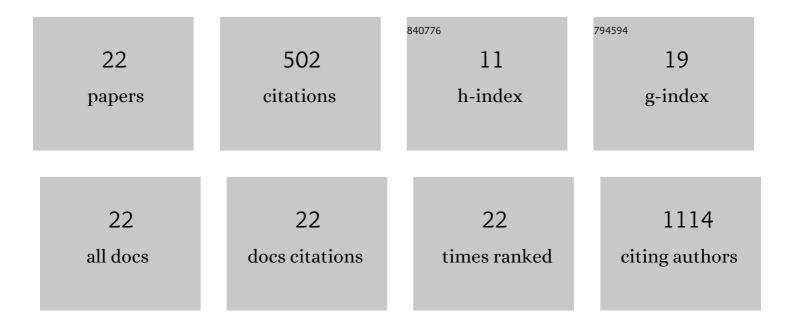
Cheng Peng

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A Metabolomics Analysis of Circulating Carotenoids and Breast Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2022, 31, 85-96.	2.5	6
2	Abstract P3-12-01: Regular aspirin use, breast tumor characteristics and long-term breast cancer survival. Cancer Research, 2022, 82, P3-12-01-P3-12-01.	0.9	0
3	A Genome-Wide Gene-Based Gene–Environment Interaction Study of Breast Cancer in More than 90,000 Women. Cancer Research Communications, 2022, 2, 211-219.	1.7	6
4	Bibliometric and visualized analysis of ocular drug delivery from 2001 to 2020. Journal of Controlled Release, 2022, 345, 625-645.	9.9	63
5	Early-Life Body Adiposity and the Breast Tumor Transcriptome. Journal of the National Cancer Institute, 2021, 113, 778-784.	6.3	9
6	Circulating carotenoids and breast cancer among high-risk individuals. American Journal of Clinical Nutrition, 2021, 113, 525-533.	4.7	13
7	Branched-Chain Amino Acids and Risk of Breast Cancer. JNCI Cancer Spectrum, 2021, 5, pkab059.	2.9	12
8	A polygenic-score-based approach for identification of gene-drug interactions stratifying breast cancer risk. American Journal of Human Genetics, 2021, 108, 1752-1764.	6.2	7
9	Vitamin A: A Potential Intervention for Breast Cancer Racial Disparities. Journal of Nutrition, 2021, 151, 3602-3603.	2.9	0
10	Prospective study of a diabetes risk reduction diet and the risk of breast cancer. American Journal of Clinical Nutrition, 2020, 112, 1492-1503.	4.7	31
11	Low dose environmental radon exposure and breast tumor gene expression. BMC Cancer, 2020, 20, 695.	2.6	5
12	Prediagnostic 25-Hydroxyvitamin D Concentrations in Relation to Tumor Molecular Alterations and Risk of Breast Cancer Recurrence. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1253-1263.	2.5	4
13	Prenatal Particulate Air Pollution and DNA Methylation in Newborns: An Epigenome-Wide Meta-Analysis. Environmental Health Perspectives, 2019, 127, 57012.	6.0	111
14	Epigenome-wide association study reveals methylation pathways associated with childhood allergic sensitization. Epigenetics, 2019, 14, 445-466.	2.7	43
15	Epigenetic age acceleration is associated with allergy and asthma in children in Project Viva. Journal of Allergy and Clinical Immunology, 2019, 143, 2263-2270.e14.	2.9	43
16	Altered cord blood mitochondrial DNA content and pregnancy lead exposure in the PROGRESS cohort. Environment International, 2019, 125, 437-444.	10.0	27
17	Psychosocial stress is associated with benign breast disease in young Chinese women: results from Project ELEFANT. Breast Cancer Research and Treatment, 2019, 173, 217-224.	2.5	8
18	Epigenome-wide association study of total serum immunoglobulin E in children: a life course approach. Clinical Epigenetics, 2018, 10, 55.	4.1	36

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#	Article	IF	CITATIONS
19	Residential Proximity to Major Roadways at Birth, DNA Methylation at Birth and Midchildhood, and Childhood Cognitive Test Scores: Project Viva(Massachusetts, USA). Environmental Health Perspectives, 2018, 126, 97006.	6.0	15
20	Associations of Annual Ambient Fine Particulate Matter Mass and Components with Mitochondrial DNA Abundance. Epidemiology, 2017, 28, 763-770.	2.7	18
21	Lack of FGF-7 Further Delays Cutaneous Wound Healing in Diabetic Mice. Plastic and Reconstructive Surgery, 2011, 128, 673e-684e.	1.4	44
22	A multi-state survival model for time to breast cancer mortality among a cohort of initially disease-free women. Cancer Epidemiology Biomarkers and Prevention, 0, , .	2.5	1