Jun Wang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5255684/publications.pdf

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		932766	1199166	
12	722	10	12	
papers	citations	h-index	g-index	
13	13	13	1424	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	MIR99AHG is a noncoding tumor suppressor gene in lung adenocarcinoma. Cell Death and Disease, 2021, 12, 424.	2.7	24
2	Anatomical Pulmonary Sublobar Resection Based on Subsegment. Annals of Thoracic Surgery, 2021, 111, e447-e450.	0.7	13
3	A modified system for classifying the bilateral superior pulmonary veins using three-dimensional computed tomography bronchography and angiography images. Journal of Thoracic Disease, 2021, 13, 5933-5941.	0.6	5
4	Denoising Autoencoder, A Deep Learning Algorithm, Aids the Identification of A Novel Molecular Signature of Lung Adenocarcinoma. Genomics, Proteomics and Bioinformatics, 2020, 18, 468-480.	3.0	18
5	Plasma RNA sequencing of extracellular RNAs reveals potential biomarkers for non-small cell lung cancer. Clinical Biochemistry, 2020, 83, 65-73.	0.8	10
6	Threeâ€dimensional navigationâ€guided thoracoscopic combined subsegmentectomy for intersegmental pulmonary nodules. Thoracic Cancer, 2019, 10, 41-46.	0.8	30
7	The Circular RNA circPRKCI Promotes Tumor Growth in Lung Adenocarcinoma. Cancer Research, 2018, 78, 2839-2851.	0.4	211
8	Modified method for distinguishing the intersegmental border for lung segmentectomy. Thoracic Cancer, 2018, 9, 330-333.	0.8	41
9	Identification of circular RNAs in human aortic valves. Gene, 2018, 642, 135-144.	1.0	13
10	Technique for tailoring complex demarcation in lung segmentectomy. Thoracic Cancer, 2018, 9, 1562-1564.	0.8	13
11	Comprehensive characterization of tissue-specific circular RNAs in the human and mouse genomes. Briefings in Bioinformatics, 2017, 18, bbw081.	3.2	251
12	Frequent alterations in cytoskeleton remodelling genes in primary and metastatic lung adenocarcinomas. Nature Communications, 2015, 6, 10131.	5.8	93