Dongying Gu

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58 888 15 27 g-index

60 1,056 ext. papers ext. citations 5.9 avg, IF L-index

#	Paper	IF	Citations
58	Genetic variants in lncRNA HOTAIR are associated with risk of colorectal cancer. <i>Mutagenesis</i> , 2015 , 30, 303-10	2.8	112
57	E2F1-induced upregulation of long noncoding RNA LINC00668 predicts a poor prognosis of gastric cancer and promotes cell proliferation through epigenetically silencing of CKIs. <i>Oncotarget</i> , 2016 , 7, 23212-26	3.3	57
56	Genome-wide analysis of long noncoding RNA signature in human colorectal cancer. <i>Gene</i> , 2015 , 556, 227-34	3.8	53
55	Clinical potential role of circulating microRNAs in early diagnosis of colorectal cancer patients. <i>Carcinogenesis</i> , 2014 , 35, 2723-30	4.6	51
54	The DNA repair gene APE1 T1349G polymorphism and cancer risk: a meta-analysis of 27 case-control studies. <i>Mutagenesis</i> , 2009 , 24, 507-12	2.8	50
53	Common genetic variation in ETV6 is associated with colorectal cancer susceptibility. <i>Nature Communications</i> , 2016 , 7, 11478	17.4	45
52	Genetic variants in noncoding PIWI-interacting RNA and colorectal cancer risk. <i>Cancer</i> , 2015 , 121, 2044-	5@ .4	43
51	Genetic variants in m6A modification genes are associated with colorectal cancer risk. <i>Carcinogenesis</i> , 2020 , 41, 8-17	4.6	30
50	Frequent KIT mutations in human gastrointestinal stromal tumors. <i>Scientific Reports</i> , 2014 , 4, 5907	4.9	30
49	Lack of association between the hOGG1 Ser326Cys polymorphism and breast cancer risk: evidence from 11 case-control studies. <i>Breast Cancer Research and Treatment</i> , 2010 , 122, 527-31	4.4	30
48	An inverse association between tea consumption and colorectal cancer risk. <i>Oncotarget</i> , 2017 , 8, 37367	-3,7;376	5 30
47	Circadian clock pathway genes associated with colorectal cancer risk and prognosis. <i>Archives of Toxicology</i> , 2018 , 92, 2681-2689	5.8	24
46	The DNA repair gene APE1 T1349G polymorphism and risk of gastric cancer in a Chinese population. <i>PLoS ONE</i> , 2011 , 6, e28971	3.7	22
45	Clinical significance of ALDH2 rs671 polymorphism in esophageal cancer: evidence from 31 case-control studies. <i>OncoTargets and Therapy</i> , 2015 , 8, 649-59	4.4	20
44	VEGF 936C>T polymorphism and breast cancer risk: evidence from 5,729 cases and 5,868 controls. Breast Cancer Research and Treatment, 2011 , 125, 489-93	4.4	20
43	Body mass index (BMI) trajectories and risk of colorectal cancer in the PLCO cohort. <i>British Journal of Cancer</i> , 2018 , 119, 130-132	8.7	15
42	Genetic mutation analysis of human gastric adenocarcinomas using ion torrent sequencing platform. <i>PLoS ONE</i> , 2014 , 9, e100442	3.7	15

(2018-2014)

