

Toshiko Tanaka

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/4388269/toshiko-tanaka-publications-by-year.pdf>

Version: 2024-04-27

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

131
papers

15,047
citations

52
h-index

122
g-index

140
ext. papers

19,147
ext. citations

10.6
avg, IF

5.07
L-index

#	Paper	IF	Citations
131	Prior psychosocial profile and perceived impact of the COVID-19 pandemic: insights from the Baltimore Longitudinal Study of Aging.. <i>Aging Clinical and Experimental Research</i> , 2022 , 1	4.8	
130	DNA methylation signature of chronic low-grade inflammation and its role in cardio-respiratory diseases.. <i>Nature Communications</i> , 2022 , 13, 2408	17.4	1
129	Understanding the Human Aging Proteome Using Epidemiological Models. <i>Methods in Molecular Biology</i> , 2022 , 173-192	1.4	
128	Metabolomic Profile of Different Dietary Patterns and Their Association with Frailty Index in Community-Dwelling Older Men and Women. <i>Nutrients</i> , 2022 , 14, 2237	6.7	0
127	Dietary Pattern Trajectories in Middle Age and Physical Function in Older Age. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2021 , 76, 513-519	6.4	2
126	A brain proteomic signature of incipient Alzheimer's disease in young ϵ carriers identifies novel drug targets. <i>Science Advances</i> , 2021 , 7, eabi8178	14.3	2
125	Adherence to the Mediterranean diet assessed by a novel dietary biomarker score and mortality in older adults: the InCHIANTI cohort study. <i>BMC Medicine</i> , 2021 , 19, 280	11.4	2
124	DNA methylation signatures reveal that distinct combinations of transcription factors specify human immune cell epigenetic identity. <i>Immunity</i> , 2021 , 54, 2465-2480.e5	32.3	2
123	Predicting physiological aging rates from a range of quantitative traits using machine learning. <i>Aging</i> , 2021 , 13, 23471-23516	5.6	0
122	Elevated Plasma Growth and Differentiation Factor 15 Predicts Incident Anemia in Older Adults Aged 60 Years and Older. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2021 , 76, 1192-1197	6.4	3
121	Association Between the Multidimensional Prognostic Index and Mortality During 15 Years of Follow-up in the InCHIANTI Study. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2021 , 76, 1678-1685	6.4	6
120	Association of Adherence to the Mediterranean-Style Diet with Lower Frailty Index in Older Adults. <i>Nutrients</i> , 2021 , 13,	6.7	4
119	Proteomics in aging research: A roadmap to clinical, translational research. <i>Aging Cell</i> , 2021 , 20, e13325	9.9	10
118	Proteomics and Epidemiological Models of Human Aging. <i>Frontiers in Physiology</i> , 2021 , 12, 674013	4.6	2
117	Genome-wide association studies identify 137 genetic loci for DNA methylation biomarkers of aging. <i>Genome Biology</i> , 2021 , 22, 194	18.3	14
116	Characterization of the plasma proteomic profile of frailty phenotype. <i>GeroScience</i> , 2021 , 43, 1029-1037	8.9	10
115	Genome-wide meta-analysis of muscle weakness identifies 15 susceptibility loci in older men and women. <i>Nature Communications</i> , 2021 , 12, 654	17.4	10

