Meena Kumari

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10886726/publications.pdf

Version: 2024-02-01

226 papers 46,339 citations

89 h-index 201 g-index

243 all docs $\begin{array}{c} 243 \\ \text{docs citations} \end{array}$

times ranked

243

50705 citing authors

#	Article	IF	CITATIONS
1	Genetic studies of body mass index yield new insights for obesity biology. Nature, 2015, 518, 197-206.	13.7	3,823
2	Discovery and refinement of loci associated with lipid levels. Nature Genetics, 2013, 45, 1274-1283.	9.4	2,641
3	New genetic loci implicated in fasting glucose homeostasis and their impact on type 2 diabetes risk. Nature Genetics, 2010, 42, 105-116.	9.4	1,982
4	Genetic variants in novel pathways influence blood pressure and cardiovascular disease risk. Nature, 2011, 478, 103-109.	13.7	1,855
5	Gene discovery and polygenic prediction from a genome-wide association study of educational attainment in 1.1 million individuals. Nature Genetics, 2018, 50, 1112-1121.	9.4	1,835
6	Defining the role of common variation in the genomic and biological architecture of adult human height. Nature Genetics, 2014, 46, 1173-1186.	9.4	1,818
7	Inflammation, obesity, stress and coronary heart disease: is interleukin-6 the link?. Atherosclerosis, 2000, 148, 209-214.	0.4	1,611
8	New genetic loci link adipose and insulin biology to body fat distribution. Nature, 2015, 518, 187-196.	13.7	1,328
9	The interleukin-6 receptor as a target for prevention of coronary heart disease: a mendelian randomisation analysis. Lancet, The, 2012, 379, 1214-1224.	6. 3	886
10	Genetic variants associated with subjective well-being, depressive symptoms, and neuroticism identified through genome-wide analyses. Nature Genetics, 2016, 48, 624-633.	9.4	870
11	Job strain as a risk factor for coronary heart disease: a collaborative meta-analysis of individual participant data. Lancet, The, 2012, 380, 1491-1497.	6.3	786
12	A genome-wide approach accounting for body mass index identifies genetic variants influencing fasting glycemic traits and insulin resistance. Nature Genetics, 2012, 44, 659-669.	9.4	762
13	Common variants associated with plasma triglycerides and risk for coronary artery disease. Nature Genetics, 2013, 45, 1345-1352.	9.4	754
14	Large-scale association analyses identify new loci influencing glycemic traits and provide insight into the underlying biological pathways. Nature Genetics, 2012, 44, 991-1005.	9.4	746
15	Assessing salivary cortisol in large-scale, epidemiological research. Psychoneuroendocrinology, 2009, 34, 1423-1436.	1.3	694
16	Genetic variation in GIPR influences the glucose and insulin responses to an oral glucose challenge. Nature Genetics, 2010, 42, 142-148.	9.4	591
17	Genome-wide meta-analysis identifies 11 new loci for anthropometric traits and provides insights into genetic architecture. Nature Genetics, 2013, 45, 501-512.	9.4	578
18	Mendelian randomization of blood lipids for coronary heart disease. European Heart Journal, 2015, 36, 539-550.	1.0	567

