John M Starr

List of Publications by Year in descending order

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ΙΟΗΝ Μ STADD

#	Article	IF	CITATIONS
1	Meta-analysis of epigenome-wide association studies of carotid intima-media thickness. European Journal of Epidemiology, 2021, 36, 1143-1155.	5.7	10
2	Genomic and phenotypic insights from an atlas of genetic effects on DNA methylation. Nature Genetics, 2021, 53, 1311-1321.	21.4	218
3	Rare and low-frequency exonic variants and gene-by-smoking interactions in pulmonary function. Scientific Reports, 2021, 11, 19365.	3.3	2
4	Meta-analysis of up to 622,409 individuals identifies 40 novel smoking behaviour associated genetic loci. Molecular Psychiatry, 2020, 25, 2392-2409.	7.9	83
5	Polygenic predictors of age-related decline in cognitive ability. Molecular Psychiatry, 2020, 25, 2584-2598.	7.9	38
6	Ageâ€dependent DNA methylation patterns on the Y chromosome in elderly males. Aging Cell, 2020, 19, e12907.	6.7	27
7	Aluminium and fluoride in drinking water in relation to later dementia risk. British Journal of Psychiatry, 2020, 216, 29-34.	2.8	37
8	Perivascular spaces in the centrum semiovale at the beginning of the 8th decade of life: effect on cognition and associations with mineral deposition. Brain Imaging and Behavior, 2020, 14, 1865-1875.	2.1	19
9	Sleep and brain morphological changes in the eighth decade of life. Sleep Medicine, 2020, 65, 152-158.	1.6	27
10	Physical frailty and decline in general and specific cognitive abilities: the Lothian Birth Cohort 1936. Journal of Epidemiology and Community Health, 2020, 74, 108-113.	3.7	12
11	Fluctuating asymmetry in brain structure and general intelligence in 73-year-olds. Intelligence, 2020, 78, 101407.	3.0	9
12	Blood pressure and cognitive function across the eighth decade: a prospective study of the Lothian Birth Cohort of 1936. BMJ Open, 2020, 10, e033990.	1.9	4
13	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. Nature Genetics, 2020, 52, 1314-1332.	21.4	91
14	DNA methylation-based measures of accelerated biological ageing and the risk of dementia in the oldest-old: a study of the Lothian Birth Cohort 1921. BMC Psychiatry, 2020, 20, 91.	2.6	24
15	Variants associated with HHIP expression have sex-differential effects on lung function. Wellcome Open Research, 2020, 5, 111.	1.8	3
16	Neurology-related protein biomarkers are associated with cognitive ability and brain volume in older age. Nature Communications, 2020, 11, 800.	12.8	42
17	Blood DNA methylation sites predict death risk in a longitudinal study of 12, 300 individuals. Aging, 2020, 12, 14092-14124.	3.1	15
18	Brain Peak Width of Skeletonized Mean Diffusivity (PSMD) and Cognitive Function in Later Life. Frontiers in Psychiatry, 2019, 10, 524.	2.6	33

