

# Long-Fei Wu

## List of Publications by Year in descending order

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Version: 2024-02-01

28  
papers

725  
citations

933447

10  
h-index

580821

25  
g-index

28  
all docs

28  
docs citations

28  
times ranked

961  
citing authors

#	ARTICLE	IF	CITATIONS
1	miR-137 regulates ferroptosis by targeting glutamine transporter SLC1A5 in melanoma. <i>Cell Death and Differentiation</i> , 2018, 25, 1457-1472.	11.2	308
2	miR-9 regulates ferroptosis by targeting glutamic oxaloacetic transaminase GOT1 in melanoma. <i>Molecular Carcinogenesis</i> , 2018, 57, 1566-1576.	2.7	125
3	Rheumatoid arthritis-associated DNA methylation sites in peripheral blood mononuclear cells. <i>Annals of the Rheumatic Diseases</i> , 2019, 78, 36-42.	0.9	75
4	Exosome: An Emerging Source of Biomarkers for Human Diseases. <i>Current Molecular Medicine</i> , 2019, 19, 387-394.	1.3	30
5	Identification of novel rheumatoid arthritis-associated MiRNA-204-5p from plasma exosomes. <i>Experimental and Molecular Medicine</i> , 2022, 54, 334-345.	7.7	27
6	Association of Plasma Irisin with Bone Mineral Density in a Large Chinese Population Using an Extreme Sampling Design. <i>Calcified Tissue International</i> , 2018, 103, 246-251.	3.1	23
7	Relative abundance of mature myostatin rather than total myostatin is negatively associated with bone mineral density in Chinese. <i>Journal of Cellular and Molecular Medicine</i> , 2018, 22, 1329-1336.	3.6	21
8	Identification of PBMC-expressed miRNAs for rheumatoid arthritis. <i>Epigenetics</i> , 2020, 15, 386-397.	2.7	16
9	SAMD9 is a (epi-) genetically regulated anti-inflammatory factor activated in RA patients. <i>Molecular and Cellular Biochemistry</i> , 2019, 456, 135-144.	3.1	15
10	Correlation analyses revealed global microRNA-mRNA expression associations in human peripheral blood mononuclear cells. <i>Molecular Genetics and Genomics</i> , 2018, 293, 95-105.	2.1	12
11	Identification and evaluation of lncRNA and mRNA integrative modules in human peripheral blood mononuclear cells. <i>Epigenomics</i> , 2017, 9, 943-954.	2.1	11
12	Multi-omics integrative analysis identified SNP-methylation-mRNA: Interaction in peripheral blood mononuclear cells. <i>Journal of Cellular and Molecular Medicine</i> , 2019, 23, 4601-4610.	3.6	10
13	Detection of lncRNA-mRNA interaction modules by integrating eQTL with weighted gene co-expression network analysis. <i>Functional and Integrative Genomics</i> , 2019, 19, 217-225.	3.5	9
14	Integrative lncRNA-mRNA co-expression network analysis identifies novel lncRNA E2F3-IT1 for rheumatoid arthritis. <i>Clinical and Translational Medicine</i> , 2021, 11, e325.	4.0	7
15	Alteration of circulating microbiome and its associated regulation role in rheumatoid arthritis: Evidence from integration of multiomics data. <i>Clinical and Translational Medicine</i> , 2020, 10, e229.	4.0	5
16	ITGA2 protein is associated with rheumatoid arthritis in Chinese and affects cellular function of T cells. <i>Clinica Chimica Acta</i> , 2021, 523, 208-215.	1.1	5
17	Association of High Serum Chemerin with Bone Mineral Density Loss and Osteoporotic Fracture in Elderly Chinese Women. <i>International Journal of Women's Health</i> , 2022, Volume 14, 107-118.	2.6	5
18	Plasma gelsolin is associated with hip BMD in Chinese postmenopausal women. <i>PLoS ONE</i> , 2018, 13, e0197732.	2.5	4

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19	Evaluation of plasma cytokine protein array profile: the highlighted PDGF-BB in rheumatoid arthritis. <i>Clinical Rheumatology</i> , 2020, 39, 3323-3330.	2.2	4
20	Integrative Analysis Confirmed the Association between Osteoprotegerin and Osteoporosis. <i>Chinese Medical Sciences Journal</i> , 2019, 34, 55.	0.4	3
21	The distribution and functional relevance analysis of runs of homozygosity (ROHs) in Chinese Han female population. <i>Molecular Genetics and Genomics</i> , 2018, 293, 197-206.	2.1	2
22	Abl interactor 1: A novel biomarker for osteoporosis in Chinese elderly men. <i>Journal of Proteomics</i> , 2019, 207, 103440.	2.4	2
23	Protein array test detected three osteoporosis related plasma inflammatory cytokines in Chinese postmenopausal women. <i>Cytokine</i> , 2020, 133, 155166.	3.2	2
24	Epigenetically-regulated RPN2 gene influences lymphocyte activation and is involved in pathogenesis of rheumatoid arthritis. <i>Gene</i> , 2022, 810, 146059.	2.2	2
25	Why SNP rs3755955 is associated with human bone mineral density? A molecular and cellular study in bone cells. <i>Molecular and Cellular Biochemistry</i> , 2021, , 1.	3.1	1
26	Body Surface Area (BSA) is a Better Osteoporosis Associated Anthropometric Parameter Than Other Anthropometric Parameters in Elderly Population. <i>Journal of Clinical Densitometry</i> , 2022, , .	1.2	1
27	Integrative analysis identified mediation effects of lncRNAs on the correlations between methylation and mRNA. <i>International Journal of Biochemistry and Cell Biology</i> , 2018, 104, 66-72.	2.8	0
28	Response to "Correspondence on "Rheumatoid arthritis-associated DNA methylation sites in peripheral blood mononuclear cells" by Wang and Niu. <i>Annals of the Rheumatic Diseases</i> , 2022, 81, e259-e259.	0.9	0