New genetic loci link adipose and insulin biology to bod

Nature 518, 187-196

DOI: 10.1038/nature14132

Citation Report

#	Article	IF	CITATIONS
3	Multi-omic profiles of human non-alcoholic fatty liver disease tissue highlight heterogenic phenotypes. Scientific Data, 2015, 2, 150068.	2.4	48
4	Cross-species gene expression analysis identifies a novel set of genes implicated in human insulin sensitivity. Npj Systems Biology and Applications, 2015, 1, 15010.	1.4	11
5	The epigenetic signature of subcutaneous fat cells is linked to altered expression of genes implicated in lipid metabolism in obese women. Clinical Epigenetics, 2015, 7, 93.	1.8	54
6	Many obesity-associated SNPs strongly associate with DNA methylation changes at proximal promoters and enhancers. Genome Medicine, 2015, 7, 103.	3.6	124
7	The direction of cross affects obesity after puberty in male but not female offspring. BMC Genomics, 2015, 16, 904.	1.2	6
8	High visceral fat with low subcutaneous fat accumulation as a determinant of atherosclerosis in patients with type 2 diabetes. Cardiovascular Diabetology, 2015, 14, 136.	2.7	61
9	Shaping fat distribution: New insights into the molecular determinants of depot- and sex-dependent adipose biology. Obesity, 2015, 23, 1345-1352.	1.5	110
10	Evaluation of the relationship between the pelvic floor muscles and insulin resistance. Diabetes, Metabolic Syndrome and Obesity: Targets and Therapy, 2015, 8, 409.	1.1	10
11	Obesity and Insulin Resistance Are the Central Issues in Prevention of and Care for Comorbidities. Healthcare (Switzerland), 2015, 3, 408-416.	1.0	5
12	New insights from monogenic diabetes for \tilde{A} ¢â,¬Å"common \tilde{A} ¢â,¬Â•type 2 diabetes. Frontiers in Genetics, 2015, 6, 251.	1.1	29
13	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
14	Discovery and Fine-Mapping of Glycaemic and Obesity-Related Trait Loci Using High-Density Imputation. PLoS Genetics, 2015, 11, e1005230.	1.5	77
15	Genetic Regulation of Puberty Timing in Humans. Neuroendocrinology, 2015, 102, 247-255.	1.2	43
16	Mendelian Randomization Study of Body Mass Index and Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1024-1031.	1.1	67
17	Can Genetics Modify the Influence of Healthy Lifestyle on Lipids in the Context of Obesity and Type 2 Diabetes?. Current Cardiovascular Risk Reports, 2015, 9, 1.	0.8	1
18	Functional Genomics Analysis of Big Data Identifies Novel Peroxisome Proliferator–Activated Receptor γ Target Single Nucleotide Polymorphisms Showing Association With Cardiometabolic Outcomes. Circulation: Cardiovascular Genetics, 2015, 8, 842-851.	5.1	1
19	Insights into the Genetic Susceptibility to Type 2 Diabetes from Genome-Wide Association Studies of Obesity-Related Traits. Current Diabetes Reports, 2015, 15, 83.	1.7	47
20	GWAS as a Driver of Gene Discovery in Cardiometabolic Diseases. Trends in Endocrinology and Metabolism, 2015, 26, 722-732.	3.1	29

#	Article	IF	Citations
21	Invited Commentary: E Pluribus Unum for Epidemiology. American Journal of Epidemiology, 2015, 183, kwv236.	1.6	1
22	Dietary fat and carbohydrate modulate the effect of the ATP-binding cassette A1 (ABCA1) R230C variant on metabolic risk parameters in premenopausal women from the Genetics of Atherosclerotic Disease (GEA) Study. Nutrition and Metabolism, 2015, 12, 45.	1.3	14
23	Genetic studies of body mass index yield new insights for obesity biology. Nature, 2015, 518, 197-206.	13.7	3,823
24	Biological interpretation of genome-wide association studies using predicted gene functions. Nature Communications, 2015, 6, 5890.	5.8	706
25	Leveraging Functional-Annotation Data in Trans-ethnic Fine-Mapping Studies. American Journal of Human Genetics, 2015, 97, 260-271.	2.6	186
26	Calcium Homeostasis and Organelle Function in the Pathogenesis of Obesity and Diabetes. Cell Metabolism, 2015, 22, 381-397.	7.2	245
27	Genetic Variation Determines PPARÎ ³ Function and Anti-diabetic Drug Response InÂVivo. Cell, 2015, 162, 33-44.	13.5	107
28	Genome-wide Analysis of Body Proportion Classifies Height-Associated Variants by Mechanism of Action and Implicates Genes Important for Skeletal Development. American Journal of Human Genetics, 2015, 96, 695-708.	2.6	67
29	Genetics of Diabetic Nephropathy: a Long Road of Discovery. Current Diabetes Reports, 2015, 15, 41.	1.7	30
30	Plexin D1 determines body fat distribution by regulating the type V collagen microenvironment in visceral adipose tissue. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 4363-4368.	3.3	61
31	Genetic Variations in MicroRNA-Binding Sites Affect MicroRNA-Mediated Regulation of Several Genes Associated With Cardio-metabolic Phenotypes. Circulation: Cardiovascular Genetics, 2015, 8, 473-486.	5.1	57
32	Apple or Pear: Size and Shape Matter. Cell Metabolism, 2015, 21, 507-508.	7.2	29
33	Strong evidence of sexual dimorphic effect of adiposity excess on insulin sensitivity. Acta Diabetologica, 2015, 52, 991-998.	1.2	4
34	Identification of shared and unique susceptibility pathways among cancers of the lung, breast, and prostate from genome-wide association studies and tissue-specific protein interactions. Human Molecular Genetics, 2015, 24, 7406-7420.	1.4	17
35	Beige adipocytes—will they beat obesity?. Nature Reviews Endocrinology, 2015, 11, 694-696.	4.3	1
36	The Genetics of Pediatric Obesity. Trends in Endocrinology and Metabolism, 2015, 26, 711-721.	3.1	66
37	Exercise genomics—a paradigm shift is needed: a commentary: TableÂ1. British Journal of Sports Medicine, 2015, 49, 1492-1496.	3.1	54
38	Genome-wide association studies of human adiposity: Zooming in on synapses. Molecular and Cellular Endocrinology, 2015, 418, 90-100.	1.6	8

