Jari Lahti

List of Publications by Year in descending order

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254 papers 20,688 citations

14124 69 h-index 124 g-index

285 all docs

285 docs citations

times ranked

285

31004 citing authors

#	Article	IF	CITATIONS
1	Positive maternal mental health during pregnancy and mental and behavioral disorders in children: A prospective pregnancy cohort study. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2023, 64, 807-816.	3.1	11
2	Maternal haemoglobin levels in pregnancy and child DNA methylation: a study in the pregnancy and childhood epigenetics consortium. Epigenetics, 2022, 17, 19-31.	1.3	3
3	Maternal Glycemic Dysregulation During Pregnancy and Neonatal Blood DNA Methylation: Meta-analyses of Epigenome-Wide Association Studies. Diabetes Care, 2022, 45, 614-623.	4.3	19
4	Reliability of a novel approach for reference-based cell type estimation in human placental DNA methylation studies. Cellular and Molecular Life Sciences, 2022, 79, 115.	2.4	7
5	Meta-analysis of epigenome-wide associations between DNA methylation at birth and childhood cognitive skills. Molecular Psychiatry, 2022, 27, 2126-2135.	4.1	13
6	Meta-analysis of epigenome-wide association studies in newborns and children show widespread sex differences in blood DNA methylation. Mutation Research - Reviews in Mutation Research, 2022, 789, 108415.	2.4	24
7	Association of social isolation, loneliness and genetic risk with incidence of dementia: UK Biobank Cohort Study. BMJ Open, 2022, 12, e053936.	0.8	16
8	Multi-ancestry genome-wide association study of gestational diabetes mellitus highlights genetic links with type 2 diabetes. Human Molecular Genetics, 2022, 31, 3377-3391.	1.4	47
9	Polygenic prediction of educational attainment within and between families from genome-wide association analyses in 3 million individuals. Nature Genetics, 2022, 54, 437-449.	9.4	215
10	The association between overnight recognition accuracy and slow oscillation-spindle coupling is moderated by BDNF Val66Met. Behavioural Brain Research, 2022, 428, 113889.	1.2	5
11	Genetic risk of type 2 diabetes modifies the effects of a lifestyle intervention aimed at the prevention of gestational and postpartum diabetes. Diabetologia, 2022, 65, 1291-1301.	2.9	4
12	Longitudinal associations of DNA methylation and sleep in children: a meta-analysis. Clinical Epigenetics, 2022, 14, .	1.8	6
13	Shared genetic risk between eating disorder†and substanceâ€use†related phenotypes: Evidence from genomeâ€wide association studies. Addiction Biology, 2021, 26, e12880.	1.4	28
14	Genome-wide association study identifies 48 common genetic variants associated with handedness. Nature Human Behaviour, 2021, 5, 59-70.	6.2	79
15	Maternal antenatal stress and mental and behavioral disorders in their children. Journal of Affective Disorders, 2021, 278, 57-65.	2.0	24
16	DNA methylation signatures of aggression and closely related constructs: A meta-analysis of epigenome-wide studies across the lifespan. Molecular Psychiatry, 2021, 26, 2148-2162.	4.1	21
17	Genome-wide association study of circulating interleukin 6 levels identifies novel loci. Human Molecular Genetics, 2021, 30, 393-409.	1.4	32
18	Maternal anxiety during pregnancy and newborn epigenome-wide DNA methylation. Molecular Psychiatry, 2021, 26, 1832-1845.	4.1	24

