## Nabila Kazmi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2636664/publications.pdf

Version: 2024-02-01

759233 940533 1,031 22 12 16 h-index citations g-index papers 22 22 22 2361 docs citations all docs times ranked citing authors

#	Article	IF	CITATIONS
1	Genomic and phenotypic insights from an atlas of genetic effects on DNA methylation. Nature Genetics, 2021, 53, 1311-1321.	21.4	218
2	Physical activity and risks of breast and colorectal cancer: a Mendelian randomisation analysis. Nature Communications, 2020, 11, 597.	12.8	193
3	Epigenome-wide meta-analysis of DNA methylation and childhood asthma. Journal of Allergy and Clinical Immunology, 2019, 143, 2062-2074.	2.9	147
4	DNA methylation changes measured in preâ€diagnostic peripheral blood samples are associated with smoking and lung cancer risk. International Journal of Cancer, 2017, 140, 50-61.	5.1	115
5	Epigenome-wide meta-analysis of blood DNA methylation in newborns and children identifies numerous loci related to gestational age. Genome Medicine, 2020, 12, 25.	8.2	81
6	Hypertensive Disorders of Pregnancy and DNA Methylation in Newborns. Hypertension, 2019, 74, 375-383.	2.7	73
7	The long-term impact of folic acid in pregnancy on offspring DNA methylation: follow-up of the Aberdeen Folic Acid Supplementation Trial (AFAST). International Journal of Epidemiology, 2018, 47, 928-937.	1.9	56
8	Associations between high blood pressure and DNA methylation. PLoS ONE, 2020, 15, e0227728.	2.5	37
9	Appraising causal relationships of dietary, nutritional and physical-activity exposures with overall and aggressive prostate cancer: two-sample Mendelian-randomization study based on 79 148 prostate-cancer cases and 61 106 controls. International Journal of Epidemiology, 2020, 49, 587-596.	1.9	36
10	DNA methylation derived systemic inflammation indices are associated with head and neck cancer development and survival. Oral Oncology, 2018, 85, 87-94.	1.5	17
11	Maternal eating disorders affect offspring cord blood DNA methylation: a prospective study. Clinical Epigenetics, 2017, 9, 120.	4.1	15
12	Genetic determinants of circulating haptoglobin concentration. Clinica Chimica Acta, 2019, 494, 138-142.	1.1	14
13	Haptoglobin genotype and outcome after aneurysmal subarachnoid haemorrhage. Journal of Neurology, Neurosurgery and Psychiatry, 2020, 91, 305-313.	1.9	11
14	Diagnosis of Coronary Heart Diseases Using Gene Expression Profiling; Stable Coronary Artery Disease, Cardiac Ischemia with and without Myocardial Necrosis. PLoS ONE, 2016, 11, e0149475.	2.5	10
15	Haptoglobin genotype and outcome after spontaneous intracerebral haemorrhage. Journal of Neurology, Neurosurgery and Psychiatry, 2020, 91, 298-304.	1.9	4
16	Rho GTPase gene expression and breast cancer risk: a Mendelian randomization analysis. Scientific Reports, 2022, 12, 1463.	3.3	4
17	Associations between high blood pressure and DNA methylation. , 2020, 15, e0227728.		O
18	Associations between high blood pressure and DNA methylation. , 2020, 15, e0227728.		0

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19	Associations between high blood pressure and DNA methylation. , 2020, 15, e0227728.		0
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