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19	Retinal microvascular features and cognitive change in the Lothianâ€Birth Cohort 1936. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2019, 11, 500-509.	2.4	8
20	Epigenome-wide association study of lung function level and its change. European Respiratory Journal, 2019, 54, 1900457.	6.7	49
21	Epigenetic signatures of smoking associate with cognitive function, brain structure, and mental and physical health outcomes in the Lothian Birth Cohort 1936. Translational Psychiatry, 2019, 9, 248.	4.8	34
22	Associations of autozygosity with a broad range of human phenotypes. Nature Communications, 2019, 10, 4957.	12.8	84
23	Predicting incident dementia $3\hat{a}\in 8$ years after brief cognitive tests in the UK Biobank prospective study of 500,000 people. Alzheimer's and Dementia, 2019, 15, 1546-1557.	0.8	28
24	Multiancestry Genome-Wide Association Study of Lipid Levels Incorporating Gene-Alcohol Interactions. American Journal of Epidemiology, 2019, 188, 1033-1054.	3.4	85
25	Sleep and cognitive aging in the eighth decade of life. Sleep, 2019, 42, .	1.1	32
26	Multi-ancestry study of blood lipid levels identifies four loci interacting with physical activity. Nature Communications, 2019, 10, 376.	12.8	64
27	Smoking does not accelerate leucocyte telomere attrition: a meta-analysis of 18 longitudinal cohorts. Royal Society Open Science, 2019, 6, 190420.	2.4	33
28	Association analyses identify 31 new risk loci for colorectal cancer susceptibility. Nature Communications, 2019, 10, 2154.	12.8	172
29	Retinal microvasculature and cerebral small vessel disease in the Lothian Birth Cohort 1936 and Mild Stroke Study. Scientific Reports, 2019, 9, 6320.	3.3	49
30	Ageing and epigenetics: linking neurodevelopmental and neurodegenerative disorders. Developmental Medicine and Child Neurology, 2019, 61, 1134-1138.	2.1	10
31	A multi-ancestry genome-wide study incorporating gene–smoking interactions identifies multiple new loci for pulse pressure and mean arterial pressure. Human Molecular Genetics, 2019, 28, 2615-2633.	2.9	31
32	Multi-ancestry genome-wide gene–smoking interaction study of 387,272 individuals identifies new loci associated with serum lipids. Nature Genetics, 2019, 51, 636-648.	21.4	112
33	Associations of Mitochondrial and Nuclear Mitochondrial Variants and Genes with Seven Metabolic Traits. American Journal of Human Genetics, 2019, 104, 112-138.	6.2	106
34	Predicting change in quality of life from age 79 to 90 in the Lothian Birth Cohort 1921. Quality of Life Research, 2019, 28, 737-749.	3.1	21
35	Older adults with intellectual disability: the National Institute for Health and Care Excellence (NICE) guidelines. Age and Ageing, 2019, 48, 14-15.	1.6	2
36	Longitudinal associations between hearing loss and general cognitive ability: The Lothian Birth Cohort 1936 Psychology and Aging, 2019, 34, 766-779.	1.6	6

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37	High-resolution magnetic resonance elastography reveals differences in subcortical gray matter viscoelasticity between young and healthy older adults. Neurobiology of Aging, 2018, 65, 158-167.	3.1	77
38	Association of Methylation Signals With Incident Coronary Heart Disease in an Epigenome-Wide Assessment of Circulating Tumor Necrosis Factor α. JAMA Cardiology, 2018, 3, 463.	6.1	33
39	Widespread associations between trait conscientiousness and thickness of brain cortical regions. NeuroImage, 2018, 176, 22-28.	4.2	22
40	A Large-Scale Multi-ancestry Genome-wide Study Accounting for Smoking Behavior Identifies Multiple Significant Loci for Blood Pressure. American Journal of Human Genetics, 2018, 102, 375-400.	6.2	123
41	Transitions across cognitive states and death among older adults in relation to education: A multistate survival model using data from six longitudinal studies. Alzheimer's and Dementia, 2018, 14, 462-472.	0.8	47
42	Meta-analysis of epigenome-wide association studies of cognitive abilities. Molecular Psychiatry, 2018, 23, 2133-2144.	7.9	68
43	Genome-Wide Meta-Analysis Unravels Interactions between Magnesium Homeostasis and Metabolic Phenotypes. Journal of the American Society of Nephrology: JASN, 2018, 29, 335-348.	6.1	34
44	Identification of a novel locus on chromosome 2q13, which predisposes to clinical vertebral fractures independently of bone density. Annals of the Rheumatic Diseases, 2018, 77, 378-385.	0.9	21
45	Brain structural differences between 73- and 92-year olds matched for childhood intelligence, social background, and intracranial volume. Neurobiology of Aging, 2018, 62, 146-158.	3.1	11
46	Green space and cognitive ageing: A retrospective life course analysis in the Lothian Birth Cohort 1936. Social Science and Medicine, 2018, 196, 56-65.	3.8	105
47	Identification of 55,000 Replicated DNA Methylation QTL. Scientific Reports, 2018, 8, 17605.	3.3	157
48	Role of cognitive ability in the association between functional health literacy and mortality in the Lothian Birth Cohort 1936: a prospective cohort study. BMJ Open, 2018, 8, e022502.	1.9	19
49	Trajectories of inflammatory biomarkers over the eighth decade and their associations with immune cell profiles and epigenetic ageing. Clinical Epigenetics, 2018, 10, 159.	4.1	30
50	Towards Standardization of Quantitative Retinal Vascular Parameters: Comparison of SIVA and VAMPIRE Measurements in the Lothian Birth Cohort 1936. Translational Vision Science and Technology, 2018, 7, 12.	2.2	55
51	Genome-wide analyses identify a role for SLC17A4 and AADAT in thyroid hormone regulation. Nature Communications, 2018, 9, 4455.	12.8	181
52	Fluid Intelligence Predicts Change in Depressive Symptoms in Later Life: The Lothian Birth Cohort 1936. Psychological Science, 2018, 29, 1984-1995.	3.3	15
53	Health literacy, cognitive ability and smoking: a cross-sectional analysis of the English Longitudinal Study of Ageing. BMJ Open, 2018, 8, e023929.	1.9	27
54	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.	6.2	326