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19	Combined effects of genotype and childhood adversity shape variability of DNA methylation across age. Translational Psychiatry, 2021, 11, 88.	2.4	27
20	Effectiveness and Predictors of Outcome for Psychotherapeutic Interventions in Clinical Settings Among Adolescents. Frontiers in Psychology, 2021, 12, 628977.	1.1	6
21	Reduced visual contrast suppression during major depressive episodes. Journal of Psychiatry and Neuroscience, 2021, 46, E222-E231.	1.4	20
22	An EPIC predictor of gestational age and its application to newborns conceived by assisted reproductive technologies. Clinical Epigenetics, 2021, 13, 82.	1.8	24
23	Characteristics of epigenetic aging across gestational and perinatal tissues. Clinical Epigenetics, 2021, 13, 97.	1.8	25
24	Maternal Hypertensive Pregnancy Disorders and Mental and Behavioral Disorders in the Offspring: a Review. Current Hypertension Reports, 2021, 23, 30.	1.5	7
25	Identifying nootropic drug targets via large-scale cognitive GWAS and transcriptomics. Neuropsychopharmacology, 2021, 46, 1788-1801.	2.8	12
26	Genomic and phenotypic insights from an atlas of genetic effects on DNA methylation. Nature Genetics, 2021, 53, 1311-1321.	9.4	218
27	The association between sleep-wake ratio and overnight picture recognition is moderated by BDNF genotype. Neurobiology of Learning and Memory, 2021, 177, 107353.	1.0	4
28	Variation in the SERPINA6/SERPINA1 locus alters morning plasma cortisol, hepatic corticosteroid binding globulin expression, gene expression in peripheral tissues, and risk of cardiovascular disease. Journal of Human Genetics, 2021, 66, 625-636.	1.1	40
29	Therapeutic Alliance: A Comparison Study between Adolescent Patients and Their Therapists. International Journal of Environmental Research and Public Health, 2021, 18, 11238.	1.2	0
30	The relationship between health-related quality of life and melancholic depressive symptoms is modified by brain insulin receptor gene network. Scientific Reports, 2021, 11, 21588.	1.6	5
31	The exercise-induced long noncoding RNA <i>CYTOR</i> promotes fast-twitch myogenesis in aging. Science Translational Medicine, 2021, 13, eabc7367.	5.8	19
32	Associations of antenatal glucocorticoid exposure with mental health in children. Psychological Medicine, 2020, 50, 247-257.	2.7	28
33	Maternal Gestational Diabetes Mellitus and Newborn DNA Methylation: Findings From the Pregnancy and Childhood Epigenetics Consortium. Diabetes Care, 2020, 43, 98-105.	4.3	145
34	Glucocorticoid exposure during hippocampal neurogenesis primes future stress response by inducing changes in DNA methylation. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 23280-23285.	3.3	141
35	Maternal depression and inflammation during pregnancy. Psychological Medicine, 2020, 50, 1839-1851.	2.7	30
36	Persistently High Levels of Maternal Antenatal Inflammation Are Associated With and Mediate the Effect of Prenatal Environmental Adversities on Neurodevelopmental Delay in the Offspring. Biological Psychiatry, 2020, 87, 898-907.	0.7	48

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37	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	3.7	200
38	Polygenic prediction of the risk of perinatal depressive symptoms. Depression and Anxiety, 2020, 37, 862-875.	2.0	12
39	DNA methylation and body mass index from birth to adolescence: meta-analyses of epigenome-wide association studies. Genome Medicine, 2020, 12, 105.	3.6	41
40	Carriership of two copies of C9orf72 hexanucleotide repeat intermediate-length alleles is a risk factor for ALS in the Finnish population. Acta Neuropathologica Communications, 2020, 8, 187.	2.4	16
41	Association between DNA methylation and ADHD symptoms from birth to school age: a prospective meta-analysis. Translational Psychiatry, 2020, 10, 398.	2.4	54
42	Eveningness associates with lower physical activity from pre- to late adolescence. Sleep Medicine, 2020, 74, 189-198.	0.8	17
43	Migrants Are Underrepresented in Mental Health and Rehabilitation Services—Survey and Register-Based Findings of Russian, Somali, and Kurdish Origin Adults in Finland. International Journal of Environmental Research and Public Health, 2020, 17, 6223.	1.2	13
44	A polyepigenetic glucocorticoid exposure score at birth and childhood mental and behavioral disorders. Neurobiology of Stress, 2020, 13, 100275.	1.9	8
45	Maternal Hypertensive Pregnancy Disorders and Mental Disorders in Children. Hypertension, 2020, 75, 1429-1438.	1.3	43
46	Early exposure to social disadvantages and later lifeÂbody mass index beyond genetic predisposition in three generations of Finnish birth cohorts. BMC Public Health, 2020, 20, 708.	1.2	9
47	Cord blood DNA methylation reflects cord blood C-reactive protein levels but not maternal levels: a longitudinal study and meta-analysis. Clinical Epigenetics, 2020, 12, 60.	1.8	9
48	Epigenome-wide meta-analysis of blood DNA methylation in newborns and children identifies numerous loci related to gestational age. Genome Medicine, 2020, 12, 25.	3.6	81
49	Polygenic impact of morningness on the overnight dynamics of sleep spindle amplitude. Genes, Brain and Behavior, 2020, 19, e12641.	1.1	1
50	Scheduled Telephone Support for Internet Cognitive Behavioral Therapy for Depression in Patients at Risk for Dropout: Pragmatic Randomized Controlled Trial. Journal of Medical Internet Research, 2020, 22, e15732.	2.1	14
51	Pleiotropic Meta-Analysis of Cognition, Education, and Schizophrenia Differentiates Roles of Early Neurodevelopmental and Adult Synaptic Pathways. American Journal of Human Genetics, 2019, 105, 334-350.	2.6	86
52	New alcohol-related genes suggest shared genetic mechanisms with neuropsychiatric disorders. Nature Human Behaviour, 2019, 3, 950-961.	6.2	75
53	Genome-wide Association Study of Change in Fasting Glucose over time in 13,807 non-diabetic European Ancestry Individuals. Scientific Reports, 2019, 9, 9439.	1.6	5
54	Disentangling the genetics of lean mass. American Journal of Clinical Nutrition, 2019, 109, 276-287.	2.2	38

