

Li-dong Wang

List of Publications by Year in descending order

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Version: 2024-02-01

29
papers

1,172
citations

759190

12
h-index

501174

28
g-index

30
all docs

30
docs citations

30
times ranked

1912
citing authors

#	ARTICLE	IF	CITATIONS
1	Characterization of E-cadherin expression in normal mucosa, dysplasia and adenocarcinoma of gastric cardia and its influence on prognosis. <i>World Journal of Gastrointestinal Oncology</i> , 2022, 14, 265-277.	2.0	1
2	Identification of a seven-cell cycle signature predicting overall survival for gastric cancer. <i>Aging</i> , 2022, 14, 3989-3999.	3.1	5
3	Long non-coding RNA ZNF667-AS1 retards the development of esophageal squamous cell carcinoma via modulation of microRNA-1290-mediated PRUNE2. <i>Translational Oncology</i> , 2022, 21, 101371.	3.7	7
4	Preoperative Prognostic Nutritional Index is a Significant Predictor of Survival in Esophageal Squamous Cell Carcinoma Patients. <i>Nutrition and Cancer</i> , 2021, 73, 215-220.	2.0	9
5	Promotion of rs3746804 (p. L267P) polymorphism to intracellular SLC52A3a trafficking and riboflavin transportation in esophageal cancer cells. <i>Amino Acids</i> , 2021, 53, 1197-1209.	2.7	1
6	Preoperative maximal voluntary ventilation, hemoglobin, albumin, lymphocytes and platelets predict postoperative survival in esophageal squamous cell carcinoma. <i>World Journal of Gastroenterology</i> , 2021, 27, 321-335.	3.3	15
7	Development and validation of a prognostic nomogram model for Chinese patients with primary small cell carcinoma of the esophagus. <i>World Journal of Clinical Cases</i> , 2021, 9, 9011-9022.	0.8	4
8	Focal amplifications are associated with chromothripsis events and diverse prognoses in gastric cardia adenocarcinoma. <i>Nature Communications</i> , 2021, 12, 6489.	12.8	27
9	MicroRNA-155 acts as a diagnostic and prognostic biomarker for oesophageal squamous cell carcinoma. <i>Artificial Cells, Nanomedicine and Biotechnology</i> , 2020, 48, 977-982.	2.8	9
10	<i>MDM2</i> and its functional polymorphism SNP309 contribute to the development of esophageal carcinoma. <i>Journal of Gene Medicine</i> , 2019, 21, e3086.	2.8	5
11	Interaction of 22 risk SNPs with <i>Helicobacter pylori</i> infection and risk of gastric cardia adenocarcinoma. <i>Future Oncology</i> , 2019, 15, 3579-3585.	2.4	3
12	SLC52A3 expression is activated by NF- κ B p65/Rel-B and serves as a prognostic biomarker in esophageal cancer. <i>Cellular and Molecular Life Sciences</i> , 2018, 75, 2643-2661.	5.4	38
13	High-throughput screening of prostate cancer risk loci by single nucleotide polymorphisms sequencing. <i>Nature Communications</i> , 2018, 9, 2022.	12.8	66
14	RNA editing of <i>SLC22A3</i> drives early tumor invasion and metastasis in familial esophageal cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E4631-E4640.	7.1	78
15	CWAS follow-up study of esophageal squamous cell carcinoma identifies potential genetic loci associated with family history of upper gastrointestinal cancer. <i>Scientific Reports</i> , 2017, 7, 4642.	3.3	11
16	Truth telling for patients with esophageal squamous cell carcinoma in Henan, China. <i>Cancer Biology and Medicine</i> , 2017, 14, 83-89.	3.0	4
17	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
18	Novel genetic locus at MHC region for esophageal squamous cell carcinoma in Chinese populations. <i>PLoS ONE</i> , 2017, 12, e0177494.	2.5	6

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19	Updated incidence rates and risk factors of esophageal cancer in Nan'ao Island, a coastal high-risk area in southern China. <i>Ecological Management and Restoration</i> , 2016, 30, n/a-n/a.	0.4	11
20	Genome-wide association study of gastric adenocarcinoma in Asia: a comparison of associations between cardia and non-cardia tumours. <i>Gut</i> , 2016, 65, 1611-1618.	12.1	99
21	Suppression of BRD4 inhibits human hepatocellular carcinoma by repressing MYC and enhancing BIM expression. <i>Oncotarget</i> , 2016, 7, 2462-2474.	1.8	90
22	ANO1 protein as a potential biomarker for esophageal cancer prognosis and precancerous lesion development prediction. <i>Oncotarget</i> , 2016, 7, 24374-24382.	1.8	38
23	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
24	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
25	A sequence variant in the phospholipase C epsilon C2 domain is associated with esophageal carcinoma and esophagitis. <i>Molecular Carcinogenesis</i> , 2013, 52, 80-86.	2.7	15
26	Correlation of telomere length shortening with TP53 somatic mutations, polymorphisms and allelic loss in breast tumors and esophageal cancer. <i>Oncology Reports</i> , 2013, 29, 226-236.	2.6	17
27	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
28	Comparative genomic hybridization analysis of genetic aberrations associated with development of esophageal squamous cell carcinoma in Henan, China. <i>World Journal of Gastroenterology</i> , 2008, 14, 1828.	3.3	28
29	Immunohistochemical studies on Waf1p21, p16, pRb and p53 in human esophageal carcinomas and neighboring epithelia from a high-risk area in northern China. , 1997, 72, 746-751.		32