114	Sex-dimorphic genetic effects and novel loci for fasting glucose and insulin variability. <i>Nature Communications</i> , 2021 , 12, 24	17.4	30
113	Genome-wide association study of circulating interleukin 6 levels identifies novel loci. <i>Human Molecular Genetics</i> , 2021 , 30, 393-409	5.6	6
112	Genome-wide association study identifies novel susceptibility loci for KIT D816V positive mastocytosis. <i>American Journal of Human Genetics</i> , 2021 , 108, 284-294	11	6
111	Genetic analysis of dietary intake identifies new loci and functional links with metabolic traits. <i>Nature Human Behaviour</i> , 2021 ,	12.8	5
110	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021 , 596, 393-397	10.4	28
109	Protein signatures of centenarians and their offspring suggest centenarians age slower than other humans. <i>Aging Cell</i> , 2021 , 20, e13290	9.9	14
108	The Mediterranean-DASH Intervention for Neurodegenerative Delay (MIND) diet is associated with physical function and grip strength in older men and women. <i>American Journal of Clinical Nutrition</i> , 2021 ,	7	1
107	Age-associated difference in circulating ACE2, the gateway for SARS-COV-2, in humans: results from the InCHIANTI study. <i>GeroScience</i> , 2021 , 43, 619-627	8.9	7
106	Interaction between Apolipoprotein E and Butyrylcholinesterase Genes on Risk of Alzheimer's Disease in a Prospective Cohort Study. <i>Journal of Alzheimers Disease</i> , 2020 , 75, 417-427	4.3	
105	Plasma proteomic signatures predict dementia and cognitive impairment. <i>Alzheimers and Dementia: Translational Research and Clinical Interventions</i> , 2020 , 6, e12018	6	6
104	Blood DNA Methylation and Aging: A Cross-Sectional Analysis and Longitudinal Validation in the InCHIANTI Study. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2020 , 75, 2051-2055	6.4	6
103	Plasma proteomic signature of the risk of developing mobility disability: A 9-year follow-up. <i>Aging Cell</i> , 2020 , 19, e13132	9.9	17
102	Comparing Analytical Methods for the Gut Microbiome and Aging: Gut Microbial Communities and Body Weight in the Osteoporotic Fractures in Men (MrOS) Study. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2020 , 75, 1267-1275	6.4	3
101	Blood Metabolite Signatures of Metabolic Syndrome in Two Cross-Cultural Older Adult Cohorts. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	10
100	Blood DNA methylation sites predict death risk in a longitudinal study of 12, 300 individuals. <i>Aging</i> , 2020 , 12, 14092-14124	5.6	6
99	Blood Metabolite Signature of Metabolic Syndrome Implicates Alterations in Amino Acid Metabolism: Findings from the Baltimore Longitudinal Study of Aging (BLSA) and the Tsuruoka Metabolomics Cohort Study (TMCS). <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	7
98	Plasma proteomic biomarker signature of age predicts health and life span. <i>ELife</i> , 2020 , 9,	8.9	18
97	Measuring biological aging in humans: A quest. <i>Aging Cell</i> , 2020 , 19, e13080	9.9	158

96	Underlying features of epigenetic aging clocks in vivo and in vitro. <i>Aging Cell</i> , 2020 , 19, e13229	9.9	37
95	A Plasma Proteomic Signature of Skeletal Muscle Mitochondrial Function. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	3
94	Elevated Plasma Growth and Differentiation Factor 15 Is Associated With Slower Gait Speed and Lower Physical Performance in Healthy Community-Dwelling Adults. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2020 , 75, 175-180	6.4	27
93	Association of dietary folate and vitamin B-12 intake with genome-wide DNA methylation in blood: a large-scale epigenome-wide association analysis in 5841 individuals. <i>American Journal of Clinical Nutrition</i> , 2019 , 110, 437-450	7	22
92	Genome-wide association study of breakfast skipping links clock regulation with food timing. <i>American Journal of Clinical Nutrition</i> , 2019 , 110, 473-484	7	22
91	An integrative cross-omics analysis of DNA methylation sites of glucose and insulin homeostasis. <i>Nature Communications</i> , 2019 , 10, 2581	17.4	31
90	Cardiovascular Health Is Associated With Disability Among Older Community Dwelling Men and Women. <i>Journal of Aging and Health</i> , 2019 , 31, 1339-1352	2.6	2
89	Genome-wide meta-analysis of macronutrient intake of 91,114 European ancestry participants from the cohorts for heart and aging research in genomic epidemiology consortium. <i>Molecular Psychiatry</i> , 2019 , 24, 1920-1932	15.1	30
88	Blood Leukocyte DNA Methylation Predicts Risk of Future Myocardial Infarction and Coronary Heart Disease. <i>Circulation</i> , 2019 , 140, 645-657	16.7	65
87	Potential Interplay between Dietary Saturated Fats and Genetic Variants of the NLRP3 Inflammasome to Modulate Insulin Resistance and Diabetes Risk: Insights from a Meta-Analysis of 19005 Individuals. <i>Molecular Nutrition and Food Research</i> , 2019 , 63, e1900226	5.9	11
86	A SERUM PROTEIN SIGNATURE OF APOE GENOTYPES IN CENTENARIANS. <i>Innovation in Aging</i> , 2019 , 3, S621-S622	0.1	78
85	Association of Methylation Signals With Incident Coronary Heart Disease in an Epigenome-Wide Assessment of Circulating Tumor Necrosis Factor α . <i>JAMA Cardiology</i> , 2018 , 3, 463-472	16.2	17
84	Skeletal muscle ex vivo mitochondrial respiration parallels decline in vivo oxidative capacity, cardiorespiratory fitness, and muscle strength: The Baltimore Longitudinal Study of Aging. <i>Aging Cell</i> , 2018 , 17, e12725	9.9	57
83	Sugar-sweetened beverage intake associations with fasting glucose and insulin concentrations are not modified by selected genetic variants in a ChREBP-FGF21 pathway: a meta-analysis. <i>Diabetologia</i> , 2018 , 61, 317-330	10.3	17
82	Genome-Wide Interactions with Dairy Intake for Body Mass Index in Adults of European Descent. <i>Molecular Nutrition and Food Research</i> , 2018 , 62, 1700347	5.9	5
81	Plasma proteomic signature of age in healthy humans. <i>Aging Cell</i> , 2018 , 17, e12799	9.9	167
80	SPARCL1 Accelerates Symptom Onset in Alzheimer's Disease and Influences Brain Structure and Function During Aging. <i>Journal of Alzheimer's Disease</i> , 2018 , 61, 401-414	4.3	12
79	Identification of a novel locus on chromosome 2q13, which predisposes to clinical vertebral fractures independently of bone density. <i>Annals of the Rheumatic Diseases</i> , 2018 , 77, 378-385	2.4	15