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55	Associations of autozygosity with a broad range of human phenotypes. Nature Communications, 2019, 10, 4957.	5.8	84
56	BDNF Val66Met polymorphism moderates the association between sleep spindles and overnight visual recognition. Behavioural Brain Research, 2019, 375, 112157.	1.2	8
57	Epigenome-wide meta-analysis of DNA methylation and childhood asthma. Journal of Allergy and Clinical Immunology, 2019, 143, 2062-2074.	1.5	147
58	Hypertensive Disorders of Pregnancy and DNA Methylation in Newborns. Hypertension, 2019, 74, 375-383.	1.3	73
59	Integrated analysis of environmental and genetic influences on cord blood DNA methylation in new-borns. Nature Communications, 2019, 10, 2548.	5.8	94
60	Epigenetic upregulation of FKBP5 by aging and stress contributes to NF-κB–driven inflammation and cardiovascular risk. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 11370-11379.	3.3	193
61	The Epigenetic Clock At Birth: Associations With Maternal Antenatal Depression And Child Psychiatric Problems. European Neuropsychopharmacology, 2019, 29, S748.	0.3	0
62	Meta-analysis of epigenome-wide association studies in neonates reveals widespread differential DNA methylation associated with birthweight. Nature Communications, 2019, 10, 1893.	5.8	140
63	The Impact of Early Life Stress on Anxiety Symptoms in Late Adulthood. Scientific Reports, 2019, 9, 4395.	1.6	53
64	Polygenic Risk: Predicting Depression Outcomes in Clinical and Epidemiological Cohorts of Youths. American Journal of Psychiatry, 2019, 176, 615-625.	4.0	67
65	The associations of daylight and melatonin receptor 1B gene rs10830963 variant with glycemic traits: the prospective PPP-Botnia study. Annals of Medicine, 2019, 51, 58-67.	1.5	7
66	Genetic risk factors for schizophrenia associate with sleep spindle activity in healthy adolescents. Journal of Sleep Research, 2019, 28, e12762.	1.7	19
67	Maternal depressive symptoms during and after pregnancy are associated with poorer sleep quantity and sleep disorders in 3.5-year-old offspring. Sleep Medicine, 2019, 56, 201-210.	0.8	32
68	Infant regulatory behavior problems during first month of life and neurobehavioral outcomes in early childhood. European Child and Adolescent Psychiatry, 2019, 28, 847-859.	2.8	13
69	REM sleep fragmentation associated with depressive symptoms and genetic risk for depression in a community-based sample of adolescents. Journal of Affective Disorders, 2019, 245, 757-763.	2.0	45
70	ADHD symptoms are associated with decreased activity of fast sleep spindles and poorer procedural overnight learning during adolescence. Neurobiology of Learning and Memory, 2019, 157, 106-113.	1.0	23
71	APOE É>4, rs405509, and rs440446 promoter and intron-1 polymorphisms and dementia risk in a cohort of elderly Finns—Helsinki Birth Cohort Study. Neurobiology of Aging, 2019, 73, 230.e5-230.e8.	1.5	2
72	Schizotypal traits are associated with sleep spindles and rapid eye movement in adolescence. Journal of Sleep Research, 2019, 28, e12692.	1.7	10

