## John B Whitfield

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Biological, clinical and population relevance of 95 loci for blood lipids. Nature, 2010, 466, 707-713.	27.8	3,249
2	Discovery and refinement of loci associated with lipid levels. Nature Genetics, 2013, 45, 1274-1283.	21.4	2,641
3	Association studies of up to 1.2 million individuals yield new insights into the genetic etiology of tobacco and alcohol use. Nature Genetics, 2019, 51, 237-244.	21.4	1,307
4	Loci influencing lipid levels and coronary heart disease risk in 16 European population cohorts. Nature Genetics, 2009, 41, 47-55.	21.4	776
5	Common variants associated with plasma triglycerides and risk for coronary artery disease. Nature Genetics, 2013, 45, 1345-1352.	21.4	754
6	Genome-wide association analyses identify 18 new loci associated with serum urate concentrations. Nature Genetics, 2013, 45, 145-154.	21.4	675
7	Meta-Analysis of 28,141 Individuals Identifies Common Variants within Five New Loci That Influence Uric Acid Concentrations. PLoS Genetics, 2009, 5, e1000504.	3.5	572
8	A catalog of genetic loci associated with kidney function from analyses of a million individuals. Nature Genetics, 2019, 51, 957-972.	21.4	549
9	Genome-wide association study identifies loci influencing concentrations of liver enzymes in plasma. Nature Genetics, 2011, 43, 1131-1138.	21.4	501
10	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. Nature Neuroscience, 2018, 21, 1656-1669.	14.8	490
11	Genetic associations at 53 loci highlight cell types and biological pathways relevant for kidney function. Nature Communications, 2016, 7, 10023.	12.8	412
12	The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679.	27.8	353
13	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
14	Seventy-five genetic loci influencing the human red blood cell. Nature, 2012, 492, 369-375.	27.8	320
15	Identification of heart rate–associated loci and their effects on cardiac conduction and rhythm disorders. Nature Genetics, 2013, 45, 621-631.	21.4	282
16	The relationship between stressful life events, the serotonin transporter (5-HTTLPR) genotype and major depression. Psychological Medicine, 2005, 35, 101-111.	4.5	265
17	Genome-wide association and genetic functional studies identify <i>autism susceptibility candidate 2</i> gene ( <i>AUTS2</i> ) in the regulation of alcohol consumption. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 7119-7124.	7.1	258
18	Target genes, variants, tissues and transcriptional pathways influencing human serum urate levels. Nature Genetics, 2019, 51, 1459-1474.	21.4	251

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19	CDT, GGT, and AST As Markers of Alcohol Use: The WHO/ISBRA Collaborative Project. Alcoholism: Clinical and Experimental Research, 2002, 26, 332-339.	2.4	247
20	Large-Scale Gene-Centric Meta-analysis across 32 Studies Identifies Multiple Lipid Loci. American Journal of Human Genetics, 2012, 91, 823-838.	6.2	227
21	Common variants in TMPRSS6 are associated with iron status and erythrocyte volume. Nature Genetics, 2009, 41, 1173-1175.	21.4	226
22	Traditional markers of excessive alcohol use. Addiction, 2003, 98, 31-43.	3.3	223
23	<i>KLB</i> is associated with alcohol drinking, and its gene product β-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.	7.1	208
24	Novel loci affecting iron homeostasis and their effects in individuals at risk for hemochromatosis. Nature Communications, 2014, 5, 4926.	12.8	192
25	Associations of ADH and ALDH2 gene variation with self report alcohol reactions, consumption and dependence: an integrated analysis. Human Molecular Genetics, 2009, 18, 580-593.	2.9	187
