

# Yanling She

## List of Publications by Year in descending order

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Version: 2024-02-01

13  
papers

194  
citations

1163117

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h-index

1125743

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13  
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13  
docs citations

13  
times ranked

274  
citing authors

#	ARTICLE	IF	CITATIONS
1	A new perspective on depression and neuroinflammation: Non-coding RNA. <i>Journal of Psychiatric Research</i> , 2022, 148, 293-306.	3.1	17
2	Knockdown of CNN3 Impairs Myoblast Proliferation, Differentiation, and Protein Synthesis via the mTOR Pathway. <i>Frontiers in Physiology</i> , 2021, 12, 659272.	2.8	9
3	Lnc-GD2H Promotes Proliferation by Forming a Feedback Loop With c-Myc and Enhances Differentiation Through Interacting With NACA to Upregulate Myog in C2C12 Myoblasts. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 671857.	3.7	3
4	Roles of lncRNAs and circRNAs in regulating skeletal muscle development. <i>Acta Physiologica</i> , 2020, 228, e13356.	3.8	33
5	Regulation of Skeletal Muscle Atrophy in Cachexia by MicroRNAs and Long Non-coding RNAs. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 577010.	3.7	16
6	Role of miRNAs and lncRNAs in dexamethasone-induced myotube atrophy <i>in vitro</i> . <i>Experimental and Therapeutic Medicine</i> , 2020, 21, 146.	1.8	4
7	Differentially expressed coding and noncoding RNAs in CoCl <sub>2</sub> -induced cytotoxicity of C2C12 cells. <i>Epigenomics</i> , 2019, 11, 423-438.	2.1	7
8	Expression patterns of regulatory lncRNAs and miRNAs in muscular atrophy models induced by starvation <i>in vitro</i> and <i>in vivo</i> . <i>Molecular Medicine Reports</i> , 2019, 20, 4175-4185.	2.4	8
9	Necrostatin-1 protects C2C12 myotubes from CoCl <sub>2</sub> -induced hypoxia. <i>International Journal of Molecular Medicine</i> , 2018, 41, 2565-2572.	4.0	26
10	Comprehensive analysis of lncRNAs and mRNAs with associated co-expression and ceRNA networks in C2C12 myoblasts and myotubes. <i>Gene</i> , 2018, 647, 164-173.	2.2	28
11	Expression of circular RNAs during C2C12 myoblast differentiation and prediction of coding potential based on the number of open reading frames and N6-methyladenosine motifs. <i>Cell Cycle</i> , 2018, 17, 1832-1845.	2.6	26
12	Network-based analysis of the molecular mechanisms of multiple myeloma and monoclonal gammopathy of undetermined significance. <i>Oncology Letters</i> , 2017, 14, 4167-4175.	1.8	16
13	Transcriptome analysis of CD34+ cells from myelodysplastic syndrome patients. <i>Leukemia Research</i> , 2017, 62, 40-50.	0.8	1