

Bao Sen Zhou

List of Publications by Year in descending order

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44
papers

987
citations

516710

16
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477307

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all docs

46
docs citations

46
times ranked

1898
citing authors

#	ARTICLE	IF	CITATIONS
1	Sarcopenia and a 5-mRNA risk module as a combined factor to predict prognosis for patients with stomach adenocarcinoma. <i>Genomics</i> , 2022, 114, 361-377.	2.9	7
2	CircRNAs as promising biomarker in diagnostic and prognostic of lung cancer: An updated meta-analysis. <i>Genomics</i> , 2021, 113, 387-397.	2.9	10
3	Integrative analysis of exosomal microRNA-149-5p in lung adenocarcinoma. <i>Aging</i> , 2021, 13, 7382-7396.	3.1	7
4	Association between Coalmine Dust and Mortality Risk of Lung Cancer: A Meta-Analysis. <i>BioMed Research International</i> , 2021, 2021, 1-9.	1.9	6
5	The epidemiological trends in the burden of lung cancer attributable to PM2.5 exposure in China. <i>BMC Public Health</i> , 2021, 21, 737.	2.9	32
6	Exosomal miR-338-3p suppresses non-small-cell lung cancer cells metastasis by inhibiting CHL1 through the MAPK signaling pathway. <i>Cell Death and Disease</i> , 2021, 12, 1030.	6.3	25
7	Application of vancomycin in patients with augmented renal clearance. <i>European Journal of Hospital Pharmacy</i> , 2020, 27, 276-279.	1.1	9
8	Intermittent vs. continuous vancomycin infusion for gram-positive infections: A systematic review and meta-analysis. <i>Journal of Infection and Public Health</i> , 2020, 13, 591-597.	4.1	20
9	Population pharmacokinetics of vancomycin in Chinese patients with augmented renal clearance. <i>Journal of Infection and Public Health</i> , 2020, 13, 68-74.	4.1	22
10	Polymorphisms in the PVT1 Gene and Susceptibility to the Lung Cancer in a Chinese Northeast Population: a Case-control Study. <i>Journal of Cancer</i> , 2020, 11, 468-478.	2.5	7
11	<p><p>GINS2 Functions as a Key Gene in Lung Adenocarcinoma by WGCNA Co-Expression Network Analysis</p></p>. <i>OncoTargets and Therapy</i> , 2020, Volume 13, 6735-6746.	2.0	13
12	Association of miR-27a polymorphism with the risk of digestive system cancers. <i>Pathology Research and Practice</i> , 2020, 216, 153115.	2.3	3
13	Polymorphisms in Neuronal Growth Regulator 1 and Otoancorin Alternate the Susceptibility to Lung Cancer in Chinese Nonsmoking Females. <i>DNA and Cell Biology</i> , 2020, 39, 1657-1663.	1.9	0
14	Coexisting EGFR and TP53 Mutations in Lung Adenocarcinoma Patients Are Associated With COMP and ITGB8 Upregulation and Poor Prognosis. <i>Frontiers in Molecular Biosciences</i> , 2020, 7, 30.	3.5	16
15	Long Noncoding RNA RAET1K Enhances CCNE1 Expression and Cell Cycle Arrest of Lung Adenocarcinoma Cell by Sponging miRNA-135a-5p. <i>Frontiers in Genetics</i> , 2020, 10, 1348.	2.3	13
16	A comparison of the burden of lung cancer attributable to tobacco exposure in China and the USA. <i>Annals of Translational Medicine</i> , 2020, 8, 1412-1412.	1.7	5
17	Association Between Two Polymorphisms in the Promoter Region of miR-143/miR-145 and the Susceptibility of Lung Cancer in Northeast Chinese Nonsmoking Females. <i>DNA and Cell Biology</i> , 2019, 38, 814-823.	1.9	13
18	A Case/Control Study:<i>AGBL1</i> Polymorphism Related to Lung Cancer Risk in Chinese Nonsmoking Females. <i>DNA and Cell Biology</i> , 2019, 38, 1452-1459.	1.9	5

