

Dongying Gu

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/5187165/dongying-gu-publications-by-year.pdf>
Version: 2024-04-09

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.
The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

58 papers	888 citations	15 h-index	27 g-index
60 ext. papers	1,056 ext. citations	5.9 avg, IF	3.78 L-index

#	Paper	IF	Citations
58	LncRNA-422 suppresses the proliferation and growth of colorectal cancer cells by targeting SFPQ.. <i>Clinical and Translational Medicine</i> , 2022 , 12, e664	5.7	1
57	Exosomal circLPA1 functions in colorectal cancer diagnosis and tumorigenesis through suppressing BRD4 via METTL3-eIF3h interaction.. <i>Molecular Cancer</i> , 2022 , 21, 49	42.1	6
56	OncoVeeMiniPDX-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	0
55	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
54	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
53	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
52	Variation rs9929218 and risk of the colorectal Cancer and adenomas: A meta-analysis. <i>BMC Cancer</i> , 2021 , 21, 190	4.8	
51	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
50	Genetic variants in m6A modification genes are associated with colorectal cancer risk. <i>Carcinogenesis</i> , 2020 , 41, 8-17	4.6	30
49	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
48	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
47	Genetic variants in circTUBB interacting with smoking can enhance colorectal cancer risk. <i>Archives of Toxicology</i> , 2020 , 94, 325-333	5.8	2
46	Association of microRNA-27a rs895819 polymorphism with the risk of cancer: An updated meta-analysis. <i>Gene</i> , 2020 , 728, 144185	3.8	4
45	IKBKB rs2272736 is Associated with Gastric Cancer Survival. <i>Pharmacogenomics and Personalized Medicine</i> , 2020 , 13, 345-352	2.1	3
44	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
43	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
42	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

41	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
40	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
39	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
38	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
37	Evaluation of GWAS-Identified Genetic Variants for Gastric Cancer Survival. <i>EBioMedicine</i> , 2018 , 33, 82-88	8.8	5
36	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
35	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
34	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
33	Circadian clock pathway genes associated with colorectal cancer risk and prognosis. <i>Archives of Toxicology</i> , 2018 , 92, 2681-2689	5.8	24
32	An inverse association between tea consumption and colorectal cancer risk. <i>Oncotarget</i> , 2017 , 8, 37367-37376	3.76	30
31	The prognostic impacts of transcription factor polymorphisms in Chinese hepatocellular carcinoma patients. <i>Oncotarget</i> , 2017 , 8, 69823-69832	3.3	1
30	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
29	Protein phosphatase magnesium-dependent 1 is a novel tumor marker and target in hepatocellular carcinoma. <i>Frontiers of Medicine</i> , 2016 , 10, 52-60	12	5
28	Functional annotation of colorectal cancer susceptibility loci identifies MLH1 rs1800734 associated with MSI patients. <i>Gut</i> , 2016 , 65, 1227-8	19.2	11
27	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
26	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	
25	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
24	Common genetic variation in ETV6 is associated with colorectal cancer susceptibility. <i>Nature Communications</i> , 2016 , 7, 11478	17.4	45

23	Genetic variants in noncoding PIWI-interacting RNA and colorectal cancer risk. <i>Cancer</i> , 2015 , 121, 2044-50.4	4.4	43
22	Genome-wide analysis of long noncoding RNA signature in human colorectal cancer. <i>Gene</i> , 2015 , 556, 227-34	3.8	53
21	Genetic variants in lncRNA HOTAIR are associated with risk of colorectal cancer. <i>Mutagenesis</i> , 2015 , 30, 303-10	2.8	112
20	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
19	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
18	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
17	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	
16	MDM2 SNP309 polymorphism is associated with colorectal cancer risk. <i>Scientific Reports</i> , 2014 , 4, 4851	4.9	12
15	Clinical potential role of circulating microRNAs in early diagnosis of colorectal cancer patients. <i>Carcinogenesis</i> , 2014 , 35, 2723-30	4.6	51
14	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
13	Frequent KIT mutations in human gastrointestinal stromal tumors. <i>Scientific Reports</i> , 2014 , 4, 5907	4.9	30
12	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
11	Genetic mutation analysis of human gastric adenocarcinomas using ion torrent sequencing platform. <i>PLoS ONE</i> , 2014 , 9, e100442	3.7	15
10	Clinical significance of POU5F1P1 rs10505477 polymorphism in Chinese gastric cancer patients receiving cisplatin-based chemotherapy after surgical resection. <i>International Journal of Molecular Sciences</i> , 2014 , 15, 12764-77	6.3	15
9	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
8	A MAP3k1 SNP predicts survival of gastric cancer in a Chinese population. <i>PLoS ONE</i> , 2014 , 9, e96083	3.7	6
7	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	
6	Comprehensive genetic mutation analysis of human gastric adenocarcinomas.. <i>Journal of Clinical Oncology</i> , 2013 , 31, 4106-4106	2.2	

5	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
4	VEGF 936C>T polymorphism and breast cancer risk: evidence from 5,729 cases and 5,868 controls. <i>Breast Cancer Research and Treatment</i> , 2011 , 125, 489-93	4.4	20
3	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
2	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
1	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