26	A Quantitative-Trait Genome-Wide Association Study of Alcoholism Risk in the Community: Findings and Implications. Biological Psychiatry, 2011, 70, 513-518.	1.3	184
27	The Role of Adiposity in Cardiometabolic Traits: A Mendelian Randomization Analysis. PLoS Medicine, 2013, 10, e1001474.	8.4	178
28	Genome-wide association study identifies novel genetic variants contributing to variation in blood metabolite levels. Nature Communications, 2015, 6, 7208.	12.8	178
29	Directional dominance on stature and cognition inÂdiverse human populations. Nature, 2015, 523, 459-462.	27.8	173
30	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. Nature Communications, 2017, 8, 14977.	12.8	169
31	Genome-wide physical activity interactions in adiposity ― A meta-analysis of 200,452 adults. PLoS Genetics, 2017, 13, e1006528.	3.5	158
32	Variants in TF and HFE Explain â^1⁄440% of Genetic Variation in Serum-Transferrin Levels. American Journal of Human Genetics, 2009, 84, 60-65.	6.2	155
33	Genome-wide association study identifies loci affecting blood copper, selenium and zinc. Human Molecular Genetics, 2013, 22, 3998-4006.	2.9	140
34	Genetic effects on alcohol dependence risk: re-evaluating the importance of psychiatric and other heritable risk factors. Psychological Medicine, 2004, 34, 1519-1530.	4.5	132
35	Adiposity as a cause of cardiovascular disease: a Mendelian randomization study. International Journal of Epidemiology, 2015, 44, 578-586.	1.9	123
36	Alcohol Dehydrogenase and Alcohol Dependence: Variation in Genotype-Associated Risk between Populations. American Journal of Human Genetics, 2002, 71, 1247-1250.	6.2	118

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37	Genome-wide meta-analysis associates HLA-DQA1/DRB1 and LPA and lifestyle factors with human longevity. Nature Communications, 2017, 8, 910.	12.8	118
38	Serum Iron Levels and the Risk of Parkinson Disease: A Mendelian Randomization Study. PLoS Medicine, 2013, 10, e1001462.	8.4	116
39	Trans-ethnic kidney function association study reveals putative causal genes and effects on kidney-specific disease aetiologies. Nature Communications, 2019, 10, 29.	12.8	113
40	A Genomewide Association Study of Nicotine and Alcohol Dependence in Australian and Dutch Populations. Twin Research and Human Genetics, 2010, 13, 10-29.	0.6	98
41	Alcohol Consumption Indices of Genetic Risk for Alcohol Dependence. Biological Psychiatry, 2009, 66, 795-800.	1.3	88
42	Should We Use Carbohydrate-deficient Transferrin instead of γ-Glutamyltransferase for Detecting Problem Drinkers? A Systematic Review and Metaanalysis. Clinical Chemistry, 2000, 46, 1894-1902.	3.2	84
43	Mining the Human Phenome Using Allelic Scores That Index Biological Intermediates. PLoS Genetics, 2013, 9, e1003919.	3.5	84
44	Associations of autozygosity with a broad range of human phenotypes. Nature Communications, 2019, 10, 4957.	12.8	84
45	Brain structure in healthy adults is related to serum transferrin and the H63D polymorphism in the <i>HFE</i> gene. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, E851-9.	7.1	83
46	The Brisbane Systems Genetics Study: Genetical Genomics Meets Complex Trait Genetics. PLoS ONE, 2012, 7, e35430.	2.5	83
47	Novel Approach Identifies SNPs in SLC2A10 and KCNK9 with Evidence for Parent-of-Origin Effect on Body Mass Index. PLoS Genetics, 2014, 10, e1004508.	3.5	80
48	Heritability and Genome-Wide Linkage in US and Australian Twins Identify Novel Genomic Regions Controlling Chromogranin A. Circulation, 2008, 118, 247-257.	1.6	79
49	Genetic Effects on Toxic and Essential Elements in Humans: Arsenic, Cadmium, Copper, Lead, Mercury, Selenium, and Zinc in Erythrocytes. Environmental Health Perspectives, 2010, 118, 776-782.	6.0	79