#	Article	IF	CITATIONS
39	Molecular insights into the aetiology of female reproductive ageing. Nature Reviews Endocrinology, 2015, 11, 725-734.	4.3	67
40	Association between waist circumference and gray matter volume in 2344 individuals from two adult community-based samples. Neurolmage, 2015, 122, 149-157.	2.1	90
41	Recent Progress in the Understanding of Obesity: Contributions of Genome-Wide Association Studies. Current Obesity Reports, 2015, 4, 401-410.	3.5	16
42	Childhood obesity a systems medicine approach. Frontiers in Bioscience - Landmark, 2016, 21, 1061-1075.	3.0	3
43	Metabolic signals in sleep regulation: recent insights. Nature and Science of Sleep, 2016, 8, 9.	1.4	20
44	Bile Acids, FXR, and Metabolic Effects of Bariatric Surgery. Journal of Obesity, 2016, 2016, 1-8.	1.1	39
45	Commentary: Two-sample Mendelian randomization: opportunities and challenges. International Journal of Epidemiology, 2016, 45, 908-915.	0.9	494
46	Integrative Tissue-Specific Functional Annotations in the Human Genome Provide Novel Insights on Many Complex Traits and Improve Signal Prioritization in Genome Wide Association Studies. PLoS Genetics, 2016, 12, e1005947.	1.5	94
47	Effects of Genetic Loci Associated with Central Obesity on Adipocyte Lipolysis. PLoS ONE, 2016, 11, e0153990.	1.1	19
48	Investigation of Genetic Variation Underlying Central Obesity amongst South Asians. PLoS ONE, 2016, 11, e0155478.	1.1	22
49	An Empirical Comparison of Joint and Stratified Frameworks for Studying G × E Interactions: Systolic Blood Pressure and Smoking in the CHARGE Geneâ€Lifestyle Interactions Working Group. Genetic Epidemiology, 2016, 40, 404-415.	0.6	18
50	CNV analysis and mutation screening indicate an important role for the <i>NPY4R</i> gene in human obesity. Obesity, 2016, 24, 970-976.	1.5	27
51	Genetic risk scores link body fat distribution with specific cardiometabolic profiles. Obesity, 2016, 24, 1778-1785.	1.5	2
52	Metabolic syndrome. Current Opinion in Lipidology, 2016, 27, 162-171.	1.2	58
53	Two-sample Mendelian randomization: avoiding the downsides of a powerful, widely applicable but potentially fallible technique. International Journal of Epidemiology, 2016, 45, 1717-1726.	0.9	458
54	"Non-modifiable―Risk Factors for Periodontitis and Diabetes. Current Oral Health Reports, 2016, 3, 270-281.	0.5	26
55	Archaic adaptive introgression in <i>TBX15/WARS2</i> . Molecular Biology and Evolution, 2017, 34, msw283.	3.5	101
56	Controlling the joint local false discovery rate is more powerful than meta-analysis methods in joint analysis of summary statistics from multiple genome-wide association studies. Bioinformatics, 2017, 33, 500-507.	1.8	21

#	ARTICLE	IF	CITATIONS
58	Phenotypic Characterization of GeneticallyÂLowered Human Lipoprotein(a) Levels. Journal of the American College of Cardiology, 2016, 68, 2761-2772.	1.2	186
59	Quantifying the extent to which index event biases influence large genetic association studies. Human Molecular Genetics, 2017, 26, ddw433.	1.4	40
60	Association of birth weight and the development of antipsychotic induced adiposity in individuals with treatment resistant schizophrenia. European Neuropsychopharmacology, 2016, 26, 972-978.	0.3	10
61	Genetic overlap between type 2 diabetes and depression in Swedish and Danish twin registries. Molecular Psychiatry, 2016, 21, 903-909.	4.1	50
62	Effect of in-situ annealing on the structural and optical properties of GeSn films grown by MBE. Journal of Alloys and Compounds, 2016, 684, 643-648.	2.8	8
63	Fat mass- and obesity-associated genotype, dietary intakes and anthropometric measures in European adults: the Food4Me study. British Journal of Nutrition, 2016, 115, 440-448.	1.2	22
64	Detection and interpretation of shared genetic influences on 42 human traits. Nature Genetics, 2016, 48, 709-717.	9.4	990
65	A short leucocyte telomere length is associated with development of insulin resistance. Diabetologia, 2016, 59, 1258-1265.	2.9	77
66	VEGFB/VEGFR1-Induced Expansion of Adipose Vasculature Counteracts Obesity and Related Metabolic Complications. Cell Metabolism, 2016, 23, 712-724.	7.2	180
67	Genome-Wide Association Studies (GWAS) of Adiposity. , 2016, , 91-109.		0
68	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
69	Established BMI-associated genetic variants and their prospective associations with BMI and other cardiometabolic traits: the GLACIER Study. International Journal of Obesity, 2016, 40, 1346-1352.	1.6	22
70	Cardiovascular Risk Factors and Ischemic Heart Disease. Circulation: Cardiovascular Genetics, 2016, 9, 279-286.	5.1	5
71	Genetic Evidence for a Link Between Favorable Adiposity and Lower Risk of Type 2 Diabetes, Hypertension, and Heart Disease. Diabetes, 2016, 65, 2448-2460.	0.3	122
72	Epigenomic elements analyses for promoters identify ESRRG as a new susceptibility gene for obesity-related traits. International Journal of Obesity, 2016, 40, 1170-1176.	1.6	21
73	Emotional Eating in Socially Subordinate Female Rhesus Monkeys. Developments in Primatology, 2016, , 141-158.	0.7	0
74	Sex and Gender Differences in Risk, Pathophysiology and Complications of Type 2 Diabetes Mellitus. Endocrine Reviews, 2016, 37, 278-316.	8.9	1,172
7 5	Recent progress in genetics, epigenetics and metagenomics unveils the pathophysiology of human obesity. Clinical Science, 2016, 130, 943-986.	1.8	281

