

Weifeng Tang

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Interleukin 1B rs16944 G>A polymorphism was associated with a decreased risk of esophageal cancer in a Chinese population. <i>Clinical Biochemistry</i> , 2013, 46, 1469-1473.	0.8	54
2	Association between the STK15 F31I Polymorphism and Cancer Susceptibility: A Meta-Analysis Involving 43,626 Subjects. <i>PLoS ONE</i> , 2013, 8, e82790.	1.1	38
3	Programmed death-1 (PD-1) polymorphisms in Chinese patients with esophageal cancer. <i>Clinical Biochemistry</i> , 2014, 47, 612-617.	0.8	35
4	Vascular endothelial growth factor A polymorphisms are associated with increased risk of coronary heart disease: a meta-analysis. <i>Oncotarget</i> , 2017, 8, 30539-30551.	0.8	33
5	Genetic variations in MTHFR and esophageal squamous cell carcinoma susceptibility in Chinese Han population. <i>Medical Oncology</i> , 2014, 31, 915.	1.2	26
6	CTLA4 tagging polymorphisms and risk of colorectal cancer: a case–control study involving 2,306 subjects. <i>OncoTargets and Therapy</i> , 2018, Volume 11, 4609-4619.	1.0	26
7	<i>Programmed death-1</i> polymorphisms is associated with risk of esophagogastric junction adenocarcinoma in the Chinese Han population: A case-control study involving 2,740 subjects. <i>Oncotarget</i> , 2017, 8, 39198-39208.	0.8	23
8	Association between <i>BTLA</i> polymorphisms and susceptibility to esophageal squamous cell carcinoma in the Chinese population. <i>Journal of Clinical Laboratory Analysis</i> , 2020, 34, e23221.	0.9	19
9	Association of CTLA-4 tagging polymorphisms and haplotypes with hepatocellular carcinoma risk. <i>Medicine (United States)</i> , 2019, 98, e16266.	0.4	17
10	Investigation of <i>IGF1</i>, <i>IGF2BP2</i>, and <i>IGFBP3</i> variants with lymph node status and esophagogastric junction adenocarcinoma risk. <i>Journal of Cellular Biochemistry</i> , 2019, 120, 5510-5518.	1.2	17
11	Lack of Association between Cytotoxic T-lymphocyte Antigen 4 (CTLA-4) -1722T/C (rs733618) Polymorphism and Cancer Risk: From a Case-Control Study to a Meta-Analysis. <i>PLoS ONE</i> , 2014, 9, e94039.	1.1	16
12	Investigation of cyclin D1 rs9344 G>A polymorphism in colorectal cancer: a meta-analysis involving 13,642 subjects. <i>OncoTargets and Therapy</i> , 2016, Volume 9, 6641-6650.	1.0	16
13	Relationship between IGF2BP2 and IGFBP3 polymorphisms and susceptibility to non-small-cell lung cancer: a case–control study in Eastern Chinese Han population. <i>Cancer Management and Research</i> , 2018, Volume 10, 2965-2975.	0.9	16
14	Relationship of PPARG, PPARGC1A, and PPARGC1B polymorphisms with susceptibility to hepatocellular carcinoma in an eastern Chinese Han population. <i>OncoTargets and Therapy</i> , 2018, Volume 11, 4651-4660.	1.0	16
15	Aurora-A V57I (rs1047972) Polymorphism and Cancer Susceptibility: A Meta-Analysis Involving 27,269 Subjects. <i>PLoS ONE</i> , 2014, 9, e90328.	1.1	16
16	TERT-CLPTM1L Rs401681 C>T Polymorphism Was Associated with a Decreased Risk of Esophageal Cancer in a Chinese Population. <i>PLoS ONE</i> , 2014, 9, e100667.	1.1	16
17	Investigation of <i>TCF7L2</i>, <i>LEP</i> and <i>LEPR</i> polymorphisms with esophageal squamous cell carcinomas. <i>Oncotarget</i> , 2017, 8, 109107-109119.	0.8	16
18	Programmed death-1 (PD-1) rs2227981 C > T polymorphism is associated with cancer susceptibility: a meta-analysis. <i>International Journal of Clinical and Experimental Medicine</i> , 2015, 8, 22278-85.	1.3	14

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19	RANK rs1805034 T>C Polymorphism Is Associated with Susceptibility of Esophageal Cancer in a Chinese Population. PLoS ONE, 2014, 9, e101705.	1.1	13
20	Uracil-DNA glycosylase (UNG) rs246079 G/A polymorphism is associated with decreased risk of esophageal cancer in a Chinese population. Medical Oncology, 2014, 31, 272.	1.2	13