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19	Association Between Three Polymorphisms in <i>BMAL1</i> Genes and Risk of Lung Cancer in a Northeast Chinese Population. <i>DNA and Cell Biology</i> , 2019, 38, 1437-1443.	1.9	4
20	LncRNA NEAT1 polymorphisms and lung cancer susceptibility in a Chinese Northeast Han Population: A case-control study. <i>Pathology Research and Practice</i> , 2019, 215, 152723.	2.3	12
21	p53/miR-30a/Sox4 feedback loop mediates cellular proliferation, apoptosis, and migration of non-small cell lung cancer. <i>Journal of Cellular Physiology</i> , 2019, 234, 22884-22895.	4.1	31
22	SNPs in LncRNA genes are associated with non-small cell lung cancer in a Chinese population. <i>Journal of Clinical Laboratory Analysis</i> , 2019, 33, e22858.	2.1	12
23	Polymorphism in <i>CYP24A1</i> Is Associated with Lung Cancer Risk: A Case-Control Study in Chinese Female Nonsmokers. <i>DNA and Cell Biology</i> , 2019, 38, 243-249.	1.9	11
24	Association between miR-146a rs2910164 polymorphism and specific cancer susceptibility: an updated meta-analysis. <i>Familial Cancer</i> , 2018, 17, 459-468.	1.9	32
25	Genetic variants in lncRNA HOTAIR are associated with lung cancer susceptibility in a Chinese Han population in China: a case–control study. <i>Cancer Management and Research</i> , 2018, Volume 10, 5209-5218.	1.9	15
26	Polymorphisms in the H19 gene and the risk of lung Cancer among female never smokers in Shenyang, China. <i>BMC Cancer</i> , 2018, 18, 893.	2.6	26
27	The lncRNA myocardial infarction associated transcript-centric competing endogenous RNA network in non-small-cell lung cancer. <i>Cancer Management and Research</i> , 2018, Volume 10, 1155-1162.	1.9	17
28	A Meta-Analysis of miR-499 rs3746444 Polymorphism for Cancer Risk of Different Systems: Evidence From 65 Case-Control Studies. <i>Frontiers in Physiology</i> , 2018, 9, 737.	2.8	20
29	The Risk Factors of Acquiring Severe Hand, Foot, and Mouth Disease: A Meta-Analysis. <i>Canadian Journal of Infectious Diseases and Medical Microbiology</i> , 2018, 2018, 1-12.	1.9	27
30	Clinically Correlated MicroRNAs in the Diagnosis of Non-Small Cell Lung Cancer: A Systematic Review and Meta-Analysis. <i>BioMed Research International</i> , 2018, 2018, 1-14.	1.9	17
31	Polymorphism in lncRNA AC008392.1 and its interaction with smoking on the risk of lung cancer in a Chinese population. <i>Cancer Management and Research</i> , 2018, Volume 10, 1377-1387.	1.9	13
32	Polymorphism in lncRNA AC016683.6 and its interaction with smoking exposure on the susceptibility of lung cancer. <i>Cancer Cell International</i> , 2018, 18, 91.	4.1	7
33	Single Nucleotide Polymorphisms in HMGB1 Correlate with Lung Cancer Risk in the Northeast Chinese Han Population. <i>Molecules</i> , 2018, 23, 832.	3.8	10
34	Association Between Long Noncoding RNA <i>MEG3</i> Polymorphisms and Lung Cancer Susceptibility in Chinese Northeast Population. <i>DNA and Cell Biology</i> , 2018, 37, 812-820.	1.9	26
35	Study on polymorphisms in <i>CHRNA5/CHRNA3/CHRNA4</i> gene cluster and the associated with the risk of non-small cell lung cancer. <i>Oncotarget</i> , 2018, 9, 2435-2444.	1.8	9
36	<i>CHL1</i> gene polymorphisms increase lung cancer susceptibility. <i>Oncotarget</i> , 2018, 9, 13545-13550.	1.8	4

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37	MiR-146a polymorphism correlates with lung cancer risk in Chinese nonsmoking females. <i>Oncotarget</i> , 2017, 8, 2275-2283.	1.8	31
38	Multiple functional SNPs in differentially expressed genes modify risk and survival of non-small cell lung cancer in chinese female non-smokers. <i>Oncotarget</i> , 2017, 8, 18924-18934.	1.8	37
39	Polymorphisms in pre-miRNA genes and cooking oil fume exposure as well as their interaction on the risk of lung cancer in a Chinese nonsmoking female population. <i>OncoTargets and Therapy</i> , 2016, 9, 395.	2.0	24
40	Polymorphisms in miR-135a-2, miR-219-2 and miR-211 as well as their interaction with cooking oil fume exposure on the risk of lung cancer in Chinese nonsmoking females: a case-control study. <i>BMC Cancer</i> , 2016, 16, 751.	2.6	14
41	Association between polymorphisms in pre-miRNA genes and risk of lung cancer in a Chinese non-smoking female population. <i>Lung Cancer</i> , 2016, 94, 15-21.	2.0	54
42	Association of MicroRNA-149 Polymorphism with Lung Cancer Risk in Chinese Non-Smoking Female: A Case-Control Study. <i>PLoS ONE</i> , 2016, 11, e0163626.	2.5	11
43	Genetic Variations in TERT-CLPTM1L Genes and Risk of Lung Cancer in Chinese Women Nonsmokers. <i>PLoS ONE</i> , 2013, 8, e64988.	2.5	23
44	Genome-wide association analysis identifies new lung cancer susceptibility loci in never-smoking women in Asia. <i>Nature Genetics</i> , 2012, 44, 1330-1335.	21.4	286