#	ARTICLE	IF	CITATIONS
76	Adiposity-Dependent Regulatory Effects on Multi-tissue Transcriptomes. American Journal of Human Genetics, 2016, 99, 567-579.	2.6	26
77	Fine mapping a major obesity locus (jObes1) using a Berlin Fat Mouse × B6N advanced intercross population. International Journal of Obesity, 2016, 40, 1784-1788.	1.6	17
78	Cystatin C and Cardiovascular Disease. Journal of the American College of Cardiology, 2016, 68, 934-945.	1.2	109
79	The importance of gene–environment interactions in human obesity. Clinical Science, 2016, 130, 1571-1597.	1.8	137
80	Human Phenotypic Diversity. Current Topics in Developmental Biology, 2016, 119, 349-390.	1.0	6
81	Expression of FBN1 during adipogenesis: Relevance to the lipodystrophy phenotype in Marfan syndrome and related conditions. Molecular Genetics and Metabolism, 2016, 119, 174-185.	0.5	29
82	Heritable components of the human fecal microbiome are associated with visceral fat. Genome Biology, 2016, 17, 189.	3.8	183
83	Association Between Low-Density Lipoprotein Cholesterol–Lowering Genetic Variants and Risk of Type 2 Diabetes. JAMA - Journal of the American Medical Association, 2016, 316, 1383.	3.8	310
84	Topological analysis of metabolic networks integrating co-segregating transcriptomes and metabolomes in type 2 diabetic rat congenic series. Genome Medicine, 2016, 8, 101.	3.6	19
85	Tissue Specificity and Sex-Specific Regulatory Variation Permit the Evolution of Sex-Biased Gene Expression. American Naturalist, 2016, 188, E74-E84.	1.0	41
86	Gene set analysis for interpreting genetic studies. Human Molecular Genetics, 2016, 25, R133-R140.	1.4	12
87	Genetics of Bariatric Surgery Outcomes. Endocrinology and Metabolism Clinics of North America, 2016, 45, 623-632.	1.2	3
88	NIH working group reportâ€"using genomic information to guide weight management: From universal to precision treatment. Obesity, 2016, 24, 14-22.	1.5	96
89	Associations of genetic risk scores based on adult adiposity pathways with childhood growth and adiposity measures. BMC Genetics, 2016, 17, 120.	2.7	21
90	Genetic Risk Score Mendelian Randomization Shows that Obesity Measured as Body Mass Index, but not Waist:Hip Ratio, Is Causal for Endometrial Cancer. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 1503-1510.	1.1	64
91	Interactions between genetic variants associated with adiposity traits and soft drinks in relation to longitudinal changes in body weight and waist circumference. American Journal of Clinical Nutrition, 2016, 104, 816-826.	2.2	44
92	Genetic support for the causal role of insulin in coronary heart disease. Diabetologia, 2016, 59, 2369-2377.	2.9	14
93	Bone morphogenetic protein 2 (<i>BMP2</i>) may contribute to partition of energy storage into visceral and subcutaneous fat depots. Obesity, 2016, 24, 2092-2100.	1.5	53

#	Article	IF	CITATIONS
94	Gene and Network Analysis of Common Variants Reveals Novel Associations in Multiple Complex Diseases. Genetics, 2016, 204, 783-798.	1.2	56
95	Cardiometabolic risk loci share downstream cis- and trans-gene regulation across tissues and diseases. Science, 2016, 353, 827-830.	6.0	241
96	A principal component meta-analysis on multiple anthropometric traits identifies novel loci for body shape. Nature Communications, 2016, 7, 13357.	5.8	74
97	Habitual coffee consumption and risk of type 2 diabetes, ischemic heart disease, depression and Alzheimer's disease: a Mendelian randomization study. Scientific Reports, 2016, 6, 36500.	1.6	55
98	Epigenetic programming of adipose-derived stem cells in low birthweight individuals. Diabetologia, 2016, 59, 2664-2673.	2.9	36
99	The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. Nature Genetics, 2016, 48, 1171-1184.	9.4	362
100	Interactions between host genetics and gut microbiome in diabetes and metabolic syndrome. Molecular Metabolism, 2016, 5, 795-803.	3.0	132
101	Genome-Wide Association Study of the Modified Stumvoll Insulin Sensitivity Index Identifies <i>BCL2</i> and <i>FAM19A2</i> as Novel Insulin Sensitivity Loci. Diabetes, 2016, 65, 3200-3211.	0.3	67
102	Non-targeted metabolomics combined with genetic analyses identifies bile acid synthesis and phospholipid metabolism as being associated with incident type 2 diabetes. Diabetologia, 2016, 59, 2114-2124.	2.9	74
103	Genome-wide association studies in East Asians identify new loci for waist-hip ratio and waist circumference. Scientific Reports, 2016, 6, 17958.	1.6	58
104	Associations of Plasma FGF2 Levels and Polymorphisms in the FGF2 Gene with Obesity Phenotypes in Han Chinese Population. Scientific Reports, 2016, 6, 19868.	1.6	19
105	Identification of mutations through dominant screening for obesity using C57BL/6 substrains. Scientific Reports, 2016, 6, 32453.	1.6	9
106	Genomic Variants Associated with Resistance to High Fat Diet Induced Obesity in a Primate Model. Scientific Reports, 2016, 6, 36123.	1.6	23
107	Applications of Systems Genetics and Biology for Obesity Using Pig Models. , 2016, , 25-42.		7
108	Signatures of Archaic Adaptive Introgression in Present-Day Human Populations. Molecular Biology and Evolution, 2017, 34, msw216.	3.5	146
111	Beyond BMI: Conceptual Issues Related to Overweight and Obese Patients. Obesity Facts, 2016, 9, 193-205.	1.6	86
112	Frailty and geography: should these two factors be added to the ABCDE contemporary guide to diabetes therapy?. Diabetes/Metabolism Research and Reviews, 2016, 32, 169-175.	1.7	17
114	Fetal haemoglobin in sickle-cell disease: from genetic epidemiology to new therapeutic strategies. Lancet, The, 2016, 387, 2554-2564.	6.3	73