41	Clinical significance of POU5F1P1 rs10505477 polymorphism in Chinese gastric cancer patients receving cisplatin-based chemotherapy after surgical resection. <i>International Journal of Molecular Sciences</i> , 2014 , 15, 12764-77	6.3	15
40	Polymorphism in one-carbon metabolism pathway affects survival of gastric cancer patients: Large and comprehensive study. <i>Oncotarget</i> , 2015 , 6, 9564-76	3.3	15
39	Polymorphism rs2682818 in miR-618 is associated with colorectal cancer susceptibility in a Han Chinese population. <i>Cancer Medicine</i> , 2018 , 7, 1194-1200	4.8	13
38	Combinations of single nucleotide polymorphisms identified in genome-wide association studies determine risk for colorectal cancer. <i>International Journal of Cancer</i> , 2019 , 145, 2661-2669	7.5	12
37	MDM2 SNP309 polymorphism is associated with colorectal cancer risk. <i>Scientific Reports</i> , 2014 , 4, 4851	4.9	12
36	Comparison of the efficacy and safety of S-1-based and capecitabine-based regimens in gastrointestinal cancer: a meta-analysis. <i>PLoS ONE</i> , 2014 , 9, e84230	3.7	12
35	Association study of genetic variants in estrogen metabolic pathway genes and colorectal cancer risk and survival. <i>Archives of Toxicology</i> , 2018 , 92, 1991-1999	5.8	12
34	Functional annotation of colorectal cancer susceptibility loci identifies MLH1 rs1800734 associated with MSI patients. <i>Gut</i> , 2016 , 65, 1227-8	19.2	11
33	TGFB1 T29C polymorphism and breast cancer risk: a meta-analysis based on 10,417 cases and 11,455 controls. <i>Breast Cancer Research and Treatment</i> , 2010 , 123, 857-61	4.4	10
32	The effects of genomic polymorphisms in one-carbon metabolism pathways on survival of gastric cancer patients received fluorouracil-based adjuvant therapy. <i>Scientific Reports</i> , 2016 , 6, 28019	4.9	9
31	A genetic variant located in the miR-532-5p-binding site of TGFBR1 is associated with the colorectal cancer risk. <i>Journal of Gastroenterology</i> , 2019 , 54, 141-148	6.9	9
30	Genetic variants in RPA1 associated with the response to oxaliplatin-based chemotherapy in colorectal cancer. <i>Journal of Gastroenterology</i> , 2019 , 54, 939-949	6.9	8
29	Functional polymorphisms in apoptosis pathway genes and survival in patients with gastric cancer. <i>Environmental and Molecular Mutagenesis</i> , 2014 , 55, 421-7	3.2	8
28	Associations of NR5A2 gene polymorphisms with the clinicopathological characteristics and survival of gastric cancer. <i>International Journal of Molecular Sciences</i> , 2014 , 15, 22902-17	6.3	8
27	Pyrotinib in the treatment of human epidermal growth factor receptor 2-positive metastatic breast cancer: A case report. <i>Medicine (United States)</i> , 2020 , 99, e20809	1.8	7
26	Sex hormones and genetic variants in hormone metabolic pathways associated with the risk of colorectal cancer. <i>Environment International</i> , 2020 , 137, 105543	12.9	7
25	Genetic variation in C12orf51 is associated with prognosis of intestinal-type gastric cancer in a Chinese population. <i>Biomedicine and Pharmacotherapy</i> , 2015 , 69, 133-8	7.5	7
24	Association of Antioxidative Enzymes Polymorphisms with Efficacy of Platin and Fluorouracil-Based Adjuvant Therapy in Gastric Cancer. <i>Cellular Physiology and Biochemistry</i> , 2018 , 48, 2247-2257	3.9	6