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19	HMG-coenzyme A reductase inhibition, type 2 diabetes, and bodyweight: evidence from genetic analysis and randomised trials. Lancet, The, 2015, 385, 351-361.	6.3	562
20	Genome-wide association analyses of risk tolerance and risky behaviors in over 1 million individuals identify hundreds of loci and shared genetic influences. Nature Genetics, 2019, 51, 245-257.	9.4	536
21	Long working hours and risk of coronary heart disease and stroke: a systematic review and meta-analysis of published and unpublished data for 603â€^838 individuals. Lancet, The, 2015, 386, 1739-1746.	6.3	529
22	Association between alcohol and cardiovascular disease: Mendelian randomisation analysis based on individual participant data. BMJ, The, 2014, 349, g4164-g4164.	3.0	528
23	Work stress and coronary heart disease: what are the mechanisms?. European Heart Journal, 2008, 29, 640-648.	1.0	507
24	Use of low-density lipoprotein cholesterol gene score to distinguish patients with polygenic and monogenic familial hypercholesterolaemia: a case-control study. Lancet, The, 2013, 381, 1293-1301.	6.3	485
25	Physical Activity Attenuates the Influence of FTO Variants on Obesity Risk: A Meta-Analysis of 218,166 Adults and 19,268 Children. PLoS Medicine, 2011, 8, e1001116.	3.9	446
26	A Prospective Study of Change in Sleep Duration: Associations with Mortality in the Whitehall II Cohort. Sleep, 2007, 30, 1659-1666.	0.6	440
27	Rare variant in scavenger receptor BI raises HDL cholesterol and increases risk of coronary heart disease. Science, 2016, 351, 1166-1171.	6.0	438
28	Gender-Specific Associations of Short Sleep Duration With Prevalent and Incident Hypertension. Hypertension, 2007, 50, 693-700.	1.3	430
29	Sex-stratified Genome-wide Association Studies Including 270,000 Individuals Show Sexual Dimorphism in Genetic Loci for Anthropometric Traits. PLoS Genetics, 2013, 9, e1003500.	1.5	371
30	The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. Nature Genetics, 2016, 48, 1171-1184.	9.4	362
31	The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679.	13.7	353
32	Impact of common genetic determinants of Hemoglobin A1c on type 2 diabetes risk and diagnosis in ancestrally diverse populations: A transethnic genome-wide meta-analysis. PLoS Medicine, 2017, 14, e1002383.	3.9	341
33	The trans-ancestral genomic architecture of glycemic traits. Nature Genetics, 2021, 53, 840-860.	9.4	341
34	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. PLoS Genetics, 2015, 11, e1005378.	1.5	331
35	Association of vitamin D status with arterial blood pressure and hypertension risk: a mendelian randomisation study. Lancet Diabetes and Endocrinology,the, 2014, 2, 719-729.	5.5	319
36	Prospective Study of Social and Other Risk Factors for Incidence of Type 2 Diabetes in the Whitehall II Study. Archives of Internal Medicine, 2004, 164, 1873.	4.3	311

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37	Causal Associations of Adiposity and Body Fat Distribution With Coronary Heart Disease, Stroke Subtypes, and Type 2 Diabetes Mellitus. Circulation, 2017, 135, 2373-2388.	1.6	304
38	Association of Diurnal Patterns in Salivary Cortisol with All-Cause and Cardiovascular Mortality: Findings from the Whitehall II Study. Journal of Clinical Endocrinology and Metabolism, 2011, 96, 1478-1485.	1.8	302
39	PCSK9 genetic variants and risk of type 2 diabetes: a mendelian randomisation study. Lancet Diabetes and Endocrinology, the, 2017, 5, 97-105.	5. 5	298
40	Trans-ancestry genome-wide association study identifies 12 genetic loci influencing blood pressure and implicates a role for DNA methylation. Nature Genetics, 2015, 47, 1282-1293.	9.4	294
41	Genetic variation near IRS1 associates with reduced adiposity and an impaired metabolic profile. Nature Genetics, 2011, 43, 753-760.	9.4	289
42	Identification of heart rate–associated loci and their effects on cardiac conduction and rhythm disorders. Nature Genetics, 2013, 45, 621-631.	9.4	282
43	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. Nature Genetics, 2014, 46, 826-836.	9.4	281
44	Utility of genetic and non-genetic risk factors in prediction of type 2 diabetes: Whitehall II prospective cohort study. BMJ: British Medical Journal, 2010, 340, b4838-b4838.	2.4	248
45	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. Nature Communications, 2016, 7, 10495.	5.8	245
46	Large-Scale Gene-Centric Meta-Analysis across 39 Studies Identifies Type 2 Diabetes Loci. American Journal of Human Genetics, 2012, 90, 410-425.	2.6	239
47	The Relationship between Smoking Status and Cortisol Secretion. Journal of Clinical Endocrinology and Metabolism, 2007, 92, 819-824.	1.8	234
48	Psychophysiological biomarkers of workplace stressors. Neuroscience and Biobehavioral Reviews, 2010, 35, 51-57.	2.9	229
49	Multivariate genome-wide analyses of the well-being spectrum. Nature Genetics, 2019, 51, 445-451.	9.4	228
50	Effects of Moderate and Vigorous Physical Activity on Heart Rate Variability in a British Study of Civil Servants. American Journal of Epidemiology, 2003, 158, 135-143.	1.6	227
51	Large-Scale Gene-Centric Meta-analysis across 32 Studies Identifies Multiple Lipid Loci. American Journal of Human Genetics, 2012, 91, 823-838.	2.6	227
52	Genomic and phenotypic insights from an atlas of genetic effects on DNA methylation. Nature Genetics, 2021, 53, 1311-1321.	9.4	218
53	Sleep epidemiologya rapidly growing field. International Journal of Epidemiology, 2011, 40, 1431-1437.	0.9	214
54	Guidance for DNA methylation studies: statistical insights from the Illumina EPIC array. BMC Genomics, 2019, 20, 366.	1.2	201