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55	Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. Nature Genetics, 2018, 50, 1412-1425.	21.4	924
56	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. Nature Communications, 2018, 9, 2098.	12.8	484
57	GWAS on family history of Alzheimer's disease. Translational Psychiatry, 2018, 8, 99.	4.8	406
58	Genome-wide association meta-analysis in 269,867 individuals identifies new genetic and functional links to intelligence. Nature Genetics, 2018, 50, 912-919.	21.4	893
59	DNA methylation and the epigenetic clock in relation to physical frailty in older people: the Lothian Birth Cohort 1936. Clinical Epigenetics, 2018, 10, 101.	4.1	62
60	DNA methylation age is associated with an altered hemostatic profile in a multiethnic meta-analysis. Blood, 2018, 132, 1842-1850.	1.4	16
61	Exome Chip Analysis Identifies Low-Frequency and Rare Variants in <i>MRPL38</i> for White Matter Hyperintensities on Brain Magnetic Resonance Imaging. Stroke, 2018, 49, 1812-1819.	2.0	17
62	DNA Methylation Signatures of Depressive Symptoms in Middle-aged and Elderly Persons. JAMA Psychiatry, 2018, 75, 949.	11.0	78
63	Cognitive function in early and later life is associated with blood glucose in older individuals: analysis of the Lothian Birth Cohort of 1936. Diabetologia, 2018, 61, 1946-1955.	6.3	22
64	Longitudinal serum S100β and brain aging in the Lothian Birth Cohort 1936. Neurobiology of Aging, 2018, 69, 274-282.	3.1	13
65	Apolipoprotein E genotype does not moderate the associations of depressive symptoms, neuroticism and allostatic load with cognitive ability and cognitive aging in the Lothian Birth Cohort 1936. PLoS ONE, 2018, 13, e0192604.	2.5	7
66	Novel genetic associations for blood pressure identified via gene-alcohol interaction in up to 570K individuals across multiple ancestries. PLoS ONE, 2018, 13, e0198166.	2.5	94
67	Cognitive ability does not predict objectively measured sedentary behavior: Evidence from three older cohorts Psychology and Aging, 2018, 33, 288-296.	1.6	12
68	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. Nature Genetics, 2018, 50, 26-41.	21.4	286
69	Meta-analysis of exome array data identifies six novel genetic loci for lung function. Wellcome Open Research, 2018, 3, 4.	1.8	19
70	Predictors of gait speed and its change over three years in community-dwelling older people. Aging, 2018, 10, 144-153.	3.1	19
71	Rare and low-frequency coding variants alter human adult height. Nature, 2017, 542, 186-190.	27.8	544
72	Impact of small vessel disease in the brain on gait and balance. Scientific Reports, 2017, 7, 41637.	3.3	86