78	Adherence to a Mediterranean Diet Protects from Cognitive Decline in the Invecchiare in Chianti Study of Aging. <i>Nutrients</i> , 2018 , 10,	6.7	23
77	Genome-wide analyses identify a role for SLC17A4 and AADAT in thyroid hormone regulation. <i>Nature Communications</i> , 2018 , 9, 4455	17.4	75
76	Genome Analyses of >200,000 Individuals Identify 58 Loci for Chronic Inflammation and Highlight Pathways that Link Inflammation and Complex Disorders. <i>American Journal of Human Genetics</i> , 2018 , 103, 691-706	11	151
75	Extracellular RNA profiles with human age. <i>Aging Cell</i> , 2018 , 17, e12785	9.9	16
74	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. <i>Nature Genetics</i> , 2017 , 49, 834-841	36.3	257
73	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. <i>Nature Communications</i> , 2017 , 8, 14977	17.4	105
72	Discovery and fine-mapping of loci associated with MUFAs through trans-ethnic meta-analysis in Chinese and European populations. <i>Journal of Lipid Research</i> , 2017 , 58, 974-981	6.3	12
71	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. <i>Biological Psychiatry</i> , 2017 , 82, 322-329	7.9	68
70	A double blind placebo controlled randomized trial of the effect of acute uric acid changes on inflammatory markers in humans: A pilot study. <i>PLoS ONE</i> , 2017 , 12, e0181100	3.7	12
69	Genome-wide association meta-analysis of fish and EPA+DHA consumption in 17 US and European cohorts. <i>PLoS ONE</i> , 2017 , 12, e0186456	3.7	15
68	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. <i>PLoS Medicine</i> , 2017 , 14, e1002383	11.6	223
67	Epigenetic clock analysis of diet, exercise, education, and lifestyle factors. <i>Aging</i> , 2017 , 9, 419-446	5.6	317
66	Genome-wide Association Study of Parental Life Span. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2017 , 72, 1407-1410	6.4	5
65	Cardiovascular Health Is Associated With Physical Function Among Older Community Dwelling Men and Women. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2017 , 72, 1710-1718	6.4	17
64	Sex-Dependent Associations of Serum Uric Acid with Brain Function During Aging. <i>Journal of Alzheimers Disease</i> , 2017 , 60, 699-706	4.3	11
63	Cross-Sectional and Longitudinal Effects of Genotypes on Individual Differences in Memory and Executive Function: Findings from the BLSA. <i>Frontiers in Aging Neuroscience</i> , 2017 , 9, 142	5.3	6
62	Genome-wide physical activity interactions in adiposity - A meta-analysis of 200,452 adults. <i>PLoS Genetics</i> , 2017 , 13, e1006528	6	103
61	Comparison of HapMap and 1000 Genomes Reference Panels in a Large-Scale Genome-Wide Association Study. <i>PLoS ONE</i> , 2017 , 12, e0167742	3.7	21

