

# Yaohua Yang

## List of Publications by Year in descending order

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Version: 2024-02-01

35  
papers

661  
citations

623574

14  
h-index

642610

23  
g-index

35  
all docs

35  
docs citations

35  
times ranked

1062  
citing authors

#	ARTICLE	IF	CITATIONS
1	2â€™-Fucosyllactose Ameliorates Chemotherapy-Induced Intestinal Mucositis by Protecting Intestinal Epithelial Cells Against Apoptosis. Cellular and Molecular Gastroenterology and Hepatology, 2022, 13, 441-457.	2.3	16
2	Prospective study of oral microbiome and gastric cancer risk among Asian, African American and European American populations. International Journal of Cancer, 2022, 150, 916-927.	2.3	17
3	The oral microbiome in relation to pancreatic cancer risk in African Americans. British Journal of Cancer, 2022, 126, 287-296.	2.9	9
4	Oral and gastric microbiome in relation to gastric intestinal metaplasia. International Journal of Cancer, 2022, 150, 928-940.	2.3	25
5	Association Between Long-Term Regular Exercise and Gut Microbiota Among Middle-Aged and Older Urban Chinese. International Journal of Sport Nutrition and Exercise Metabolism, 2022, , 1-9.	1.0	1
6	Incorporating Polygenic Risk Scores and Nongenetic Risk Factors for Breast Cancer Risk Prediction Among Asian Women. JAMA Network Open, 2022, 5, e2149030.	2.8	12
7	Large-scale Integrated Analysis of Genetics and Metabolomic Data Reveals Potential Links Between Lipids and Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2022, 31, 1216-1226.	1.1	3
8	Mendelian randomization analyses of 23 known and suspected risk factors and biomarkers for breast cancer overall and by molecular subtypes. International Journal of Cancer, 2022, 151, 372-380.	2.3	20
9	Tea Consumption and Gut Microbiome in Chinese Men and Women. Current Developments in Nutrition, 2022, 6, 1035.	0.1	0
10	Long-term diet quality is associated with gut microbiome diversity and composition among urban Chinese adults. American Journal of Clinical Nutrition, 2021, 113, 684-694.	2.2	42
11	Long-term Diet Quality and Gut Microbiome Functionality: A Prospective, Shotgun Metagenomic Study among Urban Chinese Adults. Current Developments in Nutrition, 2021, 5, nzab026.	0.1	13
12	Legume Consumption and Gut Microbiome in Elderly Chinese Men and Women. Journal of Nutrition, 2021, 151, 2399-2408.	1.3	7
13	Association of oral microbiota with lung cancer risk in a low-income population in the Southeastern USA. Cancer Causes and Control, 2021, 32, 1423-1432.	0.8	18
14	Novel strategy for disease risk prediction incorporating predicted gene expression and DNA methylation data: a multiâ€phased study of prostate cancer. Cancer Communications, 2021, 41, 1387-1397.	3.7	6
15	Integrating Genome and Methylome Data to Identify Candidate DNA Methylation Biomarkers for Pancreatic Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 2079-2087.	1.1	10
16	Variation in oral microbiome is associated with future risk of lung cancer among never-smokers. Thorax, 2021, 76, 256-263.	2.7	51
17	Sex-Specific Associations between Gut Microbiome and Non-Alcoholic Fatty Liver Disease among Urban Chinese Adults. Microorganisms, 2021, 9, 2118.	1.6	12
18	Genetically Predicted Levels of DNA Methylation Biomarkers and Breast Cancer Risk: Data From 228â€™951 Women of European Descent. Journal of the National Cancer Institute, 2020, 112, 295-304.	3.0	35

#	ARTICLE	IF	CITATIONS
19	Evaluation of associations between genetically predicted circulating protein biomarkers and breast cancer risk. <i>International Journal of Cancer</i> , 2020, 146, 2130-2138.	2.3	13
20	An integrative multi-omics analysis to identify candidate DNA methylation biomarkers related to prostate cancer risk. <i>Nature Communications</i> , 2020, 11, 3905.	5.8	28
21	Mendelian Randomization Analysis of n-6 Polyunsaturated Fatty Acid Levels and Pancreatic Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 2735-2739.	1.1	6
22	Reply to Kenyon, "Are Differences in the Oral Microbiome Due to Ancestry or Socioeconomics?". <i>MSystems</i> , 2020, 5, .	1.7	0
23	Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. <i>Nature Communications</i> , 2020, 11, 1217.	5.8	46
24	Abstract 28: Integrating genomic and transcriptomic data to identify genetic loci associated with breast cancer risk in women of African ancestry. , 2020, , .		0
25	Genetic Data from Nearly 63,000 Women of European Descent Predicts DNA Methylation Biomarkers and Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , 2019, 79, 505-517.	0.4	49
26	Re-evaluating genetic variants identified in candidate gene studies of breast cancer risk using data from nearly 280,000 women of Asian and European ancestry. <i>EBioMedicine</i> , 2019, 48, 203-211.	2.7	14
27	Oral microbiome and obesity in a large study of low-income and African-American populations. <i>Journal of Oral Microbiology</i> , 2019, 11, 1650597.	1.2	46
28	Cigarette smoking and oral microbiota in low-income and African-American populations. <i>Journal of Epidemiology and Community Health</i> , 2019, 73, 1108-1115.	2.0	26
29	Racial Differences in the Oral Microbiome: Data from Low-Income Populations of African Ancestry and European Ancestry. <i>MSystems</i> , 2019, 4, .	1.7	32
30	Prospective study of oral microbiome and colorectal cancer risk in low-income and African American populations. <i>International Journal of Cancer</i> , 2019, 144, 2381-2389.	2.3	81
31	Abstract 1649: Integrating genome, transcriptome and methylome data to identify novel genes for lung cancer: Data from over 50,000 European participants. , 2019, , .		0
32	Abstract 5314: DNA methylation quantitative trait loci and breast cancer risk: Data from nearly 230,000 women of European descent. , 2018, , .		2
33	Abstract 691: Associations of genetically-predicted circulating immunoglobulin traits with breast cancer risk. , 2018, , .		0
34	Abstract 4931: Prospective study of oral microbiome and colorectal cancer risk in low-income and African American populations. <i>Cancer Research</i> , 2017, 77, 4931-4931.	0.4	3
35	CyanOmics: an integrated database of omics for the model cyanobacterium <i>Synechococcus</i> sp. PCC 7002. <i>Database: the Journal of Biological Databases and Curation</i> , 2015, 2015, .	1.4	18