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73	Polygenic risk score of SERPINA6 / SERPINA1 associates with diurnal and stress-induced HPA axis activity in children. Psychoneuroendocrinology, 2018, 93, 1-7.	1.3	13
74	Maternal early pregnancy obesity and related pregnancy and pre-pregnancy disorders: associations with child developmental milestones in the prospective PREDO Study. International Journal of Obesity, 2018, 42, 995-1007.	1.6	39
75	Maternal depressive symptoms during and after pregnancy and child developmental milestones. Depression and Anxiety, 2018, 35, 732-741.	2.0	69
76	Association of Methylation Signals With Incident Coronary Heart Disease in an Epigenome-Wide Assessment of Circulating Tumor Necrosis Factor \hat{l}_{\pm} . JAMA Cardiology, 2018, 3, 463.	3.0	33
77	Bivariate Genome-Wide Association Study of Depressive Symptoms With Type 2 Diabetes and Quantitative Glycemic Traits. Psychosomatic Medicine, 2018, 80, 242-251.	1.3	31
78	FoxO1, A2M, and TGF- \hat{l}^2 1: three novel genes predicting depression in gene X environment interactions are identified using cross-species and cross-tissues transcriptomic and miRNomic analyses. Molecular Psychiatry, 2018, 23, 2192-2208.	4.1	73
79	The Epigenetic Clock at Birth: Associations With Maternal Antenatal Depression and Child Psychiatric Problems. Journal of the American Academy of Child and Adolescent Psychiatry, 2018, 57, 321-328.e2.	0.3	78
80	Naturally occurring circadian rhythm and sleep duration are related to executive functions in early adulthood. Journal of Sleep Research, 2018, 27, 113-119.	1.7	26
81	Cohort Profile: Pregnancy And Childhood Epigenetics (PACE) Consortium. International Journal of Epidemiology, 2018, 47, 22-23u.	0.9	105
82	Association between breastfeeding and better preserved cognitive ability in an elderly cohort of Finnish men. Psychological Medicine, 2018, 48, 939-951.	2.7	6
83	Development of Late Circadian Preference: Sleep Timing From Childhood to Late Adolescence. Journal of Pediatrics, 2018, 194, 182-189.e1.	0.9	41
84	Neurocognitive outcome in young adults born lateâ€preterm. Developmental Medicine and Child Neurology, 2018, 60, 267-274.	1.1	18
85	Personality Traits Associated with Weight Maintenance among Successful Weight Losers. American Journal of Health Behavior, 2018, 42, 78-84.	0.6	8
86	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. Nature Neuroscience, 2018, 21, 1656-1669.	7.1	490
87	Genome-wide analyses identify a role for SLC17A4 and AADAT in thyroid hormone regulation. Nature Communications, 2018, 9, 4455.	5.8	181
88	Genome Analyses of >200,000 Individuals Identify 58 Loci for Chronic Inflammation and Highlight Pathways that Link Inflammation and Complex Disorders. American Journal of Human Genetics, 2018, 103, 691-706.	2.6	326
89	Neonatal regulatory behavior problems are predicted by maternal early pregnancy overweight and obesity: findings from the prospective PREDO Study. Pediatric Research, 2018, 84, 875-881.	1.1	6
90	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. Nature Communications, 2018, 9, 2098.	5.8	484

