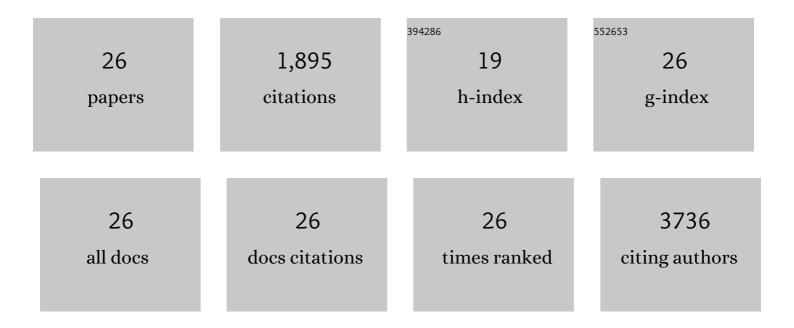
Min-Ae Song

List of Publications by Year in descending order

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MIN-AF SONG

#	Article	IF	CITATIONS
1	Saliva and Lung Microbiome Associations with Electronic Cigarette Use and Smoking. Cancer Prevention Research, 2022, 15, 435-446.	0.7	6
2	Race/ethnicity-associated blood DNA methylation differences between Japanese and European American women: an exploratory study. Clinical Epigenetics, 2021, 13, 188.	1.8	10
3	Effects of Electronic Cigarette Constituents on the Human Lung: A Pilot Clinical Trial. Cancer Prevention Research, 2020, 13, 145-152.	0.7	60
4	Biomarkers of Exposure and Effect in the Lungs of Smokers, Nonsmokers, and Electronic Cigarette Users. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 443-451.	1.1	43
5	Vitamin E Acetate in Bronchoalveolar-Lavage Fluid Associated with EVALI. New England Journal of Medicine, 2020, 382, 697-705.	13.9	510
6	Use of an Exposome Approach to Understand the Effects of Exposures From the Natural, Built, and Social Environments on Cardio-Vascular Disease Onset, Progression, and Outcomes. Frontiers in Public Health, 2020, 8, 379.	1.3	42
7	SOX9 promotes stress-responsive transcription of VGF nerve growth factor inducible gene in renal tubular epithelial cells. Journal of Biological Chemistry, 2020, 295, 16328-16341.	1.6	20
8	Lipid laden macrophages and electronic cigarettes in healthy adults. EBioMedicine, 2020, 60, 102982.	2.7	28
9	Inflammasome Adaptor ASC Is Highly Elevated in Lung Over Plasma and Relates to Inflammation and Lung Diffusion in the Absence of Speck Formation. Frontiers in Immunology, 2020, 11, 461.	2.2	10
10	DNA Methylation and Smoking: Implications for Understanding Effects of Electronic Cigarettes. Current Epidemiology Reports, 2019, 6, 148-161.	1.1	2
11	Electronic versus Combustible Cigarette Effects on Inflammasome Component Release into Human Lung. American Journal of Respiratory and Critical Care Medicine, 2019, 199, 922-925.	2.5	28
12	FoxO1 is a critical regulator of M2â€like macrophage activation in allergic asthma. Allergy: European Journal of Allergy and Clinical Immunology, 2019, 74, 535-548.	2.7	29
13	Methylation of imprinted <i>IGF2</i> regions is associated with total, visceral, and hepatic adiposity in postmenopausal women. Epigenetics, 2018, 13, 858-865.	1.3	6
14	DNA Methylation as a Biomarker of Aging in Epidemiologic Studies. Methods in Molecular Biology, 2018, 1856, 219-231.	0.4	21
15	Cigarette Filter Ventilation and its Relationship to Increasing Rates of Lung Adenocarcinoma. Journal of the National Cancer Institute, 2017, 109, .	3.0	79
16	A Review of Pulmonary Toxicity of Electronic Cigarettes in the Context of Smoking: A Focus on Inflammation. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 1175-1191.	1.1	95
17	Landscape of genome-wide age-related DNA methylation in breast tissue. Oncotarget, 2017, 8, 114648-114662.	0.8	27
18	Chemical and toxicological characteristics of conventional and low-TSNA moist snuff tobacco products. Toxicology Letters, 2016, 245, 68-77.	0.4	34

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#	Article	IF	CITATIONS
19	Genetic variation in one-carbon metabolism in relation to genome-wide DNA methylation in breast tissue from healthy women. Carcinogenesis, 2016, 37, 471-480.	1.3	22
20	Racial differences in genome-wide methylation profiling and gene expression in breast tissues from healthy women. Epigenetics, 2015, 10, 1177-1187.	1.3	49
21	Intraindividual Variation and Short-term Temporal Trend in DNA Methylation of Human Blood. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 490-497.	1.1	27
22	Genome-scale hypomethylation in the cord blood DNAs associated with early onset preeclampsia. Clinical Epigenetics, 2015, 7, 21.	1.8	41
23	Genome-wide hypermethylation coupled with promoter hypomethylation in the chorioamniotic membranes of early onset pre-eclampsia. Molecular Human Reproduction, 2014, 20, 885-904.	1.3	54
24	<i>CCAT2</i> , a novel noncoding RNA mapping to 8q24, underlies metastatic progression and chromosomal instability in colon cancer. Genome Research, 2013, 23, 1446-1461.	2.4	526
25	Elucidating the Landscape of Aberrant DNA Methylation in Hepatocellular Carcinoma. PLoS ONE, 2013, 8, e55761.	1.1	101
26	Quantification of CpG methylation at the 5′-region of XIST by pyrosequencing from human serum. Electrophoresis, 2007, 28, 2379-2384.	1.3	25