50	New alcohol-related genes suggest shared genetic mechanisms with neuropsychiatric disorders. Nature Human Behaviour, 2019, 3, 950-961.	12.0	75
51	Genetic Covariation between Serum γ-Clutamyltransferase Activity and Cardiovascular Risk Factors. Clinical Chemistry, 2002, 48, 1426-1431.	3.2	74
52	ADH single nucleotide polymorphism associations with alcohol metabolism in vivo. Human Molecular Genetics, 2009, 18, 1533-1542.	2.9	74
53	Heritabilities of Apolipoprotein and Lipid Levels in Three Countries. Twin Research and Human Genetics, 2002, 5, 87-97.	1.0	72
54	The Genetics of Alcohol Intake and of Alcohol Dependence. Alcoholism: Clinical and Experimental Research, 2004, 28, 1153-1160.	2.4	71

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55	Trans-ethnic Fine Mapping Highlights Kidney-Function Genes Linked to Salt Sensitivity. American Journal of Human Genetics, 2016, 99, 636-646.	6.2	67
56	Association Between Population Density and Genetic Risk for Schizophrenia. JAMA Psychiatry, 2018, 75, 901.	11.0	67
57	GENETIC STUDY: H2 haplotype at chromosome 17q21.31 protects against childhood sexual abuseâ€associated risk for alcohol consumption and dependence. Addiction Biology, 2010, 15, 1-11.	2.6	66
58	Age- and Sex-Specific Causal Effects of Adiposity on Cardiovascular Risk Factors. Diabetes, 2015, 64, 1841-1852.	0.6	63
59	The Role of <i>GABRA2</i> in Alcohol Dependence, Smoking, and Illicit Drug Use in an Australian Population Sample. Alcoholism: Clinical and Experimental Research, 2008, 32, 1721-1731.	2.4	61
60	CDT, GGT, and AST as markers of alcohol use: the WHO/ISBRA collaborative project. Alcoholism: Clinical and Experimental Research, 2002, 26, 332-9.	2.4	61
61	Collaboration: Group theory. Nature, 2008, 455, 720-723.	27.8	60
62	Relative importance of female-specific and non-female-specific effects on variation in iron stores between women. British Journal of Haematology, 2003, 120, 860-866.	2.5	57
63	The assessment of alcoholism in surveys of the general community: What are we measuring? Some insights from the Australian twin panel interview survey. International Review of Psychiatry, 1994, 6, 295-307.	2.8	56
64	Butyrylcholinesterase: Association with the Metabolic Syndrome and Identification of 2 Gene Loci Affecting Activity. Clinical Chemistry, 2006, 52, 1014-1020.	3.2	56
65	Genomeâ€wide Association Study and Metaâ€analysis on Alcoholâ€Associated Liver Cirrhosis Identifies Genetic Risk Factors. Hepatology, 2021, 73, 1920-1931.	7.3	54
66	Association of the gastric alcohol dehydrogenase gene ADH7 with variation in alcohol metabolism. Human Molecular Genetics, 2007, 17, 179-189.	2.9	48
67	International Genome-Wide Association Study Consortium Identifies Novel Loci Associated With Blood Pressure in Children and Adolescents. Circulation: Cardiovascular Genetics, 2016, 9, 266-278.	5.1	48
68	Serum Î <sup>3</sup> -Glutamyltransferase and Risk of Disease. Clinical Chemistry, 2007, 53, 1-2.	3.2	47
69	Meta-analysis of four new genome scans for lipid parameters and analysis of positional candidates in positive linkage regions. European Journal of Human Genetics, 2005, 13, 1143-1153.	2.8	46
70	Metabolic and Biochemical Effects of Lowâ€ŧoâ€₦oderate Alcohol Consumption. Alcoholism: Clinical and Experimental Research, 2013, 37, 575-586.	2.4	46
71	GWAS of butyrylcholinesterase activity identifies four novel loci, independent effects within BCHE and secondary associations with metabolic risk factors. Human Molecular Genetics, 2011, 20, 4504-4514.	2.9	45
72	Standardisation and use of the alcohol biomarker carbohydrate-deficient transferrin (CDT). Clinica Chimica Acta, 2016, 459, 19-24.	1.1	45

