Tuong Linh Nguyen

List of Publications by Year
in descending order

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Version: 2024-02-01

35 papers	663 citations	567281 15 h-index	610901 24 g-index
35 all docs	35 docs citations	35 times ranked	1038 citing authors

#	Article	IF	Citations
1	Blood DNA methylation score predicts breast cancer risk: applying OPERA in molecular, environmental, genetic and analytic epidemiology. Molecular Oncology, 2022, 16, 8-10.	4.6	3
2	Familial Aspects of Mammographic Density Measures Associated with Breast Cancer Risk. Cancers, 2022, 14, 1483.	3.7	6
3	Early life affects late-life health through determining DNA methylation across the lifespan: A twin study. EBioMedicine, 2022, 77, 103927.	6.1	15
4	Association of contralateral breast cancer risk with mammographic density defined at higherâ€thanâ€conventional intensity thresholds. International Journal of Cancer, 2022, 151, 1304-1309.	5.1	3
5	Weight is More Informative than Body Mass Index for Predicting Postmenopausal Breast Cancer Risk: Prospective Family Study Cohort (ProF-SC). Cancer Prevention Research, 2022, 15, 185-191.	1.5	4
6	Genome-wide and transcriptome-wide association studies of mammographic density phenotypes reveal novel loci. Breast Cancer Research, 2022, 24, 27.	5.0	15
7	Genetic Aspects of Mammographic Density Measures Associated with Breast Cancer Risk. Cancers, 2022, 14, 2767.	3.7	5
8	Novel mammogramâ€based measures improve breast cancer risk prediction beyond an established mammographic density measure. International Journal of Cancer, 2021, 148, 2193-2202.	5.1	18
9	RE: Chemopreventive Agents to Reduce Mammographic Breast Density in Premenopausal Women: A Systematic Review of Clinical Trials. JNCI Cancer Spectrum, 2021, 5, pkab051.	2.9	1
10	Towards riskâ€stratified population breast cancer screening: more than mammographic density. Medical Journal of Australia, 2021, 215, 350-351.	1.7	2
11	Mammographic texture features associated with contralateral breast cancer in the WECARE Study. Npj Breast Cancer, 2021, 7, 146.	5.2	1
12	Interval breast cancer risk associations with breast density, family history and breast tissue aging. International Journal of Cancer, 2020, 147, 375-382.	5.1	22
13	Prognostic value of metabolic tumor volume and total lesion glycolysis in breast cancer: a meta-analysis. Nuclear Medicine Communications, 2020, 41, 824-829.	1.1	9
14	Genetic and environmental causes of variation in epigenetic aging across the lifespan. Clinical Epigenetics, 2020, 12, 158.	4.1	33
15	Going Beyond Conventional Mammographic Density to Discover Novel Mammogram-Based Predictors of Breast Cancer Risk. Journal of Clinical Medicine, 2020, 9, 627.	2.4	23
16	Inference about causation between body mass index and DNA methylation in blood from a twin family study. International Journal of Obesity, 2019, 43, 243-252.	3.4	48
17	DNA methylation-based biological age, genome-wide average DNA methylation, and conventional breast cancer risk factors. Scientific Reports, 2019, 9, 15055.	3.3	18
18	Genomeâ€wide association study of peripheral blood DNA methylation and conventional mammographic density measures. International Journal of Cancer, 2019, 145, 1768-1773.	5.1	17

#	Article	IF	Citations
19	Measurement challenge: protocol for international caseâ€“control comparison of mammographic measures that predict breast cancer risk. BMJ Open, 2019, 9, e031041.	1.9	14
20	Breast Cancer Risk Associations with Digital Mammographic Density by Pixel Brightness Threshold and Mammographic System. Radiology, 2018, 286, 433-442.	7.3	29
21	Association between mammographic density and tumor marker-defined breast cancer subtypes: a caseâ€“control study. European Journal of Cancer Prevention, 2018, 27, 239-247.	1.3	13
22	Cirrus: An Automated Mammography-Based Measure of Breast Cancer Risk Based on Textural Features. JNCI Cancer Spectrum, 2018, 2, pky057.	2.9	24
23	Predicting interval and screen-detected breast cancers from mammographic density defined by different brightness thresholds. Breast Cancer Research, 2018, 20, 152.	5.0	24
24	Genome-wide average DNA methylation is determined in utero. International Journal of Epidemiology, 2018, 47, 908-916.	1.9	38
25	Causal effect of smoking on DNA methylation in peripheral blood: a twin and family study. Clinical Epigenetics, 2018, 10, 18.	4.1	95
26	Mammographic Density and Circulating Sex Hormones: a Cross-Sectional Study in Postmenopausal Korean Women. Hormones and Cancer, 2018, 9, 383-390.	4.9	2
27	Mammographic density defined by higher than conventional brightness thresholds better predicts breast cancer risk. International Journal of Epidemiology, 2017, 46, dyw212.	1.9	24
28	Twin birth changes DNA methylation of subsequent siblings. Scientific Reports, 2017, 7, 8463.	3.3	8
29	Causes of blood methylomic variation for middle-aged women measured by the HumanMethylation450 array. Epigenetics, 2017, 12, 973-981.	2.7	14
30	Comparison of the association of mammographic density and clinical factors with ductal carcinoma in situ versus invasive ductal breast cancer in Korean women. BMC Cancer, 2017, 17, 821.	2.6	5
31	Childhood body mass index and adult mammographic density measures that predict breast cancer risk. Breast Cancer Research and Treatment, 2016, 156, 163-170.	2.5	19
32	Mammographic density defined by higher than conventional brightness threshold better predicts breast cancer risk for full-field digital mammograms. Breast Cancer Research, 2015, 17, 142.	5.0	35
33	Mammographic density and risk of breast cancer in Korean women. European Journal of Cancer Prevention, 2015, 24, 422-429.	1.3	24
34	Bone mineral density and the risk of breast cancer: a case-control study of Korean women. Annals of Epidemiology, 2014, 24, 222-227.	1.9	16
35	Explaining Variance in the Cumulus Mammographic Measures That Predict Breast Cancer Risk: A Twins and Sisters Study. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 2395-2403.	2.5	36