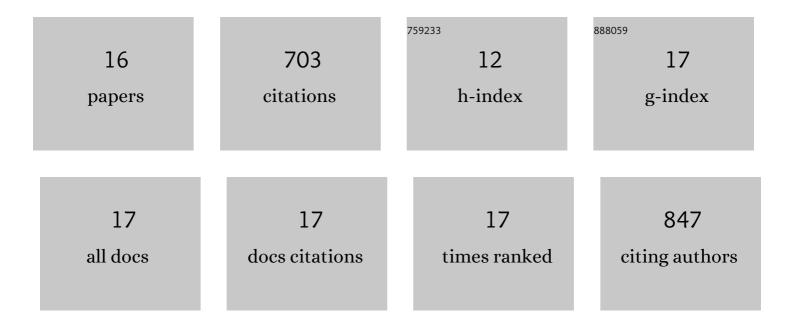
Zhongju Shi

List of Publications by Year in descending order

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7номени Shi

#	Article	IF	CITATIONS
1	Microglia-organized scar-free spinal cord repair in neonatal mice. Nature, 2020, 587, 613-618.	27.8	197
2	Programmed cell death in spinal cord injury pathogenesis and therapy. Cell Proliferation, 2021, 54, e12992.	5.3	101
3	The roles of microRNAs in spinal cord injury. International Journal of Neuroscience, 2017, 127, 1104-1115.	1.6	67
4	Effectiveness of Teriparatide on Fracture Healing: A Systematic Review and Meta-Analysis. PLoS ONE, 2016, 11, e0168691.	2.5	58
5	Epidemiological Features of Spinal Cord Injury in China: A Systematic Review. Frontiers in Neurology, 2018, 9, 683.	2.4	50
6	The emerging role of long non oding <scp>RNA</scp> in spinal cord injury. Journal of Cellular and Molecular Medicine, 2018, 22, 2055-2061.	3.6	44
7	Aquatic Exercises in the Treatment of Low Back Pain. American Journal of Physical Medicine and Rehabilitation, 2018, 97, 116-122.	1.4	42
8	Exploring the key genes and pathways of osteosarcoma with pulmonary metastasis using a gene expression microarray. Molecular Medicine Reports, 2017, 16, 7423-7431.	2.4	28
9	Signatures of altered long noncoding RNAs and messenger RNAs expression in the early acute phase of spinal cord injury. Journal of Cellular Physiology, 2019, 234, 8918-8927.	4.1	27
10	MicroRNAâ€29a regulates neural stem cell neuronal differentiation by targeting PTEN. Journal of Cellular Biochemistry, 2018, 119, 5813-5820.	2.6	26
11	Investigation of candidate long noncoding RNAs and messenger RNAs in the immediate phase of spinal cord injury based on gene expression profiles. Gene, 2018, 661, 119-125.	2.2	18
12	c-Jun Amino-Terminal Kinase is Involved in Valproic Acid-Mediated Neuronal Differentiation of Mouse Embryonic NSCs and Neurite Outgrowth of NSC-Derived Neurons. Neurochemical Research, 2017, 42, 1254-1266.	3.3	14
13	Identification and Verification of Candidate Genes Regulating Neural Stem Cells Behavior Under Hypoxia. Cellular Physiology and Biochemistry, 2018, 47, 212-222.	1.6	9
14	Exploring the key genes and pathways in enchondromas using a gene expression microarray. Oncotarget, 2017, 8, 43967-43977.	1.8	7
15	Identification of differentially expressed proteins in rats with spinal cord injury during the transitional phase using an iTRAQ-based quantitative analysis. Gene, 2018, 677, 66-76.	2.2	7
16	Systematic analysis of miRNAs in patients with postmenopausal osteoporosis. Gynecological Endocrinology, 2020, 36, 997-1001.	1.7	7