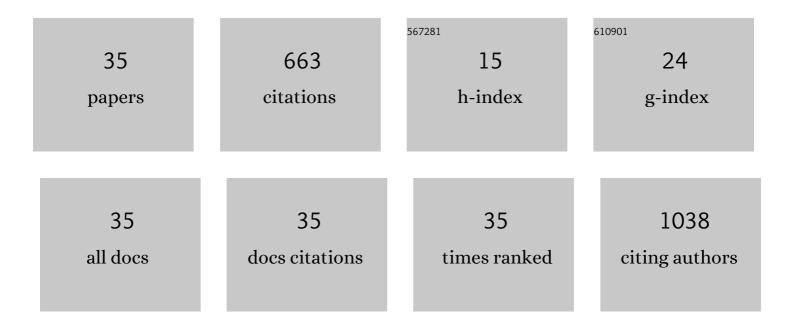
## Tuong Linh Nguyen

List of Publications by Year in descending order

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TUONG LINH NOUVEN

#	Article	IF	CITATIONS
1	Causal effect of smoking on DNA methylation in peripheral blood: a twin and family study. Clinical Epigenetics, 2018, 10, 18.	4.1	95
2	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
3	Genome-wide average DNA methylation is determined in utero. International Journal of Epidemiology, 2018, 47, 908-916.	1.9	38
4	Explaining Variance in the <i>Cumulus</i> Mammographic Measures That Predict Breast Cancer Risk: A Twins and Sisters Study. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 2395-2403.	2.5	36
5	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
6	Genetic and environmental causes of variation in epigenetic aging across the lifespan. Clinical Epigenetics, 2020, 12, 158.	4.1	33
7	Breast Cancer Risk Associations with Digital Mammographic Density by Pixel Brightness Threshold and Mammographic System. Radiology, 2018, 286, 433-442.	7.3	29
8	Mammographic density and risk of breast cancer in Korean women. European Journal of Cancer Prevention, 2015, 24, 422-429.	1.3	24
9	Mammographic density defined by higher than conventional brightness thresholds better predicts breast cancer risk. International Journal of Epidemiology, 2017, 46, dyw212.	1.9	24
10	Cirrus: An Automated Mammography-Based Measure of Breast Cancer Risk Based on Textural Features. JNCI Cancer Spectrum, 2018, 2, pky057.	2.9	24
11	Predicting interval and screen-detected breast cancers from mammographic density defined by different brightness thresholds. Breast Cancer Research, 2018, 20, 152.	5.0	24
12	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
13	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
14	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
15	DNA methylation-based biological age, genome-wide average DNA methylation, and conventional breast cancer risk factors. Scientific Reports, 2019, 9, 15055.	3.3	18
16	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
17	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
18	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

TUONG LINH NGUYEN

#	Article	IF	CITATIONS
19	Early life affects late-life health through determining DNA methylation across the lifespan: A twin study. EBioMedicine, 2022, 77, 103927.	6.1	15
20	Genome-wide and transcriptome-wide association studies of mammographic density phenotypes reveal novel loci. Breast Cancer Research, 2022, 24, 27.	5.0	15
21	Causes of blood methylomic variation for middle-aged women measured by the HumanMethylation450 array. Epigenetics, 2017, 12, 973-981.	2.7	14
22	Measurement challenge: protocol for international case–control comparison of mammographic measures that predict breast cancer risk. BMJ Open, 2019, 9, e031041.	1.9	14
23	Association between mammographic density and tumor marker-defined breast cancer subtypes: a case–control study. European Journal of Cancer Prevention, 2018, 27, 239-247.	1.3	13
24	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
25	Twin birth changes DNA methylation of subsequent siblings. Scientific Reports, 2017, 7, 8463.	3.3	8
26	Familial Aspects of Mammographic Density Measures Associated with Breast Cancer Risk. Cancers, 2022, 14, 1483.	3.7	6
27	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
28	Genetic Aspects of Mammographic Density Measures Associated with Breast Cancer Risk. Cancers, 2022, 14, 2767.	3.7	5
29	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
30	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
31	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
32	Mammographic Density and Circulating Sex Hormones: a Cross-Sectional Study in Postmenopausal Korean Women. Hormones and Cancer, 2018, 9, 383-390.	4.9	2
33	Towards riskâ€ <b>s</b> tratified population breast cancer screening: more than mammographic density. Medical Journal of Australia, 2021, 215, 350-351.	1.7	2
34	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
35	Mammographic texture features associated with contralateral breast cancer in the WECARE Study. Npj Breast Cancer, 2021, 7, 146.	5.2	1