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55	Neuroendocrine and Inflammatory Factors Associated with Positive Affect in Healthy Men and Women: The Whitehall II Study. American Journal of Epidemiology, 2007, 167, 96-102.	1.6	200
56	Job Strain as a Risk Factor for Leisure-Time Physical Inactivity: An Individual-Participant Meta-Analysis of Up to 170,000 Men and Women: The IPD-Work Consortium. American Journal of Epidemiology, 2012, 176, 1078-1089.	1.6	198
57	A Genome-Wide Association Search for Type 2 Diabetes Genes in African Americans. PLoS ONE, 2012, 7, e29202.	1.1	197
58	Gene-centric Association Signals for Lipids and Apolipoproteins Identified via the HumanCVD BeadChip. American Journal of Human Genetics, 2009, 85, 628-642.	2.6	183
59	Self-Reported Sleep Duration and Sleep Disturbance Are Independently Associated with Cortisol Secretion in the Whitehall II Study. Journal of Clinical Endocrinology and Metabolism, 2009, 94, 4801-4809.	1.8	182
60	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. Nature Communications, 2017, 8, 14977.	5.8	169
61	Association of genetic variation with systolic and diastolic blood pressure among African Americans: the Candidate Gene Association Resource study. Human Molecular Genetics, 2011, 20, 2273-2284.	1.4	168
62	Interleukin-6 and C-reactive protein as predictors of cognitive decline in late midlife. Neurology, 2014, 83, 486-493.	1.5	167
63	The CIRCORT database: Reference ranges and seasonal changes in diurnal salivary cortisol derived from a meta-dataset comprised of 15 field studies. Psychoneuroendocrinology, 2016, 73, 16-23.	1.3	160
64	Blood Pressure Loci Identified with a Gene-Centric Array. American Journal of Human Genetics, 2011, 89, 688-700.	2.6	159
65	Association of Lifecourse Socioeconomic Status with Chronic Inflammation and Type 2 Diabetes Risk: The Whitehall II Prospective Cohort Study. PLoS Medicine, 2013, 10, e1001479.	3.9	158
66	Gene-centric Meta-analysis in 87,736 Individuals of European Ancestry Identifies Multiple Blood-Pressure-Related Loci. American Journal of Human Genetics, 2014, 94, 349-360.	2.6	158
67	Genome-wide physical activity interactions in adiposity ― A meta-analysis of 200,452 adults. PLoS Genetics, 2017, 13, e1006528.	1.5	158
68	Life-course influences on health in British adults: effects of socio-economic position in childhood and adulthood. International Journal of Epidemiology, 2007, 36, 532-539.	0.9	157
69	Cross-sectional versus Prospective Associations of Sleep Duration with Changes in Relative Weight and Body Fat Distribution. American Journal of Epidemiology, 2008, 167, 321-329.	1.6	150
70	Investigating the possible causal association of smoking with depression and anxiety using Mendelian randomisation meta-analysis: the CARTA consortium. BMJ Open, 2014, 4, e006141.	0.8	150
71	Apolipoprotein E genotype, cardiovascular biomarkers and risk of stroke: Systematic review and meta-analysis of 14 015 stroke cases and pooled analysis of primary biomarker data from up to 60 883 individuals. International Journal of Epidemiology, 2013, 42, 475-492.	0.9	145
72	Gender differences in the cross-sectional relationships between sleep duration and markers of inflammation: Whitehall II study. Sleep, 2009, 32, 857-64.	0.6	143

