

Fu-You Zhou

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

746
citations

933447

10
h-index

1058476

14
g-index

15
all docs

15
docs citations

15
times ranked

1256
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of a seven-cell cycle signature predicting overall survival for gastric cancer. <i>Aging</i> , 2022, 14, 3989-3999.	3.1	5
2	PIWI-interacting RNAs: Mitochondria-based biogenesis and functions in cancer. <i>Genes and Diseases</i> , 2021, 8, 603-622.	3.4	15
3	Focal amplifications are associated with chromothripsis events and diverse prognoses in gastric cardia adenocarcinoma. <i>Nature Communications</i> , 2021, 12, 6489.	12.8	27
4	piR-823 demonstrates tumor oncogenic activity in esophageal squamous cell carcinoma through DNA methylation induction via DNA methyltransferase 3B. <i>Pathology Research and Practice</i> , 2020, 216, 152848.	2.3	31
5	Association between polymorphisms in the CYP1A1, CYP2E1 and GSTM1 genes, and smoking, alcohol and upper digestive tract carcinomas in a high incidence area of northern China. <i>Oncology Letters</i> , 2019, 18, 1267-1277.	1.8	11
6	Decreased expression of SPINT1-AS1 and SPINT1 mRNA might be independent unfavorable prognostic indicators in esophageal squamous cell carcinoma. <i>OncoTargets and Therapy</i> , 2019, Volume 12, 4755-4763.	2.0	19
7	High expression of HLA-DQA1 predicts poor outcome in patients with esophageal squamous cell carcinoma in Northern China. <i>Medicine (United States)</i> , 2019, 98, e14454.	1.0	13
8	Association of genotypes of rs671 within <i>ALDH2</i> with risk for gastric cardia adenocarcinoma in the Chinese Han population in high- and low-incidence areas. <i>Cancer Biology and Medicine</i> , 2017, 14, 60-65.	3.0	8
9	Joint analysis of three genome-wide association studies of esophageal squamous cell carcinoma in Chinese populations. <i>Nature Genetics</i> , 2014, 46, 1001-1006.	21.4	148
10	Variations in the MHC Region Confer Risk to Esophageal Squamous Cell Carcinoma on the Subjects from High-Incidence Area in Northern China. <i>PLoS ONE</i> , 2014, 9, e90438.	2.5	12
11	Association between CYP1A1 polymorphisms and esophageal cancer: a meta-analysis. <i>Molecular Biology Reports</i> , 2013, 40, 6035-6042.	2.3	13
12	Genotypic variants at 2q33 and risk of esophageal squamous cell carcinoma in China: a meta-analysis of genome-wide association studies. <i>Human Molecular Genetics</i> , 2012, 21, 2132-2141.	2.9	58
13	Genome-wide association study of esophageal squamous cell carcinoma in Chinese subjects identifies a susceptibility locus at PLCE1. <i>Nature Genetics</i> , 2010, 42, 759-763.	21.4	383