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73	Retinal microvascular network geometry and cognitive abilities in community-dwelling older people: The Lothian Birth Cohort 1936 study. British Journal of Ophthalmology, 2017, 101, 993-998.	3.9	25
74	Risk and protective factors for structural brain ageing in the eighth decade of life. Brain Structure and Function, 2017, 222, 3477-3490.	2.3	40
75	Common variants in CLDN14 are associated with differential excretion of magnesium over calcium in urine. Pflugers Archiv European Journal of Physiology, 2017, 469, 91-103.	2.8	27
76	Associations between hippocampal morphology, diffusion characteristics, and salivary cortisol in older men. Psychoneuroendocrinology, 2017, 78, 151-158.	2.7	9
77	Carotid disease at age 73 and cognitive change from age 70 to 76 years: A longitudinal cohort study. Journal of Cerebral Blood Flow and Metabolism, 2017, 37, 3042-3052.	4.3	13
78	Interaction of APOE e4 and poor glycemic control predicts white matter hyperintensity growth from 73 to 76. Neurobiology of Aging, 2017, 54, 54-58.	3.1	20
79	Cognitive Ability in Late Life and Onset of Physical Frailty: The Lothian Birth Cohort 1936. Journal of the American Geriatrics Society, 2017, 65, 1289-1295.	2.6	27
80	Hippocampal morphology and cognitive functions in community-dwelling older people: the Lothian Birth Cohort 1936. Neurobiology of Aging, 2017, 52, 1-11.	3.1	14
81	Exome-wide association study of plasma lipids in >300,000 individuals. Nature Genetics, 2017, 49, 1758-1766.	21.4	470
82	Processing speed and the relationship between Trail Making Test-B performance, cortical thinning and white matter microstructure in older adults. Cortex, 2017, 95, 92-103.	2.4	87
83	DNA Methylation Analysis Identifies Loci for Blood Pressure Regulation. American Journal of Human Genetics, 2017, 101, 888-902.	6.2	154
84	Large-Scale Cognitive GWAS Meta-Analysis Reveals Tissue-Specific Neural Expression and Potential Nootropic Drug Targets. Cell Reports, 2017, 21, 2597-2613.	6.4	103
85	A Novel Assessment and Profiling of Multidimensional Apathy in Alzheimer's Disease. Journal of Alzheimer's Disease, 2017, 60, 57-67.	2.6	39
86	Risk factors for dementia in the ninth decade of life and beyond: a study of the Lothian birth cohort 1921. BMC Psychiatry, 2017, 17, 205.	2.6	20
87	Age-related gene expression changes, and transcriptome wide association study of physical and cognitive aging traits, in the Lothian Birth Cohort 1936. Aging, 2017, 9, 2489-2503.	3.1	33
88	Association of Body Mass Index with DNA Methylation and Gene Expression in Blood Cells and Relations to Cardiometabolic Disease: A Mendelian Randomization Approach. PLoS Medicine, 2017, 14, e1002215.	8.4	246
89	Comparison of HapMap and 1000 Genomes Reference Panels in a Large-Scale Genome-Wide Association Study. PLoS ONE, 2017, 12, e0167742.	2.5	29
90	Genetically defined elevated homocysteine levels do not result in widespread changes of DNA methylation in leukocytes. PLoS ONE, 2017, 12, e0182472.	2.5	10

ARTICLE IF CITATIONS The complex genetics of gait speed: genome-wide meta-analysis approach. Aging, 2017, 9, 209-246. 3.1 Vascular and Mixed Dementia., 2017, , 2427-2434. 92 0 Associations among height, body mass index and intelligence from age 11 to age 78Âyears. BMC 93 2.7 Geriatrics, 2016, 16, 167 Exome Genotyping Identifies Pleiotropic Variants Associated with Red Blood Cell Traits. American 94 6.2 60 Journal of Human Genetics, 2016, 99, 8-21. Personality Polygenes, Positive Affect, and Life Satisfaction. Twin Research and Human Genetics, 2016, 0.6 16 19,407-417 Examining if being overweight really confers protection against dementia: Sixty-four year follow-up 96 of participants in the Glasgow University alumni cohort study. Journal of Negative Results in 1.4 6 BioMedicine, 2016, 15, 19. DNA methylation signatures of chronic low-grade inflammation are associated with complex diseases. 8.8 251 Genome Biology, 2016, 17, 255. Longitudinal telomere length shortening and cognitive and physical decline in later life: The Lothian 98 4.6 37 Birth Cohorts 1936 and 1921. Mechanisms of Ageing and Development, 2016, 154, 43-48. Predictors of ageing-related decline across multiple cognitive functions. Intelligence, 2016, 59, 115-126. 3.0 <scp>GWAS</scp> analysis of handgrip and lower body strength in older adults in the 100 6.7 51 <scp>CHARGE</scp> consortium. Aging Cell, 2016, 15, 792-800. Associations between education and brain structure at age 73 years, adjusted for age 11 IQ. Neurology, 1.1 46 2016, 87, 1820-1826. <i>KLB</i> is associated with alcohol drinking, and its gene product Î²-Klotho is necessary for FGF21 102 regulation of alcohol preference. Proceedings of the National Academy of Sciences of the United 7.1 208 States of America, 2016, 113, 14372-14377. Bilingualism, social cognition and executive functions: A tale of chickens and eggs. 1.6 53 Neuropsychologia, 2016, 91, 299-306. Trans-ancestry meta-analyses identify rare and common variants associated with blood pressure and 104 21.4 261 hypertension. Nature Genetics, 2016, 48, 1151-1161. Personality and Other Lifelong Influences on Older–Age Health and Wellbeing: Preliminary Findings 3.1 in Two Scottish Samples. European Journal of Personality, 2016, 30, 438-455. Genome-wide analysis identifies 12 loci influencing human reproductive behavior. Nature Genetics, 106 21.4 284 2016, 48, 1462-1472. Intelligence and socioeconomic position in childhood in relation to frailty and cumulative allostatic load in later life: the Lothian Birth Cohort 1936. Journal of Epidemiology and Community Health, 2016, 3.7 70, 576-582. Early-life predictors of resilience and related outcomes up to 66Âyears later in the 6-day sample of the 108 3.119 1947 Scottish mental survey. Social Psychiatry and Psychiatric Epidemiology, 2016, 51, 659-668.