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73	Associations Between Change in Sleep Duration and Inflammation: Findings on C-reactive Protein and Interleukin 6 in the Whitehall II Study. American Journal of Epidemiology, 2013, 178, 956-961.	1.6	139
74	Inflammation, Insulin Resistance, and Diabetesâ€"Mendelian Randomization Using CRP Haplotypes Points Upstream. PLoS Medicine, 2008, 5, e155.	3.9	136
75	Social and psychosocial influences on inflammatory markers and vascular function in civil servants (the Whitehall II study). American Journal of Cardiology, 2003, 92, 984-987.	0.7	126
76	Leveraging DNA-Methylation Quantitative-Trait Loci to Characterize the Relationship between Methylomic Variation, Gene Expression, and Complex Traits. American Journal of Human Genetics, 2018, 103, 654-665.	2.6	126
77	Plasma urate concentration and risk of coronary heart disease: a Mendelian randomisation analysis. Lancet Diabetes and Endocrinology,the, 2016, 4, 327-336.	5.5	122
78	GWAS and colocalization analyses implicate carotid intima-media thickness and carotid plaque loci in cardiovascular outcomes. Nature Communications, 2018, 9, 5141.	5.8	119
79	A Nonlinear Relationship of Generalized and Central Obesity with Diurnal Cortisol Secretion in the Whitehall II Study. Journal of Clinical Endocrinology and Metabolism, 2010, 95, 4415-4423.	1.8	116
80	Secretory Phospholipase A2-IIA and Cardiovascular Disease. Journal of the American College of Cardiology, 2013, 62, 1966-1976.	1.2	115
81	Assessing cortisol from hair samples in a large observational cohort: The Whitehall II study. Psychoneuroendocrinology, 2016, 73, 148-156.	1.3	114
82	<i>PLA2G7</i> Genotype, Lipoprotein-Associated Phospholipase A ₂ Activity, and Coronary Heart Disease Risk in 10 494 Cases and 15 624 Controls of European Ancestry. Circulation, 2010, 121, 2284-2293.	1.6	111
83	Cortisol secretion and fatigue: Associations in a community based cohort. Psychoneuroendocrinology, 2009, 34, 1476-1485.	1.3	109
84	Cystatin C and Cardiovascular Disease. Journal of the American College of Cardiology, 2016, 68, 934-945.	1.2	109
85	The menopausal transition was associated in a prospective study with decreased health functioning in women who report menopausal symptoms. Journal of Clinical Epidemiology, 2005, 58, 719-727.	2.4	105
86	Effect of Smoking on Blood Pressure and Resting Heart Rate. Circulation: Cardiovascular Genetics, 2015, 8, 832-841.	5.1	105
87	The Relationship between Alcohol Consumption and Cortisol Secretion in an Aging Cohort. Journal of Clinical Endocrinology and Metabolism, 2008, 93, 750-757.	1.8	101
88	Diurnal Cortisol Patterns, Future Diabetes, and Impaired Glucose Metabolism in the Whitehall II Cohort Study. Journal of Clinical Endocrinology and Metabolism, 2016, 101, 619-625.	1.8	100
89	Recalibrating the epigenetic clock: implications for assessing biological age in the human cortex. Brain, 2020, 143, 3763-3775.	3.7	100
90	Job Strain and the Risk of Stroke. Stroke, 2015, 46, 557-559.	1.0	97