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73	Genomewide Association Study of Alcohol Dependence Identifies Risk Loci Altering Ethanolâ€Response Behaviors in Model Organisms. Alcoholism: Clinical and Experimental Research, 2017, 41, 911-928.	2.4	43
74	Long-Term Stability and Heritability of Telephone Interview Measures of Alcohol Consumption and Dependence. Twin Research and Human Genetics, 2008, 11, 287-305.	0.6	42
75	Common Genetic Contributions to Alcohol and Cannabis Use and Dependence Symptomatology. Alcoholism: Clinical and Experimental Research, 2010, 34, 545-554.	2.4	42
76	Genetic covariation between serum gamma-glutamyltransferase activity and cardiovascular risk factors. Clinical Chemistry, 2002, 48, 1426-31.	3.2	42
77	Choice of Residential Location: Chance, Family Influences, or Genes?. Twin Research and Human Genetics, 2005, 8, 22-26.	0.6	41
78	COMBINING CARBOHYDRATE-DEFICIENT TRANSFERRIN AND GAMMA-GLUTAMYLTRANSFERASE TO INCREASE DIAGNOSTIC ACCURACY FOR PROBLEM DRINKING. Alcohol and Alcoholism, 2003, 38, 574-582.	1.6	40
79	Effects of Variation at the ALDH2 Locus on Alcohol Metabolism, Sensitivity, Consumption, and Dependence in Europeans. Alcoholism: Clinical and Experimental Research, 2006, 30, 1093-1100.	2.4	40
80	Toward standardization of carbohydrate-deficient transferrin (CDT) measurements: II. Performance of a laboratory network running the HPLC candidate reference measurement procedure and evaluation of a candidate reference material. Clinical Chemistry and Laboratory Medicine, 2010, 48, 1585-1592.	2.3	39
81	Measuring Carbohydrate-Deficient Transferrin by Direct Immunoassay: Factors Affecting Diagnostic Sensitivity for Excessive Alcohol Intake. Clinical Chemistry, 2008, 54, 1158-1165.	3.2	38
82	Genetic covariation of neuroticism with monoamine oxidase activity and smoking. American Journal of Medical Genetics Part A, 2001, 105, 700-706.	2.4	36
83	Understanding the role of bitter taste perception in coffee, tea and alcohol consumption through Mendelian randomization. Scientific Reports, 2018, 8, 16414.	3.3	36
84	Evidence of Genetic Effects on Blood Lead Concentration. Environmental Health Perspectives, 2007, 115, 1224-1230.	6.0	34
85	Circulating Lipids Are Associated with Alcoholic Liver Cirrhosis and Represent Potential Biomarkers for Risk Assessment. PLoS ONE, 2015, 10, e0130346.	2.5	33
86	IFCC approved HPLC reference measurement procedure for the alcohol consumption biomarker carbohydrate-deficient transferrin (CDT): Its validation and use. Clinica Chimica Acta, 2017, 465, 91-100.	1.1	33
87	Genetic comorbidity between major depression and cardioâ€metabolic traits, stratified by age at onset of major depression. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2020, 183, 309-330.	1.7	33
88	A genetic risk score and diabetes predict development of alcohol-related cirrhosis in drinkers. Journal of Hepatology, 2022, 76, 275-282.	3.7	33
89	Limitations of DSM-IV Operationalizations of Alcohol Abuse and Dependence in a Sample of Australian Twins. Twin Research and Human Genetics, 2005, 8, 574-584.	0.6	31
90	Heritability and Stability of Resting Blood Pressure in Australian Twins. Twin Research and Human Genetics, 2006, 9, 205-209.	0.6	31

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91	Genome-wide association study identifies two loci strongly affecting transferrin glycosylation. Human Molecular Genetics, 2011, 20, 3710-3717.	2.9	31
92	Hazardous alcohol consumption and other barriers to antiviral treatment among hepatitis C positive people receiving opioid maintenance treatment. Drug and Alcohol Review, 2007, 26, 231-239.	2.1	30
93	Harmonization of Measurement Results of the Alcohol Biomarker Carbohydrate-Deficient Transferrin by Use of the Toolbox of Technical Procedures of the International Consortium for Harmonization of Clinical Laboratory Results. Clinical Chemistry, 2014, 60, 945-953.	3.2	30
