

Lingmei Qian

List of Publications by Year in descending order

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Version: 2024-02-01

25
papers

488
citations

758635

12
h-index

676716

22
g-index

27
all docs

27
docs citations

27
times ranked

661
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of a novel native peptide derived from 60S ribosomal protein L23a that translationally regulates p53 to reduce myocardial ischemia-reperfusion. <i>Pharmacological Research</i> , 2022, 175, 105988.	3.1	4
2	Circular RNA Arhgap12 modulates doxorubicin-induced cardiotoxicity by sponging miR-135a-5p. <i>Life Sciences</i> , 2021, 265, 118788.	2.0	18
3	Peptide Szeto-Schiller 31 ameliorates doxorubicin-induced cardiotoxicity by inhibiting the activation of the p38MAPK signaling pathway. <i>International Journal of Molecular Medicine</i> , 2021, 47, .	1.8	15
4	Peptidomics analysis revealed that a novel peptide VMP-19 protects against AngII-induced injury in human umbilical vein endothelial cells. <i>Molecular Medicine Reports</i> , 2021, 23, .	1.1	2
5	The Impact of COVID-19 on Primary Care General Practice Consultations in a Teaching Hospital in Shanghai, China. <i>Frontiers in Medicine</i> , 2021, 8, 642496.	1.2	20
6	Exercise-induced peptide EIP-22 protect myocardial from ischaemia/reperfusion injury via activating JAK2/STAT3 signalling pathway. <i>Journal of Cellular and Molecular Medicine</i> , 2021, 25, 3560-3572.	1.6	11
7	Exercise-induced peptide TAG-23 protects cardiomyocytes from reperfusion injury through regulating PKG-cCbl interaction. <i>Basic Research in Cardiology</i> , 2021, 116, 41.	2.5	4
8	Peptidomics Analysis Reveals Peptide PDCryab1 Inhibits Doxorubicin-Induced Cardiotoxicity. <i>Oxidative Medicine and Cellular Longevity</i> , 2020, 2020, 1-23.	1.9	8
9	Expression profile of long non-coding RNAs in cardiomyocytes exposed to acute ischemic hypoxia. <i>Molecular Medicine Reports</i> , 2019, 19, 302-308.	1.1	6
10	An alternative under-valve approach to ablate right-sided accessory pathways. <i>Heart Rhythm</i> , 2019, 16, 51-56.	0.3	9
11	Long noncoding RNA uc.4 inhibits cell differentiation in heart development by altering DNA methylation. <i>Journal of Cellular Biochemistry</i> , 2019, 120, 8061-8068.	1.2	6
12	The long non-coding RNA uc.4 influences cell differentiation through the TGF-beta signaling pathway. <i>Experimental and Molecular Medicine</i> , 2018, 50, e447-e447.	3.2	24
13	Altered DNA Methylation of Long Noncoding RNA uc.167 Inhibits Cell Differentiation in Heart Development. <i>BioMed Research International</i> , 2018, 2018, 1-9.	0.9	7
14	Attenuation of Na/K-ATPase/Src/ROS amplification signal pathway with pNaktide ameliorates myocardial ischemia-reperfusion injury. <i>International Journal of Biological Macromolecules</i> , 2018, 118, 1142-1148.	3.6	33
15	Peptidomics Analysis of Transient Regeneration in the Neonatal Mouse Heart. <i>Journal of Cellular Biochemistry</i> , 2017, 118, 2828-2840.	1.2	18
16	LncRNA-uc.167 influences cell proliferation, apoptosis and differentiation of P19 cells by regulating Mef2c. <i>Gene</i> , 2016, 590, 97-108.	1.0	34
17	MicroRNA-29c overexpression inhibits proliferation and promotes apoptosis and differentiation in P19 embryonal carcinoma cells. <i>Gene</i> , 2016, 576, 304-311.	1.0	9
18	Cardiac-Specific PID1 Overexpression Enhances Pressure Overload-Induced Cardiac Hypertrophy in Mice. <i>Cellular Physiology and Biochemistry</i> , 2015, 35, 1975-1985.	1.1	8

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19	Î±-Lipoic acid ameliorates mitochondrial impairment and reverses apoptosis in FABP3-overexpressing embryonic cancer cells. <i>Journal of Bioenergetics and Biomembranes</i> , 2013, 45, 459-466.	1.0	10
20	Silencing of FABP3 Inhibits Proliferation and Promotes Apoptosis in Embryonic Carcinoma Cells. <i>Cell Biochemistry and Biophysics</i> , 2013, 66, 139-146.	0.9	15
21	Identification of maternal serum microRNAs as novel non-invasive biomarkers for prenatal detection of fetal congenital heart defects. <i>Clinica Chimica Acta</i> , 2013, 424, 66-72.	0.5	84
22	Effects of miR-19b Overexpression on Proliferation, Differentiation, Apoptosis and Wnt/Î²-Catenin Signaling Pathway in P19 Cell Model of Cardiac Differentiation In Vitro. <i>Cell Biochemistry and Biophysics</i> , 2013, 66, 709-722.	0.9	47
23	Integrated Analysis of Dysregulated lncRNA Expression in Fetal Cardiac Tissues with Ventricular Septal Defect. <i>PLoS ONE</i> , 2013, 8, e77492.	1.1	41
24	Potential role of maternal serum microRNAs as a biomarker for fetal congenital heart defects. <i>Medical Hypotheses</i> , 2011, 76, 424-426.	0.8	34
25	Overexpression of NYGGF4 (PID1) induces mitochondrial impairment in 3T3-L1 adipocytes. <i>Molecular and Cellular Biochemistry</i> , 2010, 340, 41-48.	1.4	21