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109	Vascular risk factors and progression of white matter hyperintensities in the Lothian Birth Cohort 1936. Neurobiology of Aging, 2016, 42, 116-123.	3.1	72
110	Life review in advanced age: qualitative research on the â€ [~] start in life' of 90-year-olds in the Lothian Birth Cohort 1921. BMC Geriatrics, 2016, 16, 74.	2.7	2
111	Large-Scale Exome-wide Association Analysis Identifies Loci for White Blood Cell Traits and Pleiotropy with Immune-Mediated Diseases. American Journal of Human Genetics, 2016, 99, 22-39.	6.2	50
112	A study of common Mendelian disease carriers across ageing British cohorts: meta-analyses reveal heterozygosity for alpha 1-antitrypsin deficiency increases respiratory capacity and height. Journal of Medical Genetics, 2016, 53, 280-288.	3.2	9
113	Progression of White Matter Disease and Cortical Thinning Are Not Related in Older Community-Dwelling Subjects. Stroke, 2016, 47, 410-416.	2.0	35
114	Polygenic risk for coronary artery disease is associated with cognitive ability in older adults. International Journal of Epidemiology, 2016, 45, 433-440.	1.9	16
115	Systems genetics identifies a convergent gene network for cognition and neurodevelopmental disease. Nature Neuroscience, 2016, 19, 223-232.	14.8	131
116	Polygenic risk of ischemic stroke is associated with cognitive ability. Neurology, 2016, 86, 611-618.	1.1	14
117	A meta-analysis of 120 246 individuals identifies 18 new loci for fibrinogen concentration. Human Molecular Genetics, 2016, 25, 358-370.	2.9	73
118	Meta-analysis of Genome-Wide Association Studies for Extraversion: Findings from the Genetics of Personality Consortium. Behavior Genetics, 2016, 46, 170-182.	2.1	178
119	Genome-wide association studies identify genetic loci for low von Willebrand factor levels. European Journal of Human Genetics, 2016, 24, 1035-1040.	2.8	45
120	Brain white matter structure and information processing speed in healthy older age. Brain Structure and Function, 2016, 221, 3223-3235.	2.3	75
121	Vascular and Mixed Dementia. , 2016, , 1-8.		0
122	Dietary factors and biomarkers of systemic inflammation in older people: the Lothian Birth Cohort 1936. British Journal of Nutrition, 2015, 114, 1088-1098.	2.3	37
123	<i>APOE/TOMM40</i> Genetic Loci, White Matter Hyperintensities, and Cerebral Microbleeds. International Journal of Stroke, 2015, 10, 1297-1300.	5.9	15
124	Rare and low-frequency variants and their association with plasma levels of fibrinogen, FVII, FVIII, and vWF. Blood, 2015, 126, e19-e29.	1.4	55
125	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.	1.7	25
126	Intelligence in Childhood and Atherosclerosis of the Carotid and Peripheral Arteries in Later Life: The Lothian Birth Cohort 1936. PLoS ONE, 2015, 10, e0125280.	2.5	0

