3	146