23	A MAP3k1 SNP predicts survival of gastric cancer in a Chinese population. <i>PLoS ONE</i> , 2014 , 9, e96083	3.7	6
22	Genetic variant in miR-21 binding sites is associated with colorectal cancer risk. <i>Journal of Cellular and Molecular Medicine</i> , 2019 , 23, 2012-2019	5.6	6
21	Exosomal circLPAR1 functions in colorectal cancer diagnosis and tumorigenesis through suppressing BRD4 via METTL3-eIF3h interaction <i>Molecular Cancer</i> , 2022 , 21, 49	42.1	6
20	Protein phosphatase magnesium-dependent 1lls a novel tumor marker and target in hepatocellular carcinoma. <i>Frontiers of Medicine</i> , 2016 , 10, 52-60	12	5
19	Evaluation of GWAS-Identified Genetic Variants for Gastric Cancer Survival. <i>EBioMedicine</i> , 2018 , 33, 82-8	37 .8	5
18	Vitamin B intake reduces the risk for colorectal cancer: a dose-response analysis. <i>European Journal of Nutrition</i> , 2019 , 58, 1591-1602	5.2	5
17	The efficacy and safety of irinotecan [] bevacizumab compared with oxaliplatin [] bevacizumab for metastatic colorectal cancer: A meta-analysis. <i>Medicine (United States)</i> , 2019 , 98, e17384	1.8	5
16	Association of microRNA-27a rs895819 polymorphism with the risk of cancer: An updated meta-analysis. <i>Gene</i> , 2020 , 728, 144185	3.8	4
15	A genetic variant in large tumor suppressor kinase 2 of Hippo signaling pathway contributes to prognosis of hepatocellular carcinoma. <i>OncoTargets and Therapy</i> , 2016 , 9, 1945-51	4.4	4
14	IKBKB rs2272736 is Associated with Gastric Cancer Survival. <i>Pharmacogenomics and Personalized Medicine</i> , 2020 , 13, 345-352	2.1	3
13	Genetic variants in circTUBB interacting with smoking can enhance colorectal cancer risk. <i>Archives of Toxicology</i> , 2020 , 94, 325-333	5.8	2
12	Genetic variants in Hippo signalling pathway-related genes affect the risk of colorectal cancer. <i>Archives of Toxicology</i> , 2021 , 95, 271-281	5.8	2
11	Assessment of the Diagnostic Efficiency of a Liquid Biopsy Assay for Early Detection of Gastric Cancer. <i>JAMA Network Open</i> , 2021 , 4, e2121129	10.4	2
10	LncRNA-422 suppresses the proliferation and growth of colorectal cancer cells by targeting SFPQ Clinical and Translational Medicine, 2022, 12, e664	5.7	1
9	The prognostic impacts of transcription factor polymorphisms in Chinese hepatocellular carcinoma patients. <i>Oncotarget</i> , 2017 , 8, 69823-69832	3.3	1
8	Evaluation of common genetic variants in vitamin E-related pathway genes and colorectal cancer susceptibility. <i>Archives of Toxicology</i> , 2021 , 95, 2523-2532	5.8	1
7	Circulating tumor cells: A surrogate to predict the effect of treatment and overall survival in gastric adenocarcinoma. <i>International Journal of Biological Markers</i> , 2021 , 36, 28-35	2.8	1
6	OncoVeeEMiniPDX-Guided Anticancer Treatment for Gastric Cancer Patients With Synchronous Liver Metastases: A Retrospective Cohort Analysis <i>Frontiers in Oncology</i> , 2021 , 11, 757383	5.3	O

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5	Implication of polymorphisms in the promoter region of apoptosis-related genes in survival of gastric cancer patients <i>Journal of Clinical Oncology</i> , 2014 , 32, e15062-e15062	2.2
4	Effect of polymorphism in one-carbon metabolism pathway on survival of gastric cancer patients in a large and comprehensive study <i>Journal of Clinical Oncology</i> , 2015 , 33, e15091-e15091	2.2
3	Effect of H3K27 acetylation activated-long noncoding RNA CCAT1 on cell proliferation and migration by regulating SPRY4 and HOXB13 expression in esophageal squamous carcinoma cell <i>Journal of Clinical Oncology</i> , 2016 , 34, e15566-e15566	2.2
2	Comprehensive genetic mutation analysis of human gastric adenocarcinomas <i>Journal of Clinical Oncology</i> , 2013 , 31, 4106-4106	2.2
1	Variation rs9929218 and risk of the colorectal Cancer and adenomas: A meta-analysis. <i>BMC Cancer</i> , 2021 , 21, 190	4.8