#	Article	IF	CITATIONS
115	Deep subcutaneous adipose tissue lipid unsaturation associates with intramyocellular lipid content. Metabolism: Clinical and Experimental, 2016, 65, 1230-1237.	1.5	9
116	Detailed phenotyping identifies genes with pleiotropic effects on body composition. BMC Genomics, 2016, 17, 224.	1.2	52
117	Role of Adiponectin in Coronary Heart Disease Risk. Circulation Research, 2016, 119, 491-499.	2.0	77
118	Numerous Genes in Loci Associated With Body Fat Distribution Are Linked to Adipose Function. Diabetes, 2016, 65, 433-437.	0.3	50
119	Rare variant associations with waist-to-hip ratio in European-American and African-American women from the NHLBI-Exome Sequencing Project. European Journal of Human Genetics, 2016, 24, 1181-1187.	1.4	5
120	Nutritional habits, lifestyle, and genetic predisposition in cardiovascular and metabolic traits in Turkish population. Nutrition, 2016, 32, 693-701.	1.1	8
121	The Framingham Heart Study — 67 years of discovery in metabolic disease. Nature Reviews Endocrinology, 2016, 12, 177-183.	4.3	48
122	Uncovering the Genetic Architectures of Quantitative Traits. Computational and Structural Biotechnology Journal, 2016, 14, 28-34.	1.9	39
123	Sex differences in the association of cord blood insulin with subcutaneous adipose tissue in neonates. International Journal of Obesity, 2016, 40, 538-542.	1.6	19
124	Genome-wide association studies suggest sex-specific loci associated with abdominal and visceral fat. International Journal of Obesity, 2016, 40, 662-674.	1.6	74
125	Perspectives in Polycystic Ovary Syndrome: From Hair to Eternity. Journal of Clinical Endocrinology and Metabolism, 2016, 101, 759-768.	1.8	71
126	Integration of summary data from GWAS and eQTL studies predicts complex trait gene targets. Nature Genetics, 2016, 48, 481-487.	9.4	1,757
127	Comprehensive analysis of schizophrenia-associated loci highlights ion channel pathways and biologically plausible candidate causal genes. Human Molecular Genetics, 2016, 25, 1247-1254.	1.4	69
128	Type 2 Diabetes Genetic Predisposition, Obesity, and All-Cause Mortality Risk in the U.S.: A Multiethnic Analysis. Diabetes Care, 2016, 39, 539-546.	4.3	38
129	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. Nature Communications, 2016, 7, 10495.	5.8	245
130	Imbalanced insulin action in chronic over nutrition: Clinical harm, molecular mechanisms, and a way forward. Atherosclerosis, 2016, 247, 225-282.	0.4	67
131	Genetics and Pathogenesis of Inflammatory Bowel Disease. Annual Review of Pathology: Mechanisms of Disease, 2016, 11, 127-148.	9.6	201
132	Genome-wide meta-analysis uncovers novel loci influencing circulating leptin levels. Nature Communications, 2016, 7, 10494.	5.8	153

#	Article	IF	CITATIONS
133	Identification of Genetic and Environmental Factors Predicting Metabolically Healthy Obesity in Children: Data From the BCAMS Study. Journal of Clinical Endocrinology and Metabolism, 2016, 101, 1816-1825.	1.8	59
134	Influence of ADRB2 Gln27Glu and ADRB3 Trp64Arg polymorphisms on body weight and body composition changes after a controlled weight-loss intervention. Applied Physiology, Nutrition and Metabolism, 2016, 41, 307-314.	0.9	16
135	Assessment of established HDL-C loci for association with HDL-C levels and type 2 diabetes in Pima Indians. Diabetologia, 2016, 59, 481-491.	2.9	16
136	A large genome-wide association study of age-related macular degeneration highlights contributions of rare and common variants. Nature Genetics, 2016, 48, 134-143.	9.4	1,167
137	Machine Learning in Genomic Medicine: A Review of Computational Problems and Data Sets. Proceedings of the IEEE, 2016, 104, 176-197.	16.4	186
138	Is there a causal role for homocysteine concentration in blood pressure? A Mendelian randomization study. American Journal of Clinical Nutrition, 2016, 103, 39-49.	2.2	35
139	Association of polymorphisms in 5-HTT (SLC6A4) and MAOA genes with measures of obesity in young adults of Portuguese origin. Archives of Physiology and Biochemistry, 2016, 122, 8-13.	1.0	18
140	Genome-wide association analysis identifies three new susceptibility loci for childhood body mass index. Human Molecular Genetics, 2016, 25, 389-403.	1.4	275
141	GM3 ganglioside and phosphatidylethanolamine-containing lipids are adipose tissue markers of insulin resistance in obese women. International Journal of Obesity, 2016, 40, 706-713.	1.6	28
142	Evidence for three genetic loci involved in both anorexia nervosa risk and variation of body mass index. Molecular Psychiatry, 2017, 22, 192-201.	4.1	63
143	Associations between body mass index-related genetic variants and adult body composition: The Fenland cohort study. International Journal of Obesity, 2017, 41, 613-619.	1.6	14
144	Translating the biology of adipokines in atherosclerosis and cardiovascular diseases: Gaps and open questions. Nutrition, Metabolism and Cardiovascular Diseases, 2017, 27, 379-395.	1.1	52
145	A cellular model for the investigation of depot specific human adipocyte biology. Adipocyte, 2017, 6, 40-55.	1.3	21
146	Molecular Profiling of Human <scp>Induced Pluripotent Stem Cell</scp> â€Derived Hypothalamic Neurones Provides Developmental Insights into Genetic Loci for Body Weight Regulation. Journal of Neuroendocrinology, 2017, 29, .	1.2	4
147	The Metabolic Syndrome in Men study: a resource for studies of metabolic and cardiovascular diseases. Journal of Lipid Research, 2017, 58, 481-493.	2.0	147
148	The most important questions in cancer research and clinical oncologyâ€"Question 2â€"5. Obesity-related cancers: more questions than answers. Chinese Journal of Cancer, 2017, 36, 18.	4.9	21
149	What have human experimental overfeeding studies taught us about adipose tissue expansion and susceptibility to obesity and metabolic complications?. International Journal of Obesity, 2017, 41, 853-865.	1.6	93
150	Store-Operated Ca 2+ Entry Controls Induction of Lipolysis and the Transcriptional Reprogramming to Lipid Metabolism. Cell Metabolism, 2017, 25, 698-712.	7.2	131