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91	Systematic underestimation of the epigenetic clock and age acceleration in older subjects. Genome Biology, 2019, 20, 283.	3.8	97
92	Separating the Mechanism-Based and Off-Target Actions of Cholesteryl Ester Transfer Protein Inhibitors With <i>CETP</i> Gene Polymorphisms. Circulation, 2010, 121, 52-62.	1.6	96
93	Genetic loci associated with heart rate variability and their effects on cardiac disease risk. Nature Communications, 2017, 8, 15805.	5.8	95
94	Comparative analysis of genome-wide association studies signals for lipids, diabetes, and coronary heart disease: Cardiovascular Biomarker Genetics Collaboration. European Heart Journal, 2012, 33, 393-407.	1.0	93
95	Psychological coping styles and cortisol over the day in healthy older adults. Psychoneuroendocrinology, 2008, 33, 601-611.	1.3	91
96	Sixty-Five Common Genetic Variants and Prediction of Type 2 Diabetes. Diabetes, 2015, 64, 1830-1840.	0.3	91
97	The Association of C-Reactive Protein and CRP Genotype with Coronary Heart Disease: Findings from Five Studies with 4,610 Cases amongst 18,637 Participants. PLoS ONE, 2008, 3, e3011.	1.1	90
98	<i>ANGPTL4</i> E40K and T266M. Arteriosclerosis, Thrombosis, and Vascular Biology, 2008, 28, 2319-2325.	1.1	89
99	Association of Diurnal Patterns in Salivary Cortisol With Type 2 Diabetes in the Whitehall II Study. Journal of Clinical Endocrinology and Metabolism, 2014, 99, 4625-4631.	1.8	89
100	Causal Effect of Plasminogen Activator Inhibitor Type 1 on Coronary Heart Disease. Journal of the American Heart Association, 2017, 6, .	1.6	89
101	Genome-wide analysis of health-related biomarkers in the UK Household Longitudinal Study reveals novel associations. Scientific Reports, 2017, 7, 11008.	1.6	88
102	Sex-dimorphic genetic effects and novel loci for fasting glucose and insulin variability. Nature Communications, 2021, 12, 24.	5.8	87
103	Genetics of cortisol secretion and depressive symptoms: A candidate gene and genome wide association approach. Psychoneuroendocrinology, 2011, 36, 1053-1061.	1.3	85
104	The Joint Effect of Sleep Duration and Disturbed Sleep on Cause-Specific Mortality: Results from the Whitehall II Cohort Study. PLoS ONE, 2014, 9, e91965.	1.1	85
105	Socioeconomic Position and DNA Methylation Age Acceleration Across the Life Course. American Journal of Epidemiology, 2018, 187, 2346-2354.	1.6	81
106	Identifying patterns in cortisol secretion in an older population. Findings from the Whitehall II study. Psychoneuroendocrinology, 2010, 35, 1091-1099.	1.3	79
107	Recurrent short sleep, chronic insomnia symptoms and salivary cortisol: A 10-year follow-up in the Whitehall II study. Psychoneuroendocrinology, 2016, 68, 91-99.	1.3	79
108	Long working hours as a risk factor for atrial fibrillation: a multi-cohort study. European Heart Journal, 2017, 38, 2621-2628.	1.0	76

