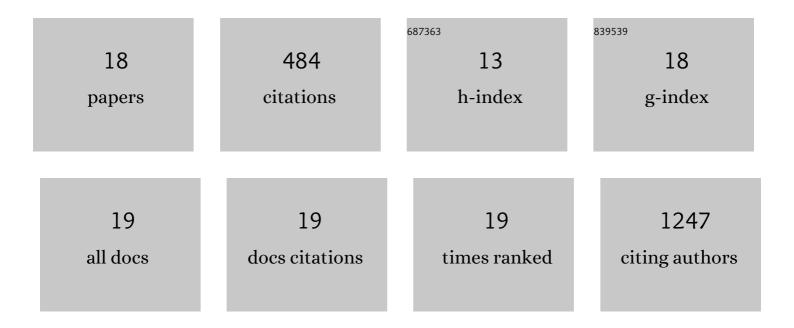
Hongwei Tang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10529475/publications.pdf Version: 2024-02-01



HONGWEI TANG

#	Article	IF	CITATIONS
1	Body Mass Index and Obesity- and Diabetes-Associated Genotypes and Risk for Pancreatic Cancer. Cancer Epidemiology Biomarkers and Prevention, 2011, 20, 779-792.	2.5	79
2	Insights into Pancreatic Cancer Etiology from Pathway Analysis of Genome-Wide Association Study Data. PLoS ONE, 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. Carcinogenesis, 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. Scientific Reports, 2015, 5, 17102.	3.3	36
5	Antioxidant genes, diabetes and dietary antioxidants in association with risk of pancreatic cancer. Carcinogenesis, 2010, 31, 607-613.	2.8	35
6	Glucose metabolism gene polymorphisms and clinical outcome in pancreatic cancer. Cancer, 2011, 117, 480-491.	4.1	32
7	Genes–Environment Interactions in Obesity- and Diabetes-Associated Pancreatic Cancer: A GWAS Data Analysis. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 98-106.	2.5	32
8	Glucose Metabolism Gene Variants Modulate the Risk of Pancreatic Cancer. Cancer Prevention Research, 2011, 4, 758-766.	1.5	25
9	Dietary N-nitroso compounds and risk of pancreatic cancer: results from a large case–control study. Carcinogenesis, 2019, 40, 254-262.	2.8	25
10	Genetic polymorphisms associated with pancreatic cancer survival: a genomeâ€wide association study. International Journal of Cancer, 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. Cancer Medicine, 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. Genetic Epidemiology, 2014, 38, 638-651.	1.3	16
13	A powerful and dataâ€adaptive test for rareâ€variant–based geneâ€environment interaction analysis. Statistics in Medicine, 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. Biology of Blood and Marrow Transplantation, 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. Journal of Nutrition, 2019, 149, 1443-1450.	2.9	9
16	Genome-Wide Gene–Diabetes and Gene–Obesity Interaction Scan in 8,255 Cases and 11,900 Controls from PanScan and PanC4 Consortia. Cancer Epidemiology Biomarkers and Prevention, 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. PLoS ONE, 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. Genetic Epidemiology, 2020, 44, 880-892.	1.3	0