

Hongwei Tang

List of Publications by Year in descending order

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Version: 2024-02-01

18
papers

484
citations

687363

13
h-index

839539

18
g-index

19
all docs

19
docs citations

19
times ranked

1247
citing authors

#	ARTICLE	IF	CITATIONS
1	Body Mass Index and Obesity- and Diabetes-Associated Genotypes and Risk for Pancreatic Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 779-792.	2.5	79
2	Insights into Pancreatic Cancer Etiology from Pathway Analysis of Genome-Wide Association Study Data. <i>PLoS ONE</i> , 2012, 7, e46887.	2.5	68
3	Axonal guidance signaling pathway interacting with smoking in modifying the risk of pancreatic cancer: a gene- and pathway-based interaction analysis of GWAS data. <i>Carcinogenesis</i> , 2014, 35, 1039-1045.	2.8	41
4	Effect of Diabetes Mellitus on Survival in Patients with Pancreatic Cancer: A Systematic Review and Meta-analysis. <i>Scientific Reports</i> , 2015, 5, 17102.	3.3	36
5	Antioxidant genes, diabetes and dietary antioxidants in association with risk of pancreatic cancer. <i>Carcinogenesis</i> , 2010, 31, 607-613.	2.8	35
6	Glucose metabolism gene polymorphisms and clinical outcome in pancreatic cancer. <i>Cancer</i> , 2011, 117, 480-491.	4.1	32
7	Genesâ€“Environment Interactions in Obesity- and Diabetes-Associated Pancreatic Cancer: A GWAS Data Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 98-106.	2.5	32
8	Glucose Metabolism Gene Variants Modulate the Risk of Pancreatic Cancer. <i>Cancer Prevention Research</i> , 2011, 4, 758-766.	1.5	25
9	Dietary N-nitroso compounds and risk of pancreatic cancer: results from a large caseâ€“control study. <i>Carcinogenesis</i> , 2019, 40, 254-262.	2.8	25
10	Genetic polymorphisms associated with pancreatic cancer survival: a genomeâ€“wide association study. <i>International Journal of Cancer</i> , 2017, 141, 678-686.	5.1	23
11	<scp>ABO</scp> nonâ€“O type as a risk factor for thrombosis in patients with pancreatic cancer. <i>Cancer Medicine</i> , 2015, 4, 1651-1658.	2.8	18
12	Functional Logistic Regression Approach to Detecting Gene by Longitudinal Environmental Exposure Interaction in a Caseâ€“Control Study. <i>Genetic Epidemiology</i> , 2014, 38, 638-651.	1.3	16
13	A powerful and dataâ€“adaptive test for rareâ€“variantâ€“based geneâ€“environment interaction analysis. <i>Statistics in Medicine</i> , 2019, 38, 1230-1244.	1.6	15
14	Impact of Polymorphic Variations of Gemcitabine Metabolism, DNA Damage Repair, and Drug-Resistance Genes on the Effect of High-Dose Chemotherapy for Relapsed or Refractory Lymphoid Malignancies. <i>Biology of Blood and Marrow Transplantation</i> , 2016, 22, 843-849.	2.0	9
15	Vitamin C and Vitamin E Mitigate the Risk of Pancreatic Ductal Adenocarcinoma from Meat-Derived Mutagen Exposure in Adults in a Case-Control Study. <i>Journal of Nutrition</i> , 2019, 149, 1443-1450.	2.9	9
16	Genome-Wide Geneâ€“Diabetes and Geneâ€“Obesity Interaction Scan in 8,255 Cases and 11,900 Controls from PanScan and PanC4 Consortia. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1784-1791.	2.5	5
17	Genetic Variants in DNA Double-Strand Break Repair Genes and Risk of Salivary Gland Carcinoma: A Case-Control Study. <i>PLoS ONE</i> , 2015, 10, e0128753.	2.5	4
18	Incorporating multiple sets of eQTL weights into geneâ€“byâ€“environment interaction analysis identifies novel susceptibility loci for pancreatic cancer. <i>Genetic Epidemiology</i> , 2020, 44, 880-892.	1.3	0