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109	Social isolation in childhood and adult inflammation: Evidence from the National Child Development Study. Psychoneuroendocrinology, 2014, 50, 85-94.	1.3	75
110	The transferability of lipid loci across African, Asian and European cohorts. Nature Communications, 2019, 10, 4330.	5.8	75
111	Effects of Socioeconomic Position on Inflammatory and Hemostatic Markers: A Life-Course Analysis in the 1958 British Birth Cohort. American Journal of Epidemiology, 2008, 167, 1332-1341.	1.6	68
112	Parental separation in childhood and adult inflammation: The importance of material and psychosocial pathways. Psychoneuroendocrinology, 2013, 38, 2476-2484.	1.3	68
113	De-standardization and gender convergence in work–family life courses in Great Britain: A multi-channel sequence analysis. Advances in Life Course Research, 2015, 26, 60-75.	0.8	68
114	Examining Overweight and Obesity as Risk Factors for Common Mental Disorders Using Fat Mass and Obesity-Associated (FTO) Genotype-Instrumented Analysis: The Whitehall II Study, 1985-2004. American Journal of Epidemiology, 2011, 173, 421-429.	1.6	66
115	The Relationship Between Plasma Angiopoietin-like Protein 4 Levels, Angiopoietin-like Protein 4 Genotype, and Coronary Heart Disease Risk. Arteriosclerosis, Thrombosis, and Vascular Biology, 2010, 30, 2277-2282.	1.1	64
116	Social Determinants of von Willebrand Factor. Arteriosclerosis, Thrombosis, and Vascular Biology, 2000, 20, 1842-1847.	1.1	62
117	Measures of Social Position and Cortisol Secretion in an Aging Population: Findings From the Whitehall II Study. Psychosomatic Medicine, 2010, 72, 27-34.	1.3	62
118	Lifecourse influences on health among British adults: Effects of region of residence in childhood and adulthood. International Journal of Epidemiology, 2007, 36, 522-531.	0.9	61
119	Socio-economic trajectories and cardiovascular disease mortality in older people: the English Longitudinal Study of Ageing. International Journal of Epidemiology, 2018, 47, 36-46.	0.9	61
120	Meta-analysis of Gene-Level Associations for Rare Variants Based on Single-Variant Statistics. American Journal of Human Genetics, 2013, 93, 236-248.	2.6	60
121	Dysregulation of the hypothalamic pituitary adrenal (HPA) axis and physical performance at older ages: An individual participant meta-analysis. Psychoneuroendocrinology, 2013, 38, 40-49.	1.3	60
122	Cholesteryl Ester Transfer Protein (CETP) Polymorphisms Affect mRNA Splicing, HDL Levels, and Sex-Dependent Cardiovascular Risk. PLoS ONE, 2012, 7, e31930.	1.1	59
123	Duration of depressive symptoms and mortality risk: The English Longitudinal Study of Ageing (ELSA). British Journal of Psychiatry, 2016, 208, 337-342.	1.7	59
124	Are Flexible Work Arrangements Associated with Lower Levels of Chronic Stress-Related Biomarkers? A Study of 6025 Employees in the UK Household Longitudinal Study. Sociology, 2019, 53, 779-799.	1.7	58
125	Change in Sleep Duration and Type 2 Diabetes: The Whitehall II Study. Diabetes Care, 2015, 38, 1467-1472.	4.3	56
126	Heavier smoking may lead to a relative increase in waist circumference: evidence for a causal relationship from a Mendelian randomisation meta-analysis. The CARTA consortium: TableÂ1. BMJ Open, 2015, 5, e008808.	0.8	53