60	The complex genetics of gait speed: genome-wide meta-analysis approach. <i>Aging</i> , 2017 , 9, 209-246	5.6	16
59	52 Genetic Loci Influencing Myocardial Mass. <i>Journal of the American College of Cardiology</i> , 2016 , 68, 1435-1448	15.1	76
58	The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. <i>Nature Genetics</i> , 2016 , 48, 1171-1184	36.3	251
57	Association Between Accelerated Multimorbidity and Age-Related Cognitive Decline in Older Baltimore Longitudinal Study of Aging Participants without Dementia. <i>Journal of the American Geriatrics Society</i> , 2016 , 64, 965-72	5.6	58
56	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. <i>Nature Communications</i> , 2016 , 7, 10495	17.4	180
55	Genome-wide meta-analysis uncovers novel loci influencing circulating leptin levels. <i>Nature Communications</i> , 2016 , 7, 10494	17.4	107
54	Change in Epigenome-Wide DNA Methylation Over 9 Years and Subsequent Mortality: Results From the InCHIANTI Study. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2016 , 71, 1029-35	6.4	28
53	A meta-analysis of 120 246 individuals identifies 18 new loci for fibrinogen concentration. <i>Human Molecular Genetics</i> , 2016 , 25, 358-70	5.6	54
52	State- and trait-dependent associations of vitamin-D with brain function during aging. <i>Neurobiology of Aging</i> , 2016 , 39, 38-45	5.6	18
51	DNA methylation-based measures of biological age: meta-analysis predicting time to death. <i>Aging</i> , 2016 , 8, 1844-1865	5.6	531
50	Associations between Common Variants in Iron-Related Genes with Haematological Traits in Populations of African Ancestry. <i>PLoS ONE</i> , 2016 , 11, e0157996	3.7	11
49	DNA methylation signatures of chronic low-grade inflammation are associated with complex diseases. <i>Genome Biology</i> , 2016 , 17, 255	18.3	171
48	Fine-mapping, novel loci identification, and SNP association transferability in a genome-wide association study of QRS duration in African Americans. <i>Human Molecular Genetics</i> , 2016 , 25, 4350-4368	5.6	20
47	GWAS analysis of handgrip and lower body strength in older adults in the CHARGE consortium. <i>Aging Cell</i> , 2016 , 15, 792-800	9.9	33
46	Genomewide meta-analysis identifies loci associated with IGF-I and IGFBP-3 levels with impact on age-related traits. <i>Aging Cell</i> , 2016 , 15, 811-24	9.9	71
45	Gene-Environment Interactions of Circadian-Related Genes for Cardiometabolic Traits. <i>Diabetes Care</i> , 2015 , 38, 1456-66	14.6	36
44	GWAS of longevity in CHARGE consortium confirms APOE and FOXO3 candidacy. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2015 , 70, 110-8	6.4	188
43	Whole-genome sequencing to understand the genetic architecture of common gene expression and biomarker phenotypes. <i>Human Molecular Genetics</i> , 2015 , 24, 1504-12	5.6	7

42	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. <i>Nature Genetics</i> , 2015 , 47, 1294-1303	36.3	226
41	Genetic variation at MECOM, TERT, JAK2 and HBS1L-MYB predisposes to myeloproliferative neoplasms. <i>Nature Communications</i> , 2015 , 6, 6691	17.4	120
40	Effects of a behavioral intervention that emphasizes spices and herbs on adherence to recommended sodium intake: results of the SPICE randomized clinical trial. <i>American Journal of Clinical Nutrition</i> , 2015 , 102, 671-9	7	36
39	Genome-wide studies of verbal declarative memory in nondemented older people: the Cohorts for Heart and Aging Research in Genomic Epidemiology consortium. <i>Biological Psychiatry</i> , 2015 , 77, 749-63	7.9	48
38	Genome-wide meta-analysis identifies six novel loci associated with habitual coffee consumption. <i>Molecular Psychiatry</i> , 2015 , 20, 647-656	15.1	167
37	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. <i>PLoS Genetics</i> , 2015 , 11, e1005378	6	220
36	New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015 , 518, 187-196	50.4	920
35	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015 , 518, 197-206	50.4	2687
34	Dietary fatty acids modulate associations between genetic variants and circulating fatty acids in plasma and erythrocyte membranes: Meta-analysis of nine studies in the CHARGE consortium. <i>Molecular Nutrition and Food Research</i> , 2015 , 59, 1373-83	5.9	32
33	Novel loci affecting iron homeostasis and their effects in individuals at risk for hemochromatosis. <i>Nature Communications</i> , 2014 , 5, 4926	17.4	121
32	Meta-analysis of genome-wide association studies for circulating phyloquinone concentrations. <i>American Journal of Clinical Nutrition</i> , 2014 , 100, 1462-9	7	27
31	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014 , 46, 1173-86	36.3	1339
30	Genetics of Energy and Macronutrient Intake in Humans. <i>Current Nutrition Reports</i> , 2014 , 3, 170-177	6	3
29	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. <i>Nature Genetics</i> , 2014 , 46, 826-36	36.3	199
28	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. <i>PLoS ONE</i> , 2014 , 9, e1111156	3.7	5
27	Genome-wide association study of plasma N6 polyunsaturated fatty acids within the cohorts for heart and aging research in genomic epidemiology consortium. <i>Circulation: Cardiovascular Genetics</i> , 2014 , 7, 321-331		112
26	FTO genetic variants, dietary intake and body mass index: insights from 177,330 individuals. <i>Human Molecular Genetics</i> , 2014 , 23, 6961-72	5.6	120
25	PRKCZ methylation is associated with sunlight exposure in a North American but not a Mediterranean population. <i>Chronobiology International</i> , 2014 , 31, 1034-40	3.6	10