94	Is Alcohol-Related Flushing a Protective Factor for Alcoholism in Caucasians?. Alcoholism: Clinical and Experimental Research, 1995, 19, 582-592.	2.4	29
95	Platelet Adenylyl Cyclase Activity as a Trait Marker of Alcohol Dependence. Alcoholism: Clinical and Experimental Research, 2000, 24, 810-821.	2.4	29
96	Evidence for a QTL on chromosome 19 influencing LDL cholesterol levels in the general population. European Journal of Human Genetics, 2003, 11, 845-850.	2.8	29
97	A Genome-Wide Screen for Interactions Reveals a New Locus on 4p15 Modifying the Effect of Waist-to-Hip Ratio on Total Cholesterol. PLoS Genetics, 2011, 7, e1002333.	3.5	29
98	Brief Report: Genetics of Alcoholic Cirrhosis— <scp>G</scp> enom <scp>ALC</scp> Multinational Study. Alcoholism: Clinical and Experimental Research, 2015, 39, 836-842.	2.4	29
99	Toward standardization of carbohydrate-deficient transferrin (CDT) measurements: III. Performance of native serum and serum spiked with disialotransferrin proves that harmonization of CDT assays is possible. Clinical Chemistry and Laboratory Medicine, 2013, 51, 991-6.	2.3	28
100	Genome-wide association study of blood lead shows multiple associations near ALAD. Human Molecular Genetics, 2015, 24, 3871-3879.	2.9	28
101	Genetic insights into cardiometabolic risk factors. Clinical Biochemist Reviews, 2014, 35, 15-36.	3.3	28
102	Old insects in new order. Nature, 2002, 417, 29-29.	27.8	27
103	Genetic and Non-Genetic Factors Affecting Birth-Weight and Adult Body Mass Index. Twin Research and Human Genetics, 2001, 4, 365-370.	1.0	26
104	ADH Genotype Does Not Modify the Effects of Alcohol on High-Density Lipoprotein. Alcoholism: Clinical and Experimental Research, 2003, 27, 509-514.	2.4	26
105	Loci affecting gamma-glutamyl transferase in adults and adolescents show age × SNP interaction and cardiometabolic disease associations. Human Molecular Genetics, 2012, 21, 446-455.	2.9	26
106	Serum cholesterol and variant in cholesterol-related gene CETP predict white matter microstructure. Neurobiology of Aging, 2014, 35, 2504-2513.	3.1	26
107	Genetics of Serum Dehydroepiandrosterone Sulfate and Its Relationship to Insulin in a Population-Based Cohort of Twin Subjects. Journal of Clinical Endocrinology and Metabolism, 2002, 87, 682-686.	3.6	25
108	Obesity, Diabetes, Coffee, Tea, and Cannabis Use Alter Risk for Alcohol-Related Cirrhosis in 2 Large Cohorts of High-Risk Drinkers. American Journal of Gastroenterology, 2021, 116, 106-115.	0.4	25

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109	Genetic architecture of circulating lipid levels. European Journal of Human Genetics, 2011, 19, 813-819.	2.8	23
110	Iron and hepcidin as risk factors in atherosclerosis: what do the genes say?. BMC Genetics, 2015, 16, 79.	2.7	23
111	Serum iron level and kidney function: a Mendelian randomization study. Nephrology Dialysis Transplantation, 2016, 32, gfw215.	0.7	23
112	Autosomal linkage analysis for cannabis use behaviors in Australian adults. Drug and Alcohol Dependence, 2008, 98, 185-190.	3.2	22
113	Neuropeptide Y (NPY). Journal of the American College of Cardiology, 2012, 60, 1678-1689.	2.8	22
114	Choice of Residential Location: Chance, Family Influences, or Genes?. Twin Research and Human Genetics, 2005, 8, 22-26.	0.6	21
115	A Multivariate Assessment of Alcohol Consumption. International Journal of Epidemiology, 1981, 10, 281-288.	1.9	20
116	Longitudinal Genetic Analysis of Plasma Lipids. Twin Research and Human Genetics, 2006, 9, 550-557.	0.6	20
117	Genetic and Non-Genetic Factors Affecting Birth-Weight and Adult Body Mass Index. Twin Research and Human Genetics, 2001, 4, 365-370.	1.0	20