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127	Bigmelon: tools for analysing large DNA methylation datasets. Bioinformatics, 2019, 35, 981-986.	1.8	49
128	Sleep and biomarkers in the English Longitudinal Study of Ageing: Associations with C-reactive protein, fibrinogen, dehydroepiandrosterone sulfate and hemoglobin. Psychoneuroendocrinology, 2013, 38, 1484-1493.	1.3	48
129	Identification of the <i>BCAR1-CFDP1-TMEM170A</i> Locus as a Determinant of Carotid Intima-Media Thickness and Coronary Artery Disease Risk. Circulation: Cardiovascular Genetics, 2012, 5, 656-665.	5.1	47
130	Job insecurity and risk of diabetes: a meta-analysis of individual participant data. Cmaj, 2016, 188, E447-E455.	0.9	47
131	Positive affect and distressed affect over the day in older people Psychology and Aging, 2011, 26, 956-965.	1.4	46
132	Stratification by Smoking Status Reveals an Association of CHRNA5-A3-B4 Genotype with Body Mass Index in Never Smokers. PLoS Genetics, 2014, 10, e1004799.	1.5	45
133	Gene-Centric Analysis Identifies Variants Associated With Interleukin-6 Levels and Shared Pathways With Other Inflammation Markers. Circulation: Cardiovascular Genetics, 2013, 6, 163-170.	5.1	44
134	Unemployment, underweight, and obesity: Findings from Understanding Society (UKHLS). Preventive Medicine, 2017, 97, 19-25.	1.6	44
135	The relationship between physical activity, sleep duration and depressive symptoms in older adults: The English Longitudinal Study of Ageing (ELSA). Preventive Medicine Reports, 2016, 4, 512-516.	0.8	43
136	BMI and Waist Circumference as Predictors of Wellâ€being in Older Adults: Findings From the English Longitudinal Study of Ageing. Obesity, 2010, 18, 1981-1987.	1.5	40
137	Genetic Determinants of Circulating Interleukin-1 Receptor Antagonist Levels and Their Association With Glycemic Traits. Diabetes, 2014, 63, 4343-4359.	0.3	40
138	Lifecourse socioeconomic status and type 2 diabetes: the role of chronic inflammation in the English Longitudinal Study of Ageing. Scientific Reports, 2016, 6, 24780.	1.6	40
139	Does High C-reactive Protein Concentration Increase Atherosclerosis? The Whitehall II Study. PLoS ONE, 2008, 3, e3013.	1.1	39
140	Population Genomics of Cardiometabolic Traits: Design of the University College London-London School of Hygiene and Tropical Medicine-Edinburgh-Bristol (UCLEB) Consortium. PLoS ONE, 2013, 8, e71345.	1.1	39
141	Causal Relevance of Blood Lipid Fractions in the Development of Carotid Atherosclerosis. Circulation: Cardiovascular Genetics, 2013, 6, 63-72.	5.1	36
142	Circulating Fatty Acids and Risk of Coronary Heart Disease and Stroke: Individual Participant Data Metaâ€Analysis in Up to 16Â126 Participants. Journal of the American Heart Association, 2020, 9, e013131.	1.6	36
143	Social isolation and diurnal cortisol patterns in an ageing cohort. Psychoneuroendocrinology, 2013, 38, 2737-2745.	1.3	35
144	Effect of smoking on physical and cognitive capability in later life: a multicohort study using observational and genetic approaches. BMJ Open, 2015, 5, e008393.	0.8	35