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91	Habitual coffee consumption and cognitive function: a Mendelian randomization meta-analysis in up to 415,530 participants. Scientific Reports, 2018, 8, 7526.	1.6	36
92	127. The Epigenetic Clock at Birth: Associations With Maternal Antenatal Depression and Child Psychiatric Problems. Biological Psychiatry, 2018, 83, S52.	0.7	0
93	Genome-wide association meta-analysis in 269,867 individuals identifies new genetic and functional links to intelligence. Nature Genetics, 2018, 50, 912-919.	9.4	893
94	Multi-Trait Analysis of GWAS and Biological Insights Into Cognition: A Response to Hill (2018). Twin Research and Human Genetics, 2018, 21, 394-397.	0.3	3
95	DNA Methylation Signatures of Depressive Symptoms in Middle-aged and Elderly Persons. JAMA Psychiatry, 2018, 75, 949.	6.0	78
96	The epigenetic clock and pubertal, neuroendocrine, psychiatric, and cognitive outcomes in adolescents. Clinical Epigenetics, 2018, 10, 96.	1.8	43
97	Melatonin receptor 1B gene rs10830963 polymorphism, depressive symptoms and glycaemic traits. Annals of Medicine, 2018, 50, 704-712.	1.5	6
98	Visual tool for real-time monitoring of membrane fouling via Raman spectroscopy and process model based on principal component analysis. Scientific Reports, 2018, 8, 11057.	1.6	10
99	Circadian preference and sleep timing from childhood to adolescence in relation to genetic variants from a genome-wide association study. Sleep Medicine, 2018, 50, 36-41.	0.8	18
100	Cognitive ability in young adulthood predicts risk of early-onset dementia in Finnish men. Neurology, 2018, 91, e171-e179.	1.5	6
101	Food and nutrient intakes by temperament traits: findings in the Helsinki Birth Cohort Study. European Journal of Clinical Nutrition, 2018, 72, 1136-1141.	1.3	1
102	GWAS meta-analysis reveals novel loci and genetic correlates for general cognitive function: a report from the COGENT consortium. Molecular Psychiatry, 2017, 22, 336-345.	4.1	194
103	Autism spectrum traits and visual processing in young adults with very low birth weight: the Helsinki Study of Very Low Birth Weight adults. Journal of Developmental Origins of Health and Disease, 2017, 8, 161-167.	0.7	5
104	Eveningness as a risk for behavioral problems in late adolescence. Chronobiology International, 2017, 34, 225-234.	0.9	45
105	141. FOXO1, A2M and TGFB1: Three Novel Genes Predicting Depression in Gene X Environment Interactions Are Identified Using Cross-Species and Cross-Tissues Transcriptomic and Mirnomic Analyses. Biological Psychiatry, 2017, 81, 859.	0.7	0
106	Associations between maternal risk factors of adverse pregnancy and birth outcomes and the offspring epigenetic clock of gestational age at birth. Clinical Epigenetics, 2017, 9, 49.	1.8	68
107	Maternal Licorice Consumption During Pregnancy and Pubertal, Cognitive, and Psychiatric Outcomes in Children. American Journal of Epidemiology, 2017, 185, 317-328.	1.6	44
108	Psychometric properties of the Finnish version of the Young Person's Clinical Outcomes in Routine Evaluation (YP-CORE) questionnaire. Nordic Journal of Psychiatry, 2017, 71, 250-255.	0.7	8

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109	Early Life Stress, FKBP5 Polymorphisms, and Quantitative Glycemic Traits. Psychosomatic Medicine, 2017, 79, 524-532.	1.3	6
110	Single dose of mirtazapine modulates whole-brain functional connectivity during emotional narrative processing. Psychiatry Research - Neuroimaging, 2017, 263, 61-69.	0.9	7
111	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. Biological Psychiatry, 2017, 82, 322-329.	0.7	84
112	Growth after late-preterm birth and adult cognitive, academic, and mental health outcomes. Pediatric Research, 2017, 81, 767-774.	1.1	25
113	CNV-association meta-analysis in 191,161 European adults reveals new loci associated with anthropometric traits. Nature Communications, 2017, 8, 744.	5. 8	64
114	Vitamin D and cognitive function: A Mendelian randomisation study. Scientific Reports, 2017, 7, 13230.	1.6	50
115	Large meta-analysis of genome-wide association studies identifies five loci for lean body mass. Nature Communications, 2017, 8, 80.	5. 8	147
116	Large-Scale Cognitive GWAS Meta-Analysis Reveals Tissue-Specific Neural Expression and Potential Nootropic Drug Targets. Cell Reports, 2017, 21, 2597-2613.	2.9	103
117	Circadian preference towards morningness is associated with lower slow sleep spindle amplitude and intensity in adolescents. Scientific Reports, 2017, 7, 14619.	1.6	14
118	SIRT6 polymorphism rs117385980 is associated with longevity and healthy aging in Finnish men. BMC Medical Genetics, 2017, 18, 41.	2.1	21
119	Maternal Depressive Symptoms During and After Pregnancy and Psychiatric Problems in Children. Journal of the American Academy of Child and Adolescent Psychiatry, 2017, 56, 30-39.e7.	0.3	106
120	An epigenome-wide association study meta-analysis of educational attainment. Molecular Psychiatry, 2017, 22, 1680-1690.	4.1	70
121	Nutrition after preterm birth and adult neurocognitive outcomes. PLoS ONE, 2017, 12, e0185632.	1.1	29
122	Genome-wide association meta-analysis of fish and EPA+DHA consumption in 17 US and European cohorts. PLoS ONE, 2017, 12, e0186456.	1.1	18
123	Maternal depressive symptoms during and after pregnancy are associated with attention-deficit/hyperactivity disorder symptoms in their 3- to 6-year-old children. PLoS ONE, 2017, 12, e0190248.	1.1	63
124	RÃikkönen et al. Respond to "Maternal Stress and Offspring Health― American Journal of Epidemiology, 2017, 185, 333-334.	1.6	1
125	Identification of a novel proinsulin-associated SNP and demonstration that proinsulin is unlikely to be a causal factor in subclinical vascular remodelling using Mendelian randomisation. Atherosclerosis, 2017, 266, 196-204.	0.4	3
126	Maternal prenatal anxiety and child COMT genotype predict working memory and symptoms of ADHD. PLoS ONE, 2017, 12, e0177506.	1.1	35