118	Effects of high alcohol intake, alcoholâ€related symptoms and smoking on mortality. Addiction, 2018, 113, 158-166.	3.3	19
119	Associations between polygenic risk for tobacco and alcohol use and liability to tobacco and alcohol use, and psychiatric disorders in an independent sample of 13,999 Australian adults. Drug and Alcohol Dependence, 2019, 205, 107704.	3.2	19
120	Genome-Wide Scan for Blood Pressure in Australian and Dutch Subjects Suggests Linkage at 5P, 14Q, and 17P. Hypertension, 2007, 49, 832-838.	2.7	18
121	Nosy neighbours. Nature, 2002, 419, 242-243.	27.8	17
122	The cost of leafing. Nature, 2006, 444, 539-541.	27.8	17
123	Differential and shared genetic effects on kidney function between diabetic and non-diabetic individuals. Communications Biology, 2022, 5, .	4.4	17
124	Reprint of Standardisation and use of the alcohol biomarker carbohydrate-deficient transferrin (CDT). Clinica Chimica Acta, 2017, 467, 15-20.	1.1	16
125	An indifference to boundaries. Nature, 2008, 451, 872-873.	27.8	15
126	Biomarker and Genomic Risk Factors for Liver Function Test Abnormality in Hazardous Drinkers. Alcoholism: Clinical and Experimental Research, 2019, 43, 473-482.	2.4	15

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127	Functional Relevance of Human ADH Polymorphism. Alcoholism: Clinical and Experimental Research, 2001, 25, 157S-163S.	2.4	15
128	A COMMUNITY SCREENING TEST FOR HIGH ALCOHOL CONSUMPTION USING BIOCHEMICAL AND HAEMATOLOGICAL MEASURES. Alcohol and Alcoholism, 1991, 26, 337-346.	1.6	14
129	The law of the jungle. Nature, 2003, 421, 8-9.	27.8	13
130	Can We Identify Genes For Alcohol Consumption In Samples Ascertained For Heterogeneous Purposes?. Alcoholism: Clinical and Experimental Research, 2009, 33, 729-739.	2.4	13
131	Identification of novel loci affecting circulating chromogranins and related peptides. Human Molecular Genetics, 2016, 26, ddw380.	2.9	13
132	Evaluation of laboratory tests for cirrhosis and for alcohol use, in the context of alcoholic cirrhosis. Alcohol, 2018, 66, 1-7.	1.7	13
133	Metabolomics reveals a link between homocysteine and lipid metabolism and leukocyte telomere length: the ENGAGE consortium. Scientific Reports, 2019, 9, 11623.	3.3	13
134	Pessimism is associated with greater all-cause and cardiovascular mortality, but optimism is not protective. Scientific Reports, 2020, 10, 12609.	3.3	13
135	Too hot to handle. Nature, 2003, 425, 338-339.	27.8	12
136	Linkage Analysis of Alcohol Dependence Symptoms in the Community. Alcoholism: Clinical and Experimental Research, 2010, 34, 158-163.	2.4	12
137	Blood copper and risk of cardiometabolic diseases: a Mendelian randomization study. Human Molecular Genetics, 2022, 31, 783-791.	2.9	12
138	Alcohol and gene interactions. Clinical Chemistry and Laboratory Medicine, 2005, 43, 480-7.	2.3	11
139	Association Between In Vivo Alcohol Metabolism and Genetic Variation in Pathways that Metabolize the Carbon Skeleton of Ethanol and <scp>NADH</scp> Reoxidation in the Alcohol Challenge Twin Study. Alcoholism: Clinical and Experimental Research, 2012, 36, 2074-2085.	2.4	11
140	Gut reaction. Nature, 2003, 423, 583-584.	27.8	10
141	ERYTHROCYTE ALDEHYDE DEHYDROGENASE ACTIVITY: LACK OF ASSOCIATION WITH ALCOHOL USE AND DEPENDENCE OR ALCOHOL REACTIONS IN AUSTRALIAN TWINS. Alcohol and Alcoholism, 2005, 40, 343-348.	1.6	10
142	No Genetic Overlap Between Circulating Iron Levels and Alzheimer's Disease. Journal of Alzheimer's Disease, 2017, 59, 85-99.	2.6	10
143	Investigating the relationship between iron and depression. Journal of Psychiatric Research, 2017, 94, 148-155.	3.1	10
144	Platelet Adenylyl Cyclase Activity as a Trait Marker of Alcohol Dependence. Alcoholism: Clinical and Experimental Research, 2000, 24, 810-821.	2.4	10

