

Ghazaleh Fatemifar

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3506926/publications.pdf>

Version: 2024-02-01

16
papers

2,713
citations

623734

14
h-index

940533

16
g-index

21
all docs

21
docs citations

21
times ranked

7157
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association and Mendelian randomisation analysis provide insights into the pathogenesis of heart failure. <i>Nature Communications</i> , 2020, 11, 163.	12.8	466
2	UK phenomics platform for developing and validating electronic health record phenotypes: CALIBER. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2019, 26, 1545-1559.	4.4	143
3	Phenome-wide association analysis of LDL-cholesterol lowering genetic variants in PCSK9. <i>BMC Cardiovascular Disorders</i> , 2019, 19, 240.	1.7	22
4	Associations Between Measures of Sarcopenic Obesity and Risk of Cardiovascular Disease and Mortality: A Cohort Study and Mendelian Randomization Analysis Using the UK Biobank. <i>Journal of the American Heart Association</i> , 2019, 8, e011638.	3.7	75
5	The relationship between sleep duration, cognition and dementia: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2019, 48, 849-860.	1.9	83
6	Assessing potential shared genetic aetiology between body mass index and sleep duration in 142,209 individuals. <i>Genetic Epidemiology</i> , 2019, 43, 207-214.	1.3	7
7	Causal Associations of Adiposity and Body Fat Distribution With Coronary Heart Disease, Stroke Subtypes, and Type 2 Diabetes Mellitus. <i>Circulation</i> , 2017, 135, 2373-2388.	1.6	304
8	Genome-wide analysis of health-related biomarkers in the UK Household Longitudinal Study reveals novel associations. <i>Scientific Reports</i> , 2017, 7, 11008.	3.3	88
9	Genetic variants associated with subjective well-being, depressive symptoms, and neuroticism identified through genome-wide analyses. <i>Nature Genetics</i> , 2016, 48, 624-633.	21.4	870
10	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. <i>Nature Communications</i> , 2016, 7, 10495.	12.8	245
11	Metabolomic Profiling of Statin Use and Genetic Inhibition of HMG-CoA Reductase. <i>Journal of the American College of Cardiology</i> , 2016, 67, 1200-1210.	2.8	173
12	The Association between Primary Tooth Emergence and Anthropometric Measures in Young Adults: Findings from a Large Prospective Cohort Study. <i>PLoS ONE</i> , 2014, 9, e96355.	2.5	9
13	Genome-wide association study of primary tooth eruption identifies pleiotropic loci associated with height and craniofacial distances. <i>Human Molecular Genetics</i> , 2013, 22, 3807-3817.	2.9	84
14	Postnatal Growth and DNA Methylation Are Associated With Differential Gene Expression of the TACSTD2 Gene and Childhood Fat Mass. <i>Diabetes</i> , 2012, 61, 391-400.	0.6	55
15	Genome-Wide Association Study Identifies Four Loci Associated with Eruption of Permanent Teeth. <i>PLoS Genetics</i> , 2011, 7, e1002275.	3.5	42
16	Evaluation of the Association of <i>IGF2BP2</i> Variants With Type 2 Diabetes in French Caucasians. <i>Diabetes</i> , 2008, 57, 1992-1996.	0.6	26