21	Investigation of <i>LEP</i> and <i>LEPR</i> polymorphisms with the risk of hepatocellular carcinoma: a case–control study in Eastern Chinese Han population. OncoTargets and Therapy, 2018, Volume 11, 2083-2089.	1.0	13
22	<p>Association of miRNA-499 rs3746444 A>G variants with adenocarcinoma of esophagogastric junction (AEG) risk and lymph node status</p>. OncoTargets and Therapy, 2019, Volume 12, 6245-6252.	1.0	13
23	Investigation of <i>Cytotoxic T-lymphocyte antigen-4</i> polymorphisms in non-small cell lung cancer: a case–control study. Oncotarget, 2017, 8, 76634-76643.	0.8	13
24	Association of the interleukin-18 receptor 1 and interleukin-18 receptor accessory protein polymorphisms with the risk of esophageal cancer. Biomedical Reports, 2016, 4, 227-235.	0.9	12
25	Leptin rs2167270 G&A (G19A) polymorphism may decrease the risk of cancer: A case–control study and meta–analysis involving 19&989 subjects. Journal of Cellular Biochemistry, 2019, 120, 10998-11007.	1.2	12
26	Associations Between <i>microRNA</i> Polymorphisms and Development of Coronary Artery Disease: A Case–Control Study. DNA and Cell Biology, 2020, 39, 25-36.	0.9	12
27	The relationship between <i>IGF2BP2</i> and <i>PPARG</i> polymorphisms and susceptibility to esophageal squamous-cell carcinomas in the eastern Chinese Han population. OncoTargets and Therapy, 2017, Volume 10, 5525-5532.	1.0	11
28	Investigation of <i>leptin receptor</i>rs1137101 G&A polymorphism with cancer risk: evidence from 35936 subjects. Bioscience Reports, 2019, 39, .	1.1	11
29	Lack of Association Between CTLA-4 Genetic Polymorphisms and Noncardiac Gastric Cancer in a Chinese Population. DNA and Cell Biology, 2019, 38, 443-448.	0.9	11
30	Investigation of <i>BTLA</i> tagging variants with risk of esophagogastric junction adenocarcinoma. Bioscience Reports, 2019, 39, .	1.1	11
31	Association of PPARG rs 1801282 C>G polymorphism with risk of colorectal cancer: from a case-control study to a meta-analysis. Oncotarget, 2017, 8, 100558-100569.	0.8	11
32	Methylenetetrahydrofolate reductase tagging polymorphisms are associated with risk of non-small cell lung cancer in eastern Chinese Han population. Oncotarget, 2017, 8, 110326-110336.	0.8	11
33	miRNA-146a rs2910164 C&G polymorphism increased the risk of esophagogastric junction adenocarcinoma: a case–control study involving 2,740 participants. Cancer Management and Research, 2018, Volume 10, 1657-1664.	0.9	10
34	Association of ALDH3B2 gene polymorphism and risk factors with susceptibility of esophageal squamous cell carcinoma in a Chinese population: a case-control study involving 2,358 subjects. Oncotarget, 2017, 8, 110153-110165.	0.8	10
35	p21 rs3176352 G>C and p73 rs1801173 C>T Polymorphisms Are Associated with an Increased Risk of Esophageal Cancer in a Chinese Population. PLoS ONE, 2014, 9, e96958.	1.1	9
36	Investigation of the Association of <i>miRNA-499</i>, <i>miRNA-146a</i>, <i>miRNA-196a2</i> Loci with Hepatocellular Carcinoma Risk: A Case–Control Study Involving 1507 Subjects. DNA and Cell Biology, 2020, 39, 379-388.	0.9	9

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37	Association between methylenetetrahydrofolate reductase tagging polymorphisms and susceptibility of hepatocellular carcinoma: a caseâ€“control study. <i>Bioscience Reports</i> , 2019, 39, .	1.1	9
38	Investigation of <i>methylenetetrahydrofolate reductase</i> tagging polymorphisms with colorectal cancer in Chinese Han population. <i>Oncotarget</i> , 2017, 8, 63518-63527.	0.8	9
39	N-acetyltransferase 2 Polymorphisms and Risk of Esophageal Cancer in a Chinese Population. <i>PLoS ONE</i> , 2014, 9, e87783.	1.1	8
40	<i>Methylenetetrahydrofolate reductase</i> C677T (Ala>Val, rs1801133 C>T) polymorphism decreases the susceptibility of hepatocellular carcinoma: a meta-analysis involving 12,628 subjects. <i>Bioscience Reports</i> , 2020, 40, .	1.1	8