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127	Body image and eating behavior in young adults born preterm. International Journal of Eating Disorders, 2016, 49, 572-580.	2.1	5
128	Common mental disorders in young adults born late-preterm. Psychological Medicine, 2016, 46, 2227-2238.	2.7	16
129	Somatic, positive and negative domains of the Center for Epidemiological Studies Depression (CES-D) scale: a meta-analysis of genome-wide association studies. Psychological Medicine, 2016, 46, 1613-1623.	2.7	17
130	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
131	Genome-wide association study identifies 74 loci associated with educational attainment. Nature, 2016, 533, 539-542.	13.7	1,204
132	An epigenetic clock for gestational age at birth based on blood methylation data. Genome Biology, 2016, 17, 206.	3.8	193
133	Sleep and Lipid Profile During Transition from Childhood to Adolescence. Journal of Pediatrics, 2016, 177, 173-178.e1.	0.9	28
134	<i>KLB</i> is associated with alcohol drinking, and its gene product \hat{l}^2 -Klotho is necessary for FGF21 regulation of alcohol preference. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 14372-14377.	3.3	208
135	A principal component meta-analysis on multiple anthropometric traits identifies novel loci for body shape. Nature Communications, 2016, 7, 13357.	5.8	74
136	Prediction and Prevention of Preeclampsia and Intrauterine Growth Restriction (PREDO) study. International Journal of Epidemiology, 2016, 46, dyw154.	0.9	53
137	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. Nature Genetics, 2016, 48, 1462-1472.	9.4	284
138	Genetic variants linked to education predict longevity. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 13366-13371.	3.3	110
139	Heritability and Genome-Wide Association Analyses of Sleep Duration in Children: The EAGLE Consortium. Sleep, 2016, 39, 1859-1869.	0.6	34
140	Intellectual ability in young adulthood as an antecedent of physical functioning in older age. Age and Ageing, 2016, 45, 724-728.	0.7	2
141	APOE and aging-related cognitive change in a longitudinal cohort of men. Neurobiology of Aging, 2016, 44, 151-158.	1.5	8
142	Associations Between Self-Reported and Objectively Recorded Early Life Stress, FKBP5 Polymorphisms, and Depressive Symptoms in Midlife. Biological Psychiatry, 2016, 80, 869-877.	0.7	29
143	A single dose of mirtazapine attenuates neural responses to self-referential processing. Journal of Psychopharmacology, 2016, 30, 23-32.	2.0	13
144	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. Nature Communications, 2016, 7, 10495.	5.8	245