24	Genetic diversity is a predictor of mortality in humans. <i>BMC Genetics</i> , 2014 , 15, 159	2.6	6
23	Common genetic loci influencing plasma homocysteine concentrations and their effect on risk of coronary artery disease. <i>American Journal of Clinical Nutrition</i> , 2013 , 98, 668-76	7	122
22	Genome-wide meta-analysis of observational studies shows common genetic variants associated with macronutrient intake. <i>American Journal of Clinical Nutrition</i> , 2013 , 97, 1395-402	7	161
21	Novel locus including FGF21 is associated with dietary macronutrient intake. <i>Human Molecular Genetics</i> , 2013 , 22, 1895-902	5.6	134
20	Vitamin D receptor and megalin gene polymorphisms are associated with central adiposity status and changes among US adults. <i>Journal of Nutritional Science</i> , 2013 , 2, e33	2.7	14
19	Caffeine, alcohol and overall nutrient adequacy are associated with longitudinal cognitive performance among US adults. <i>FASEB Journal</i> , 2013 , 27, 346.4	0.9	
18	Seventy-five genetic loci influencing the human red blood cell. <i>Nature</i> , 2012 , 492, 369-75	50.4	257
17	A genome-wide approach accounting for body mass index identifies genetic variants influencing fasting glycemic traits and insulin resistance. <i>Nature Genetics</i> , 2012 , 44, 659-69	36.3	615
16	A higher adherence to a Mediterranean-style diet is inversely associated with the development of frailty in community-dwelling elderly men and women. <i>Journal of Nutrition</i> , 2012 , 142, 2161-6	4.1	161
15	Large-scale association analyses identify new loci influencing glycemic traits and provide insight into the underlying biological pathways. <i>Nature Genetics</i> , 2012 , 44, 991-1005	36.3	621
14	Mediterranean diet and mobility decline in older persons. <i>Experimental Gerontology</i> , 2011 , 46, 303-8	4.5	108
13	Contribution of Diet and Genes to Polyunsaturated Fatty Acid Composition. <i>Current Cardiovascular Risk Reports</i> , 2011 , 5, 45-51	0.9	
12	CUBN is a gene locus for albuminuria. <i>Journal of the American Society of Nephrology: JASN</i> , 2011 , 22, 555-1707	17.7	170
11	Novel association to the proprotein convertase PCSK7 gene locus revealed by analysing soluble transferrin receptor (sTfR) levels. <i>Human Molecular Genetics</i> , 2011 , 20, 1042-7	5.6	51
10	Identification of a common variant in the TFR2 gene implicated in the physiological regulation of serum iron levels. <i>Human Molecular Genetics</i> , 2011 , 20, 1232-40	5.6	59
9	Multiple loci are associated with white blood cell phenotypes. <i>PLoS Genetics</i> , 2011 , 7, e1002113	6	92
8	Genetic loci associated with plasma phospholipid n-3 fatty acids: a meta-analysis of genome-wide association studies from the CHARGE Consortium. <i>PLoS Genetics</i> , 2011 , 7, e1002193	6	257
7	Genetic variation in GIPR influences the glucose and insulin responses to an oral glucose challenge. <i>Nature Genetics</i> , 2010 , 42, 142-8	36.3	527

6	Common variants at 10 genomic loci influence hemoglobin A _{1c} levels via glyceimic and nonglyceimic pathways. <i>Diabetes</i> , 2010 , 59, 3229-39	0.9	314
5	A genome-wide association analysis of serum iron concentrations. <i>Blood</i> , 2010 , 115, 94-6	2.2	117
4	Genome-wide association analysis of total cholesterol and high-density lipoprotein cholesterol levels using the Framingham heart study data. <i>BMC Medical Genetics</i> , 2010 , 11, 55	2.1	64
3	Genome-wide association study of plasma polyunsaturated fatty acids in the InCHIANTI Study. <i>PLoS Genetics</i> , 2009 , 5, e1000338	6	300
2	Genome-wide association study of vitamin B6, vitamin B12, folate, and homocysteine blood concentrations. <i>American Journal of Human Genetics</i> , 2009 , 84, 477-82	11	193
1	Multi-trait genome-wide association meta-analysis of dietary intake identifies new loci and genetic and functional links with metabolic traits		3