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127	Serum cholesterol and cognitive functions: the Lothian Birth Cohort 1936. International Psychogeriatrics, 2015, 27, 439-453.	1.0	22
128	Meta-analysis of Genome-wide Association Studies for Neuroticism, and the Polygenic Association With Major Depressive Disorder. JAMA Psychiatry, 2015, 72, 642.	11.0	289
129	Coupled Changes in Brain White Matter Microstructure and Fluid Intelligence in Later Life. Journal of Neuroscience, 2015, 35, 8672-8682.	3.6	97
130	Beyond a bigger brain: Multivariable structural brain imaging and intelligence. Intelligence, 2015, 51, 47-56.	3.0	101
131	Retinal Vascular Fractal Dimension, Childhood IQ, and Cognitive Ability in Old Age: The Lothian Birth Cohort Study 1936. PLoS ONE, 2015, 10, e0121119.	2.5	26
132	Brain volumetric changes and cognitive ageing during the eighth decade of life. Human Brain Mapping, 2015, 36, 4910-4925.	3.6	79
133	Sixteen new lung function signals identified through 1000 Genomes Project reference panel imputation. Nature Communications, 2015, 6, 8658.	12.8	108
134	Effect of smoking on physical and cognitive capability in later life: a multicohort study using observational and genetic approaches. BMJ Open, 2015, 5, e008393.	1.9	35
135	Do personality traits moderate the manifestation of type 2 diabetes genetic risk?. Journal of Psychosomatic Research, 2015, 79, 303-308.	2.6	13
136	Does the Order of Item Difficulty of the Addenbrooke's Cognitive Examination Add Anything to Subdomain Scores in the Clinical Assessment of Dementia. Dementia and Geriatric Cognitive Disorders Extra, 2015, 5, 155-169.	1.3	4
137	Association Between Psychological Distress and Liver Disease Mortality: A Meta-analysis of Individual Study Participants. Gastroenterology, 2015, 148, 958-966.e4.	1.3	85
138	Cognitive consequences of overweight and obesity in the ninth decade of life?. Age and Ageing, 2015, 44, 59-65.	1.6	41
139	Pulmonary function as a risk factor for dementia death: an individual participant meta-analysis of six UK general population cohort studies. Journal of Epidemiology and Community Health, 2015, 69, 550-556.	3.7	34
140	Directional dominance on stature and cognition inÂdiverse human populations. Nature, 2015, 523, 459-462.	27.8	173
141	Improving Phenotypic Prediction by Combining Genetic and Epigenetic Associations. American Journal of Human Genetics, 2015, 97, 75-85.	6.2	116
142	DNA methylation age of blood predicts all-cause mortality in later life. Genome Biology, 2015, 16, 25.	8.8	928
143	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.	21.4	294
144	Brain iron deposits and lifespan cognitive ability. Age, 2015, 37, 100.	3.0	24

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145	The epigenetic clock is correlated with physical and cognitive fitness in the Lothian Birth Cohort 1936. International Journal of Epidemiology, 2015, 44, 1388-1396.	1.9	472
146	Association of allostatic load with brain structure and cognitive ability in later life. Neurobiology of Aging, 2015, 36, 1390-1399.	3.1	67
147	Total MRI load of cerebral small vessel disease and cognitive ability in older people. Neurobiology of Aging, 2015, 36, 2806-2811.	3.1	199
148	Genes From a Translational Analysis Support a Multifactorial Nature of White Matter Hyperintensities. Stroke, 2015, 46, 341-347.	2.0	33
149	Post-mortem brain analyses of the Lothian Birth Cohort 1936: extending lifetime cognitive and brain phenotyping to the level of the synapse. Acta Neuropathologica Communications, 2015, 3, 53.	5.2	25
150	Compensation or inhibitory failure? Testing hypotheses of age-related right frontal lobe involvement in verbal memory ability using structural and diffusion MRI. Cortex, 2015, 63, 4-15.	2.4	19
151	Target risk factors for dementia prevention: a systematic review and Delphi consensus study on the evidence from observational studies. International Journal of Geriatric Psychiatry, 2015, 30, 234-246.	2.7	363
152	White matter hyperintensities and normal-appearing white matter integrity in the aging brain. Neurobiology of Aging, 2015, 36, 909-918.	3.1	224
153	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.	1.3	67
154	Prediction of general hospital admission in people with dementia: Cohort study. British Journal of Psychiatry, 2015, 206, 153-159.	2.8	28
155	Genetic Basis of a Cognitive Complexity Metric. PLoS ONE, 2015, 10, e0123886.	2.5	22
156	Large-Scale Genome-Wide Association Studies and Meta-Analyses of Longitudinal Change in Adult Lung Function. PLoS ONE, 2014, 9, e100776.	2.5	52
157	No Evidence for Genome-Wide Interactions on Plasma Fibrinogen by Smoking, Alcohol Consumption and Body Mass Index: Results from Meta-Analyses of 80,607 Subjects. PLoS ONE, 2014, 9, e111156.	2.5	8
158	Grip Strength across the Life Course: Normative Data from Twelve British Studies. PLoS ONE, 2014, 9, e113637.	2.5	734
159	Polygenic Risk for Alzheimer's Disease is not Associated with Cognitive Ability or Cognitive Aging in Non-Demented Older People. Journal of Alzheimer's Disease, 2014, 39, 565-574.	2.6	63
160	Life course influences of physical and cognitive function and personality on attitudes to aging in the Lothian Birth Cohort 1936. International Psychogeriatrics, 2014, 26, 1417-1430.	1.0	30
161	Structure and correlates of cognitive aging in a narrow age cohort Psychology and Aging, 2014, 29, 236-249.	1.6	62
162	Potential effect of skull thickening on the associations between cognition and brain atrophy in ageing. Age and Ageing, 2014, 43, 712-716.	1.6	6