#	Article	IF	Citations
151	Rare Loss-of-Function Variants in <i>NPC1</i> Predispose to Human Obesity. Diabetes, 2017, 66, 935-947.	0.3	54
152	Genetically predicted milk consumption and bone health, ischemic heart disease and type 2 diabetes: a Mendelian randomization study. European Journal of Clinical Nutrition, 2017, 71, 1008-1012.	1.3	44
153	Genomic variants at $20p11$ associated with body fat mass in the European population. Obesity, 2017 , 25 , $757-764$.	1.5	10
154	Recent Advances in Human Genetics and Epigenetics of Adiposity: Pathway to Precision Medicine?. Gastroenterology, 2017, 152, 1695-1706.	0.6	34
155	Genetic Control of Fatty Acid \hat{l}^2 -Oxidation in Chronic Obstructive Pulmonary Disease. American Journal of Respiratory Cell and Molecular Biology, 2017, 56, 738-748.	1.4	55
156	A molecular census of arcuate hypothalamus and median eminence cell types. Nature Neuroscience, 2017, 20, 484-496.	7.1	635
157	Sexual dimorphisms in genetic loci linked to body fat distribution. Bioscience Reports, 2017, 37, .	1.1	58
158	Inflammation, metaflammation and immunometabolic disorders. Nature, 2017, 542, 177-185.	13.7	1,502
159	A Genomewide Association Study Identifies Two Sexâ€Specific Loci, at <i>SPTB</i> and <i>IZUMO3</i> , Influencing Pediatric Bone Mineral Density at Multiple Skeletal Sites. Journal of Bone and Mineral Research, 2017, 32, 1274-1281.	3.1	30
160	Reducing VEGF-B Signaling Ameliorates Renal Lipotoxicity and Protects against Diabetic Kidney Disease. Cell Metabolism, 2017, 25, 713-726.	7.2	115
161	Genetic Regulation of Adipose Gene Expression and Cardio-Metabolic Traits. American Journal of Human Genetics, 2017, 100, 428-443.	2.6	141
162	Genetic Association of Waist-to-Hip Ratio With Cardiometabolic Traits, Type 2 Diabetes, and Coronary Heart Disease. JAMA - Journal of the American Medical Association, 2017, 317, 626.	3.8	313
163	Trans-ethnic fine-mapping of genetic loci for body mass index in the diverse ancestral populations of the Population Architecture using Genomics and Epidemiology (PAGE) Study reveals evidence for multiple signals at established loci. Human Genetics, 2017, 136, 771-800.	1.8	31
165	Missing heritability: is the gap closing? An analysis of 32 complex traits in the Lifelines Cohort Study. European Journal of Human Genetics, 2017, 25, 877-885.	1.4	67
166	Flow cytometric single cell analysis reveals heterogeneity between adipose depots. Adipocyte, 2017, 6, 112-123.	1.3	26
167	Genetic Variation at the Sulfonylurea Receptor, Type 2 Diabetes, and Coronary Heart Disease. Diabetes, 2017, 66, 2310-2315.	0.3	20
168	Causal Associations of Adiposity and Body Fat Distribution With Coronary Heart Disease, Stroke Subtypes, and Type 2 Diabetes Mellitus. Circulation, 2017, 135, 2373-2388.	1.6	304
169	Differential investment in body girths by sex: Evidence from 3D photonic scanning in a Thai cohort. American Journal of Physical Anthropology, 2017, 163, 696-706.	2.1	3