41	Investigation of IL-4, IL-10, and HVEM polymorphisms with esophageal squamous cell carcinoma: a caseâ€“control study involving 1929 participants. <i>Bioscience Reports</i> , 2020, 40, .	1.1	8
42	Associations between CTLA-4 +49 A/G (rs231775) polymorphism and cancer risk: a meta-analysis based on 52 case-control studies. <i>International Journal of Clinical and Experimental Medicine</i> , 2015, 8, 6835-51.	1.3	8
43	<p>Leptin rs7799039 (G2548A) polymorphism is associated with cancer risk: a meta-analysis involving 25,799 subjects</p>. <i>OncoTargets and Therapy</i> , 2019, Volume 12, 2879-2890.	1.0	7
44	Investigation of <i>ICOS, CD28</i> and <i>CD80</i> polymorphisms with the risk of hepatocellular carcinoma: a caseâ€“control study in eastern Chinese population. <i>Bioscience Reports</i> , 2019, 39, .	1.1	7
45	Association between <i>microRNA-146a, -499a</i> and <i>-196a-2</i> SNPs and non-small cell lung cancer: a caseâ€“control study involving 2249 subjects. <i>Bioscience Reports</i> , 2021, 41, .	1.1	7
46	<i>TCF7L2</i> rs290481 T>C polymorphism is associated with an increased risk of type 2 diabetes mellitus and fasting plasma glucose level. <i>Oncotarget</i> , 2017, 8, 77000-77008.	0.8	7
47	Association between the CD28 IVS3 +17T>C (rs3116496) polymorphism and cancer susceptibility: a meta-analysis involving 8,843 subjects. <i>International Journal of Clinical and Experimental Medicine</i> , 2015, 8, 17353-61.	1.3	7
48	Cyclooxygenase-2 -1195G>A (rs689466) polymorphism and cancer susceptibility: an updated meta-analysis involving 50,672 subjects. <i>International Journal of Clinical and Experimental Medicine</i> , 2015, 8, 12448-62.	1.3	6
49	Peroxisome proliferator-activated receptor gamma rs1801282 C>G polymorphism is associated with polycystic ovary syndrome susceptibility: a meta-analysis involving 7,069 subjects. <i>International Journal of Clinical and Experimental Medicine</i> , 2015, 8, 17418-29.	1.3	6
50	Genetic variations in MTHFR and gastric cardia adenocarcinoma susceptibility in the Chinese Han population. <i>International Journal of Clinical and Experimental Medicine</i> , 2015, 8, 18936-44.	1.3	6
51	Forkhead box A1 (<i>FOXA1</i>) tagging polymorphisms and esophageal cancer risk in a Chinese population: a fine-mapping study. <i>Biomarkers</i> , 2016, 21, 523-529.	0.9	5
52	Association between <i>Paraoxonase</i> 1 polymorphisms and risk of esophagogastric junction adenocarcinoma: a case-control study involving 2,740 subjects. <i>Oncotarget</i> , 2017, 8, 101095-101102.	0.8	5
53	Association of metabolismâ€“related genes polymorphisms with adenocarcinoma of the oesophagogastric junction: Evidence from 2261 subjects. <i>Journal of Cellular Biochemistry</i> , 2019, 120, 18689-18701.	1.2	5
54	Association between apurinic/apyrimidinic endonuclease 1 rs1760944 T>G polymorphism and susceptibility of cancer: a meta-analysis involving 21764 subjects. <i>Bioscience Reports</i> , 2019, 39, .	1.1	5

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55	Polymorphisms of VDR gene and risk of gastric cardiac adenocarcinoma in Chinese population. <i>Oncotarget</i> , 2017, 8, 45531-45543.	0.8	5
56	S100A14 rs11548103 G>A polymorphism is associated with a decreased risk of esophageal cancer in a Chinese population. <i>Oncotarget</i> , 2017, 8, 86917-86923.	0.8	5
57	Methylenetetrahydrofolate reductase tagging polymorphisms are associated with risk of esophagogastric junction adenocarcinoma: a case-control study involving 2,740 Chinese Han subjects. <i>Oncotarget</i> , 2017, 8, 111482-111494.	0.8	5
58	Association of BTLA Polymorphisms with Susceptibility to Non-Small-Cell Lung Cancer in the Chinese Population. <i>BioMed Research International</i> , 2021, 2021, 1-10.	0.9	4