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163	Lawton IADL scale in dementia: can item response theory make it more informative?. Age and Ageing, 2014, 43, 491-495.	1.6	20
164	Common genetic variants associated with cognitive performance identified using the proxy-phenotype method. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 13790-13794.	7.1	244
165	Urinary incontinence in people with Alzheimer's disease. International Journal of Geriatric Psychiatry, 2014, 29, 107-109.	2.7	12
166	Genetic diversity is a predictor of mortality in humans. BMC Genetics, 2014, 15, 159.	2.7	12
167	Skin Aging and Oxidative Stress. , 2014, , 15-22.		6
168	Association of Low-Frequency and Rare Coding-Sequence Variants with Blood Lipids and Coronary Heart Disease in 56,000 Whites and Blacks. American Journal of Human Genetics, 2014, 94, 223-232.	6.2	287
169	Height in relation to dementia death: individual participant meta-analysis of 18 UK prospective cohort studies. British Journal of Psychiatry, 2014, 205, 348-354.	2.8	49
170	Genome-wide association analysis identifies six new loci associated with forced vital capacity. Nature Genetics, 2014, 46, 669-677.	21.4	131
171	Quantitative multi-modal MRI of the Hippocampus and cognitive ability in community-dwelling older subjects. Cortex, 2014, 53, 34-44.	2.4	22
172	Genetic and environmental exposures constrain epigenetic drift over the human life course. Genome Research, 2014, 24, 1725-1733.	5.5	152
173	Multi-site study of additive genetic effects on fractional anisotropy of cerebral white matter: Comparing meta and megaanalytical approaches for data pooling. NeuroImage, 2014, 95, 136-150.	4.2	127
174	Alzheimer's disease susceptibility genes APOE and TOMM40, and brain white matter integrity in the Lothian Birth Cohort 1936. Neurobiology of Aging, 2014, 35, 1513.e25-1513.e33.	3.1	58
175	The Attitudes to Ageing Questionnaire: Mokken Scaling Analysis. PLoS ONE, 2014, 9, e99100.	2.5	12
176	Lower ankle-brachial index is related to worse cognitive performance in old age Neuropsychology, 2014, 28, 281-289.	1.3	16
177	Personality, health, and brain integrity: The Lothian Birth Cohort Study 1936 Health Psychology, 2014, 33, 1477-1486.	1.6	38
178	Towards understanding the links between health literacy and physical health Health Psychology, 2014, 33, 164-173.	1.6	91
179	Functional Gene Group Analysis Indicates No Role for Heterotrimeric G Proteins in Cognitive Ability. PLoS ONE, 2014, 9, e91690.	2.5	3
180	Does the Addenbrooke's Cognitive Examinationâ€revised add to the Miniâ€Mental State Examination in established Alzheimer disease? Results from a national dementia research register. International Journal of Geriatric Psychiatry, 2013, 28, 351-355.	2.7	31

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181	Editorial. Archives of Gerontology and Geriatrics, 2013, 56, 405-406.	3.0	0
182	Brain white matter damage in aging and cognitive ability in youth and older age. Neurobiology of Aging, 2013, 34, 2740-2747.	3.1	83
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