#	Article	IF	Citations
170	Interaction between FTO rs9939609 and the Native American-origin ABCA1 rs9282541 affects BMI in the admixed Mexican population. BMC Medical Genetics, 2017, 18, 46.	2.1	12
171	Obesity: a chronic relapsing progressive disease process. A position statement of the World Obesity Federation. Obesity Reviews, 2017, 18, 715-723.	3.1	846
172	FGF21 Is a Sugar-Induced Hormone Associated with Sweet Intake and Preference in Humans. Cell Metabolism, 2017, 25, 1045-1053.e6.	7.2	169
173	Little Fish, Big Data: Zebrafish as a Model for Cardiovascular and Metabolic Disease. Physiological Reviews, 2017, 97, 889-938.	13.1	250
174	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. Nature Communications, 2017, 8, 14977.	5.8	169
175	The Immune Response to Implanted Materials and Devices. , 2017, , .		17
176	Multiethnic genome-wide meta-analysis of ectopic fat depots identifies loci associated with adipocyte development and differentiation. Nature Genetics, 2017, 49, 125-130.	9.4	116
177	Obesity. Nature Reviews Disease Primers, 2017, 3, 17034.	18.1	766
178	Assessing the effect of obesity-related traits on multiple myeloma using a Mendelian randomisation approach. Blood Cancer Journal, 2017, 7, e573-e573.	2.8	12
179	Genetic architecture of epigenetic and neuronal ageing rates in human brain regions. Nature Communications, 2017, 8, 15353.	5.8	92
180	The biomarker and causal roles of homoarginine in the development of cardiometabolic diseases: an observational and Mendelian randomization analysis. Scientific Reports, 2017, 7, 1130.	1.6	18
181	Reduction of HbA1c levels by fucoxanthin-enriched akamoku oil possibly involves the thrifty allele of uncoupling protein 1 ($\langle i\rangle$ UCP1 $\langle i\rangle$): a randomised controlled trial in normal-weight and obese Japanese adults. Journal of Nutritional Science, 2017, 6, e5.	0.7	37
182	Genetic Basis for Sex Differences in Obesity and Lipid Metabolism. Annual Review of Nutrition, 2017, 37, 225-245.	4.3	191
183	Genetic determination of body fat distribution and the attributive influence on metabolism. Obesity, 2017, 25, 1277-1283.	1.5	15
184	Causal Effect of Plasminogen Activator Inhibitor Type 1 on Coronary Heart Disease. Journal of the American Heart Association, 2017, 6, .	1.6	89
185	Whole-Genome Sequencing Coupled to Imputation Discovers Genetic Signals for Anthropometric Traits. American Journal of Human Genetics, 2017, 100, 865-884.	2.6	131
186	An Expanded Genome-Wide Association Study of Type 2 Diabetes in Europeans. Diabetes, 2017, 66, 2888-2902.	0.3	615
187	Adipose Tissue Biology. , 2017, , .		7

#	ARTICLE	IF	CITATIONS
188	Association analyses of East Asian individuals and trans-ancestry analyses with European individuals reveal new loci associated with cholesterol and triglyceride levels. Human Molecular Genetics, 2017, 26, 1770-1784.	1.4	135
189	Ovarian hormones and obesity. Human Reproduction Update, 2017, 23, 300-321.	5.2	229
190	Obesity-induced hypoadiponectinaemia: the opposite influences of central and peripheral fat compartments. International Journal of Epidemiology, 2017, 46, 2044-2055.	0.9	25
191	Genetic variants associated with type 2 diabetes and adiposity and risk of intracranial and abdominal aortic aneurysms. European Journal of Human Genetics, 2017, 25, 758-762.	1.4	13
192	Human Demographic History Impacts Genetic Risk Prediction across Diverse Populations. American Journal of Human Genetics, 2017, 100, 635-649.	2.6	1,120
193	Obesity, genes, and sleep habits. American Journal of Clinical Nutrition, 2017, 105, 779-780.	2.2	1
194	Genetic determinants of adiponectin regulation revealed by pregnancy. Obesity, 2017, 25, 935-944.	1.5	10
195	Genetic Mechanisms Leading to Sex Differences Across Common Diseases and Anthropometric Traits. Genetics, 2017, 205, 979-992.	1.2	64
196	Genetics of obesity: can an old dog teach us new tricks?. Diabetologia, 2017, 60, 778-783.	2.9	23
197	Genetic determinants of serum vitamin B12 and their relation to body mass index. European Journal of Epidemiology, 2017, 32, 125-134.	2.5	35
198	Genetic Predictors of ≥5% Weight Loss by Multidisciplinary Advice to Severely Obese Subjects. Journal of Nutrigenetics and Nutrigenomics, 2017, 10, 32-42.	1.8	8
199	IRS1 DNA promoter methylation and expression in human adipose tissue are related to fat distribution and metabolic traits. Scientific Reports, 2017, 7, 12369.	1.6	16
200	Local Genetic Correlation Gives Insights into the Shared Genetic Architecture of Complex Traits. American Journal of Human Genetics, 2017, 101, 737-751.	2.6	220
201	Convergence between biological, behavioural and genetic determinants of obesity. Nature Reviews Genetics, 2017, 18, 731-748.	7.7	83
202	Crosstalk between KCNK3-Mediated Ion Current and Adrenergic Signaling Regulates Adipose Thermogenesis and Obesity. Cell, 2017, 171, 836-848.e13.	13.5	69
203	New Blood Pressure–Associated Loci Identified in Meta-Analyses of 475 000 Individuals. Circulation: Cardiovascular Genetics, 2017, 10, .	5.1	48
204	The genetic underpinnings of body fat distribution. Expert Review of Endocrinology and Metabolism, 2017, 12, 417-427.	1.2	3
205	Association between gestational diabetes mellitus exposure and childhood adiposity is not substantially explained by offspring genetic risk of obesity. Diabetic Medicine, 2017, 34, 1696-1700.	1.2	11