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145	Work-family life courses and markers of stress and inflammation in mid-life: evidence from the National Child Development Study. International Journal of Epidemiology, 2016, 45, 1247-1259.	0.9	35
146	Investigating the causal effect of smoking on hay fever and asthma: a Mendelian randomization meta-analysis in the CARTA consortium. Scientific Reports, 2017, 7, 2224.	1.6	35
147	Circulating Apolipoprotein E Concentration and Cardiovascular Disease Risk: Meta-analysis of Results from Three Studies. PLoS Medicine, 2016, 13, e1002146.	3.9	35
148	No evidence of a longitudinal association between diurnal cortisol patterns and cognition. Neurobiology of Aging, 2014, 35, 2239-2245.	1.5	34
149	Characterising sex differences of autosomal DNA methylation in whole blood using the Illumina EPIC array. Clinical Epigenetics, 2022, 14, 62.	1.8	34
150	Influence of common genetic variation on blood lipid levels, cardiovascular risk, and coronary events in two British prospective cohort studies. European Heart Journal, 2013, 34, 972-981.	1.0	33
151	Sexâ€Specific Effects of Adiponectin on Carotid Intimaâ€Media Thickness and Incident Cardiovascular Disease. Journal of the American Heart Association, 2015, 4, e001853.	1.6	33
152	Genome-wide association study of circulating interleukin 6 levels identifies novel loci. Human Molecular Genetics, 2021, 30, 393-409.	1.4	32
153	Filaggrin gene mutations are associated with asthma and eczema in later life. Journal of Allergy and Clinical Immunology, 2008, 122, 834-836.	1.5	30
154	Is There an Association between Work Stress and Diurnal Cortisol Patterns? Findings from the Whitehall II Study. PLoS ONE, 2013, 8, e81020.	1.1	29
155	Genetic Variants Associated with von Willebrand Factor Levels in Healthy Men and Women Identified Using the HumanCVD BeadChip. Annals of Human Genetics, 2011, 75, 456-467.	0.3	28
156	Alcohol consumption and cognitive performance: a $\langle scp \rangle M \langle scp \rangle$ endelian randomization study. Addiction, 2014, 109, 1462-1471.	1.7	27
157	A systematic review and meta-analysis of 130,000 individuals shows smoking does not modify the association of APOE genotype on risk of coronary heart disease. Atherosclerosis, 2014, 237, 5-12.	0.4	27
158	Marginal role for 53 common genetic variants in cardiovascular disease prediction. Heart, 2016, 102, 1640-1647.	1.2	27
159	Proinflammatory genotype is associated with the frailty phenotype in the English Longitudinal Study of Ageing. Aging Clinical and Experimental Research, 2016, 28, 413-421.	1.4	27
160	Unemployment and inflammatory markers in England, Wales and Scotland, 1998–2012: Meta-analysis of results from 12 studies. Brain, Behavior, and Immunity, 2017, 64, 91-102.	2.0	26
161	Metabolic Profiling of Adiponectin Levels in Adults. Circulation: Cardiovascular Genetics, 2017, 10, .	5.1	26
162	Dysregulation of the hypothalamic pituitary adrenal (HPA) axis and cognitive capability at older ages: individual participant meta-analysis of five cohorts. Scientific Reports, 2019, 9, 4555.	1.6	26

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163	Glycemia, Insulin Resistance, Insulin Secretion, and Risk of Depressive Symptoms in Middle Age. Diabetes Care, 2013, 36, 928-934.	4.3	25
164	Elevated inflammatory biomarkers during unemployment: modification by age and country in the UK. Journal of Epidemiology and Community Health, 2015, 69, 673-679.	2.0	25
165	No Interactions Between Previously Associated 2-Hour Glucose Gene Variants and Physical Activity or BMI on 2-Hour Glucose Levels. Diabetes, 2012, 61, 1291-1296.	0.3	23
166	Associations between APOE and low-density lipoprotein cholesterol genotypes and cognitive and physical capability: the HALCyon programme. Age, 2014, 36, 9673.	3.0	23
167	Relationships Between Sleep Duration and von Willebrand Factor, Factor VII, and Fibrinogen. Arteriosclerosis, Thrombosis, and Vascular Biology, 2010, 30, 2032-2038.	1.1	22
168	Novel Genetic Approach to Investigate the Role of Plasma Secretory Phospholipase A2 (sPLA) Tj ETQq0 0 0 rgBT 144-150.	Overlock 5.1	10 Tf 50 547 22
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170	Socio-economic inequalities in C-reactive protein and fibrinogen across the adult age span: Findings from Understanding Society. Scientific Reports, 2017, 7, 2641.	1.6	22
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