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145	Genetics of Serum Dehydroepiandrosterone Sulfate and Its Relationship to Insulin in a Population-Based Cohort of Twin Subjects. Journal of Clinical Endocrinology and Metabolism, 2002, 87, 682-686.	3.6	10
146	It's lifeisn't it?. Nature, 2004, 430, 288-290.	27.8	9
147	Mathematical biology centre launched. Nature, 2008, 455, 11-11.	27.8	9
148	Origin of life: Nascence man. Nature, 2009, 459, 316-319.	27.8	9
149	An assessment of the genetic relationship between alcohol metabolism and alcoholism risk in Australian twins of European ancestry. Behavior Genetics, 1999, 29, 463-472.	2.1	8
150	Fine mapping the CETP region reveals a common intronic insertion associated to HDL-C. Npj Aging and Mechanisms of Disease, 2015, 1, 15011.	4.5	8
151	Genetic Variation Within a Metabolic Motif in the Chromogranin A Promoter: Pleiotropic Influence on Cardiometabolic Risk Traits in Twins. American Journal of Hypertension, 2012, 25, 29-40.	2.0	6
152	Association and genetic overlap between clinical chemistry tests and migraine. Cephalalgia, 2021, 41, 1208-1221.	3.9	6
153	Evidence of Differential Allelic Effects between Adolescents and Adults for Plasma High-Density Lipoprotein. PLoS ONE, 2012, 7, e35605.	2.5	6
154	Birthweights in Same-sex and Opposite-sex Twin Pregnancies. Twin Research and Human Genetics, 2002, 5, 310-310.	1.0	5
155	Genome-wide linkage scan for loci influencing plasma triglycerides. Clinica Chimica Acta, 2006, 374, 87-92.	1.1	5
156	Validity of the Grossarth-Maticek and Eysenck personality-stress model of disease: An empirical prospective cohort study. Personality and Individual Differences, 2020, 157, 109797.	2.9	4
157	Base invaders. Nature, 2006, 439, 130-131.	27.8	3
158	From microscope to multiplex - An MRI scanner darkly. Nature, 2006, 441, 922-924.	27.8	3
159	Molecular biology and genetics in clinical chemistry and laboratory medicine. Clinical Chemistry and Laboratory Medicine, 2010, 48, 431-4.	2.3	3
160	Genetics and molecular biology in laboratory medicine, 1963–2013. Clinical Chemistry and Laboratory Medicine, 2013, 51, 113-117.	2.3	3
161	Comparison of Familial, Polygenic and Biochemical Predictors of Mortality. Twin Research and Human Genetics, 2020, 23, 307-315.	0.6	2
162	Transferrin Saturation and Mortality. Clinical Chemistry, 2011, 57, 921-923.	3.2	1

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163	Identifying candidate gene effects by restricting search space in a multivariate genetic analysis of white matter microstructure. , 2014, , .		1
164	Two-locus Linkage Analysis Applied to Putative Quantitative Trait Loci for Lipoprotein(a) Levels. Twin Research and Human Genetics, 2003, 6, 322-324.	1.0	1
165	Letter to the Editor. Drug and Alcohol Review, 1994, 13, 347-347.	2.1	0
166	A scientific perspective on harm reduction. Drug and Alcohol Review, 1996, 15, 117-119.	2.1	0
167	Eat me!. Nature, 2000, 406, 840-840.	27.8	0
168	Lovely grub. Nature, 2000, 408, 422-422.	27.8	0
169	Locking horns. Nature, 2002, 415, 956-956.	27.8	0
170	Corporate chiefs told to follow animal urges. Nature, 2002, 420, 724-724.	27.8	0
171	P3-010: Assessment of genetic overlap between serum iron levels and risk of Alzheimer's disease. , 2015, 11, P623-P623.		0
172	Genetics of Biochemical Phenotypes. Twin Research and Human Genetics, 2020, 23, 77-79.	0.6	0
173	Commentary: Causation versus association for fetal effects of maternal alcohol use. International Journal of Epidemiology, 2021, 49, 1995-1997.	1.9	0
174	Co-Inheritance of Variation in All-Cause Mortality and Biochemical Risk Factors. Twin Research and Human Genetics, 0, , 1-8.	0.6	0