#	Article	IF	CITATIONS
206	Population genetics of sexual conflict in the genomic era. Nature Reviews Genetics, 2017, 18, 721-730.	7.7	106
207	Mendelian randomisation analysis provides no evidence for a relationship between adult height and testicular cancer risk. Andrology, 2017, 5, 914-922.	1.9	4
208	Human genetic variation and the gut microbiome in disease. Nature Reviews Genetics, 2017, 18, 690-699.	7.7	383
209	A Mendelian Randomization Study of Metabolite Profiles, Fasting Glucose, and Type 2 Diabetes. Diabetes, 2017, 66, 2915-2926.	0.3	40
210	Foundations of Immunometabolism and Implications for Metabolic Health and Disease. Immunity, 2017, 47, 406-420.	6.6	340
211	Systems approach to the pharmacological actions of HDAC inhibitors reveals EP300 activities and convergent mechanisms of regulation in diabetes. Epigenetics, 2017, 12, 991-1003.	1.3	20
212	Catching new targets in metabolic disease with a zebrafish. Current Opinion in Pharmacology, 2017, 37, 41-50.	1.7	29
213	Evaluating the Causal Relation of ApoA-IV with Disease-Related Traits - A Bidirectional Two-sample Mendelian Randomization Study. Scientific Reports, 2017, 7, 8734.	1.6	13
214	Epidemiology in Germanyâ€"general development and personal experience. European Journal of Epidemiology, 2017, 32, 635-656.	2.5	3
215	Epigenetic Regulation of PLIN 1 in Obese Women and its Relation to Lipolysis. Scientific Reports, 2017, 7, 10152.	1.6	19
216	Thyroid function and ischemic heart disease: a Mendelian randomization study. Scientific Reports, 2017, 7, 8515.	1.6	31
217	Association of Genetic Variants Related to Serum Calcium Levels With Coronary Artery Disease and Myocardial Infarction. JAMA - Journal of the American Medical Association, 2017, 318, 371.	3.8	165
218	Genetics of Nonsyndromic Human Obesity, With Suggestions for New Studies From Work in Mouse Models., 2017,, 455-476.		0
219	Obesity and the Risk for Type 2 Diabetes. , 2017, , 677-689.		0
220	Genetic analysis in UK Biobank links insulin resistance and transendothelial migration pathways to coronary artery disease. Nature Genetics, 2017, 49, 1392-1397.	9.4	190
221	Mechanisms of Type 2 Diabetes Risk Loci. Current Diabetes Reports, 2017, 17, 72.	1.7	39
222	The Role of Obesity, Type 2 Diabetes, and Metabolic Factors in Pancreatic Cancer: A Mendelian Randomization Study. Journal of the National Cancer Institute, 2017, 109, .	3.0	185
223	Elucidating the role of plexin D1 in body fat distribution and susceptibility to metabolic disease using a zebrafish model system. Adipocyte, 2017, 6, 277-283.	1.3	7

#	Article	IF	CITATIONS
224	The role of vascular endothelial growth factor-B in metabolic homoeostasis: current evidence. Bioscience Reports, 2017, 37, .	1.1	27
225	Sport disciplines, types of sports, and waist circumference in young adulthood – a populationâ€based twin study. European Journal of Sport Science, 2017, 17, 1184-1193.	1.4	2
226	A Selection Operator for Summary Association Statistics Reveals Allelic Heterogeneity of Complex Traits. American Journal of Human Genetics, 2017, 101, 903-912.	2.6	9
227	Cellular Mechanisms Driving Sex Differences in Adipose Tissue Biology and Body Shape in Humans and Mouse Models. Advances in Experimental Medicine and Biology, 2017, 1043, 29-51.	0.8	61
228	Rationale and design of GENEiUS: a prospective observational study on the genetic and environmental determinants of body mass index evolution in Canadian undergraduate students. BMJ Open, 2017, 7, e019365.	0.8	7
229	The metabolic syndrome- associated small G protein ARL15 plays a role in adipocyte differentiation and adiponectin secretion. Scientific Reports, 2017, 7, 17593.	1.6	24
230	Novel Common Variants Associated with Obesity and Type 2 Diabetes Detected Using a cFDR Method. Scientific Reports, 2017, 7, 16397.	1.6	11
231	Meis1 effects on motor phenotypes and the sensorimotor system in mice. DMM Disease Models and Mechanisms, 2017, 10, 981-991.	1.2	25
232	Normalized periprostatic fat MRI measurements can predict prostate cancer aggressiveness in men undergoing radical prostatectomy for clinically localised disease. Scientific Reports, 2017, 7, 4630.	1.6	24
233	Identification of genetic loci associated with abdominal visceral adiposity in Korean populations. Genes and Genomics, 2017, 39, 541-548.	0.5	2
234	Aging and Caloric Restriction Research: A Biological Perspective With Translational Potential. EBioMedicine, 2017, 21, 37-44.	2.7	115
235	Identification of 15 novel risk loci for coronary artery disease and genetic risk of recurrent events, atrial fibrillation and heart failure. Scientific Reports, 2017, 7, 2761.	1.6	81
236	The genomic landscape of African populations in health and disease. Human Molecular Genetics, 2017, 26, R225-R236.	1.4	64
237	Unveiling genomic regions that underlie differences between Afec-Assaf sheep and its parental Awassi breed. Genetics Selection Evolution, 2017, 49, 19.	1.2	31
238	Genetic pleiotropy between age-related macular degeneration and 16 complex diseases and traits. Genome Medicine, 2017, 9, 29.	3.6	52
239	Association of gene coding variation and resting metabolic rate in a multi-ethnic sample of children and adults. BMC Obesity, 2017, 4, 12.	3.1	6
240	Mechanistic Pathways of Sex Differences in Cardiovascular Disease. Physiological Reviews, 2017, 97, 1-37.	13.1	444
241	Transcriptional Dynamics During Human Adipogenesis and Its Link to Adipose Morphology and Distribution. Diabetes, 2017, 66, 218-230.	0.3	27