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145	Genome-wide meta-analysis uncovers novel loci influencing circulating leptin levels. Nature Communications, 2016, 7, 10494.	5.8	153
146	A meta-analysis of 120 246 individuals identifies 18 new loci for fibrinogen concentration. Human Molecular Genetics, 2016, 25, 358-370.	1.4	73
147	Meta-analysis of Genome-Wide Association Studies for Extraversion: Findings from the Genetics of Personality Consortium. Behavior Genetics, 2016, 46, 170-182.	1.4	178
148	GWAS for executive function and processing speed suggests involvement of the CADM2 gene. Molecular Psychiatry, 2016, 21, 189-197.	4.1	134
149	The Associations of Borderline Personality Disorder Symptoms, Five-Factor Model Personality Dimensions, and Personality Fragmentation among Depressed Inpatients. Journal of Psychiatry, 2016, 19,	0.1	2
150	ISDN2014_0385: REMOVED: A bivariate Genome Wide Association Study (GWAS) of depressive symptoms and lipid levels has identified pleiotropic gene loci. International Journal of Developmental Neuroscience, 2015, 47, 113-114.	0.7	0
151	Additive effects of maternal depressive symptoms during pregnancy and three years after childbirth on offspring psychiatric symptoms in early childhood. Psychoneuroendocrinology, 2015, 61, 12.	1.3	1
152	Independent evidence for an association between general cognitive ability and a genetic locus for educational attainment. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 363-373.	1.1	25
153	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
154	Infant Growth after Preterm Birth and Mental Health in Young Adulthood. PLoS ONE, 2015, 10, e0137092.	1.1	10
155	Late preterm birth, post-term birth, and abnormal fetal growth as risk factors for severe mental disorders from early to late adulthood. Psychological Medicine, 2015, 45, 985-999.	2.7	51
156	Meta-analysis of Genome-wide Association Studies for Neuroticism, and the Polygenic Association With Major Depressive Disorder. JAMA Psychiatry, 2015, 72, 642.	6.0	289
157	The association between lower educational attainment and depression owing to shared genetic effects? Results in ~25 000 subjects. Molecular Psychiatry, 2015, 20, 735-743.	4.1	59
158	Gene $\tilde{A}-$ dietary pattern interactions in obesity: analysis of up to 68 317 adults of European ancestry. Human Molecular Genetics, 2015, 24, 4728-4738.	1.4	84
159	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
160	Insulin Sensitivity and Secretory Response in Adults Born Preterm: The Helsinki Study of Very Low Birth Weight Adults. Journal of Clinical Endocrinology and Metabolism, 2015, 100, 244-250.	1.8	67
161	Telomere length and hypothalamic–pituitary–adrenal axis response to stress in elderly adults. Psychoneuroendocrinology, 2015, 53, 179-184.	1.3	17
162	Genetic contributions to variation in general cognitive function: a meta-analysis of genome-wide association studies in the CHARGE consortium (N=53 949). Molecular Psychiatry, 2015, 20, 183-192.	4.1	344

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163	Directional dominance on stature and cognition inÂdiverse human populations. Nature, 2015, 523, 459-462.	13.7	173
164	Poor sleep and neurocognitive function in early adolescence. Sleep Medicine, 2015, 16, 1207-1212.	0.8	75
165	Gene-Environment Interactions of Circadian-Related Genes for Cardiometabolic Traits. Diabetes Care, 2015, 38, 1456-1466.	4.3	52
166	Infant motor development and cognitive performance in early old age: the Helsinki Birth Cohort Study. Age, 2015, 37, 9785.	3.0	10
167	Late Preterm Birth and Neurocognitive Performance in Late Adulthood: A Birth Cohort Study. Pediatrics, 2015, 135, e818-e825.	1.0	76
168	Effect of Smoking on Blood Pressure and Resting Heart Rate. Circulation: Cardiovascular Genetics, 2015, 8, 832-841.	5.1	105
169	Consumption of meat is associated with higher fasting glucose and insulin concentrations regardless of glucose and insulin genetic risk scores: a meta-analysis of 50,345 Caucasians. American Journal of Clinical Nutrition, 2015, 102, 1266-1278.	2.2	69
170	Habitual sleep duration is associated with BMI and macronutrient intake and may be modified by CLOCK genetic variants. American Journal of Clinical Nutrition, 2015, 101, 135-143.	2.2	93
171	Novel loci associated with usual sleep duration: the CHARGE Consortium Genome-Wide Association Study. Molecular Psychiatry, 2015, 20, 1232-1239.	4.1	112
172	Genome-wide Studies of Verbal Declarative Memory in Nondemented Older People: The Cohorts for Heart and Aging Research in Genomic Epidemiology Consortium. Biological Psychiatry, 2015, 77, 749-763.	0.7	67
173	Genome-wide meta-analysis identifies six novel loci associated with habitual coffee consumption. Molecular Psychiatry, 2015, 20, 647-656.	4.1	235
174	Physical Activity, Body Composition and Metabolic Syndrome in Young Adults. PLoS ONE, 2015, 10, e0126737.	1.1	39
175	Prenatal and Childhood Growth, and Hospitalization for Alcohol Use Disorders in Adulthood: The Helsinki Birth Cohort Study. PLoS ONE, 2014, 9, e87404.	1.1	3
176	The Effect of Personality on Daily Life Emotional Processes. PLoS ONE, 2014, 9, e110907.	1.1	71
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