59	The correlation of microRNA-499 rs3746444 T>C locus with the susceptibility of gastric cancer: from a case"control study to a meta-analysis. <i>Bioscience Reports</i> , 2021, 41, .	1.1	4
60	Relationship of microRNA locus with type 2 diabetes mellitus: a case"control study. <i>Endocrine Connections</i> , 2021, 10, 1393-1402.	0.8	4
61	Lack of association between cyclin D1 A870G (rs9344) polymorphism and esophageal squamous cell carcinoma risk: case-control study and meta-analysis. <i>International Journal of Clinical and Experimental Medicine</i> , 2015, 8, 12685-95.	1.3	4
62	IL-15 receptor alpha rs2228059 A>C polymorphism was associated with a decreased risk of esophageal cancer in a Chinese population. <i>Molecular Biology Reports</i> , 2014, 41, 1951-1957.	1.0	3
63	<i>PADI4</i> rs2240337 G>A polymorphism is associated with susceptibility of esophageal squamous cell carcinoma in a Chinese population. <i>Oncotarget</i> , 2017, 8, 93655-93671.	0.8	3
64	MiRNA-146a rs2910164 Confers a Susceptibility to Digestive System Cancer: A Meta-Analysis Involving 59,098 Subjects. <i>Immunological Investigations</i> , 2020, , 1-20.	1.0	3
65	Lack of association between interleukin-2 (IL-2) gene rs2069762 polymorphism and cancer risk: a meta-analysis. <i>International Journal of Clinical and Experimental Medicine</i> , 2015, 8, 12557-65.	1.3	3
66	Lack of Association Between <i>PDCD-1</i> Polymorphisms and Colorectal Cancer Risk: A Case-Control Study. <i>Immunological Investigations</i> , 2022, 51, 1867-1882.	1.0	3
67	Association between <i>miR’a</i> rs2910164, <i>miRÄa2</i> rs11614913, and <i>miRǳ</i> rs3746444 polymorphisms and the risk of esophageal carcinoma: A case"control study. <i>Cancer Medicine</i> , 2022, 11, 3949-3959.	1.3	3
68	Lack of association between miR-146a rs2910164 C/G locus and colorectal cancer: from a case"control study to a meta-analysis. <i>Bioscience Reports</i> , 2021, 41, .	1.1	2
69	Evaluation of ICAM-1 rs5498 and rs3093030 Polymorphisms in Chinese Patients with Colorectal Cancer. <i>DNA and Cell Biology</i> , 2021, 40, 384-392.	0.9	2
70	Variant TP53BP1 rs560191 G>C is associated with risk of gastric cardia adenocarcinoma in a Chinese Han population. <i>Chinese Journal of Cancer Research: Official Journal of China Anti-Cancer Association</i> , Beijing Institute for Cancer Research, 2015, 27, 156-62.	0.7	2
71	IGFBP3 polymorphisms and risk of esophageal cancer in a Chinese population. <i>International Journal of Clinical and Experimental Medicine</i> , 2015, 8, 17006-14.	1.3	2
72	Association of Long NoncodingRNAsPolymorphisms with the Risk of Esophagogastric Junction Adenocarcinoma: A Three-Center Study of 1063 Cases and 1677 Controls. <i>DNA and Cell Biology</i> , 2020, 39, 828-835.	0.9	1

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73	Associations of interleukin-4 and interleukin-4 receptor loci with esophageal squamous cell carcinoma susceptibility. <i>International Immunopharmacology</i> , 2021, 97, 107659.	1.7	1
74	IL18 rs360719 A>G, IL18R1 rs13015714 G>T, IL18RAP rs917997 C>T and IL28B rs8099917 T>G polymorphisms and risk of gastric cardiac adenocarcinoma. <i>Molecular and Clinical Oncology</i> , 2017, 7, 1101-1106.	0.4	0
75	<p>The Relationship Between Single Nucleotide Polymorphisms of SMAD3/SMAD6 and Risk of Esophageal Squamous Cell Carcinoma in Chinese Population</p>. <i>Pharmacogenomics and Personalized Medicine</i> , 2020, Volume 13, 355-363.	0.4	0
76	Insulin receptor substrate-1 (IRS-1) rs1801278G>A polymorphism is associated with polycystic ovary syndrome susceptibility: a meta-analysis. <i>International Journal of Clinical and Experimental Medicine</i> , 2015, 8, 17451-60.	1.3	0
77	Association between polymorphisms and risk of colorectal cancer: a case-control study involving 2,606 subjects. <i>International Journal of Clinical and Experimental Pathology</i> , 2018, 11, 2822-2830.	0.5	0