#	Article	IF	CITATIONS
242	Dissecting the genetics of complex traits using summary association statistics. Nature Reviews Genetics, 2017, 18, 117-127.	7.7	379
243	Integrative genomic analysis implicates limited peripheral adipose storage capacity in the pathogenesis of human insulin resistance. Nature Genetics, 2017, 49, 17-26.	9.4	452
244	Appearance and Internal Aging. , 2017, , 331-340.		0
245	Genome-wide DNA promoter methylation and transcriptome analysis in human adipose tissue unravels novel candidate genes for obesity. Molecular Metabolism, 2017, 6, 86-100.	3.0	84
246	Can food addiction replace binge eating assessment in obesity clinics?. Egyptian Journal of Medical Human Genetics, 2017, 18, 181-185.	0.5	9
247	LD Hub: a centralized database and web interface to perform LD score regression that maximizes the potential of summary level GWAS data for SNP heritability and genetic correlation analysis. Bioinformatics, 2017, 33, 272-279.	1.8	822
248	A genetic link between prepregnancy body mass index, postpartum weight retention, and offspring weight in early childhood. Obesity, 2017, 25, 236-243.	1.5	14
249	Diabetes Mellitus in Developing Countries and Underserved Communities. , 2017, , .		17
250	Nutrigenomics in the modern era. Proceedings of the Nutrition Society, 2017, 76, 265-275.	0.4	65
251	Generalization and fine mapping of European ancestry-based central adiposity variants in African ancestry populations. International Journal of Obesity, 2017, 41, 324-331.	1.6	16
252	The Genetic Landscape of Renal Complications in Type 1 Diabetes. Journal of the American Society of Nephrology: JASN, 2017, 28, 557-574.	3.0	101
253	Genotype-based recall to study metabolic effects of genetic variation: a pilot study of <i>PPARG < /i>Pro12Ala carriers. Upsala Journal of Medical Sciences, 2017, 122, 234-242.</i>	0.4	5
254	Plasma levels of the anti-coagulation protein C and the risk of ischaemic heart disease. Thrombosis and Haemostasis, 2017, 117, 262-268.	1.8	7
255	Mouse Models of Human GWAS Hits for Obesity and Diabetes in the Post Genomic Era: Time for Reevaluation. Frontiers in Endocrinology, 2017, 8, 11.	1.5	9
256	Angiopoietin-2 in white adipose tissue improves metabolic homeostasis through enhanced angiogenesis. ELife, 2017, 6, .	2.8	56
257	Defective STIM-mediated store operated Ca2+ entry in hepatocytes leads to metabolic dysfunction in obesity. ELife, 2017, 6, .	2.8	46
258	Alcohol consumption and its interaction with adiposity-associated genetic variants in relation to subsequent changes in waist circumference and body weight. Nutrition Journal, 2017, 16, 51.	1.5	6
259	Genetic and epigenetic studies of adiposity and cardiometabolic disease. Genome Medicine, 2017, 9, 82.	3.6	13

#	Article	IF	CITATIONS
260	A putative causal relationship between genetically determined female body shape and posttraumatic stress disorder. Genome Medicine, 2017, 9, 99.	3.6	31
261	Role of mitochondrial dysfunction and dysregulation of Ca2+ homeostasis in the pathophysiology of insulin resistance and type 2 diabetes. Journal of Biomedical Science, 2017, 24, 70.	2.6	82
262	Metabolomics and Molecular Imaging in the Post-genomic Era., 2017,, 12-21.		0
263	Modifiable pathways in Alzheimer's disease: Mendelian randomisation analysis. BMJ: British Medical Journal, 2017, 359, j5375.	2.4	239
264	The Omics of Obesity., 0,,.		1
265	Role of prediabetes in stroke. Neuropsychiatric Disease and Treatment, 2017, Volume 13, 259-267.	1.0	25
266	Physiology and Pathophysiology of Adipose Tissue-Derived Cytokine Networks., 2017,, 33-50.		2
267	Sex and Gender Specific Aspects—From Cells to Cardiovascular Disease. , 2017, , 341-362.		1
268	FAM13A and POM121C are candidate genes for fasting insulin: functional follow-up analysis of a genome-wide association study. Diabetologia, 2018, 61, 1112-1123.	2.9	24
269	Heterogeneity of adipose tissue in development and metabolic function. Journal of Experimental Biology, 2018, 221, .	0.8	147
270	Transcriptional Regulatory Mechanisms in Adipose and Muscle Tissue Associated with Composite Glucometabolic Phenotypes. Obesity, 2018, 26, 559-569.	1.5	10
271	Childhood BMI and Adult Type 2 Diabetes, Coronary Artery Diseases, Chronic Kidney Disease, and Cardiometabolic Traits: A Mendelian Randomization Analysis. Diabetes Care, 2018, 41, 1089-1096.	4.3	95
272	The genetics of adiposity. Current Opinion in Genetics and Development, 2018, 50, 86-95.	1.5	103
273	HLA-B*07, HLA-DRB1*07, HLA-DRB1*12, and HLA-C*03:02 Strongly Associate With BMI: Data From 1.3 Million Healthy Chinese Adults. Diabetes, 2018, 67, 861-871.	0.3	9
274	Integrative analysis of omics summary data reveals putative mechanisms underlying complex traits. Nature Communications, 2018, 9, 918.	5.8	250
275	Implications of publicly available genomic data resources in searching for therapeutic targets of obesity and type 2 diabetes. Experimental and Molecular Medicine, 2018, 50, 1-13.	3.2	2
276	Sex differences in obesity, lipid metabolism, and inflammationâ€"A role for the sex chromosomes?. Molecular Metabolism, 2018, 15, 35-44.	3.0	145
277	Genomeâ€wide interaction with the insulin secretion locus <i>MTNR1B</i> reveals <i>CMIP</i> as a novel type 2 diabetes susceptibility gene in African Americans. Genetic Epidemiology, 2018, 42, 559-570.	0.6	17

#	Article	IF	CITATIONS
278	Biological Insights Into Muscular Strength: Genetic Findings in the UK Biobank. Scientific Reports, 2018, 8, 6451.	1.6	78
279	Integration of human adipocyte chromosomal interactions with adipose gene expression prioritizes obesity-related genes from GWAS. Nature Communications, 2018, 9, 1512.	5 . 8	75
280	Refining the accuracy of validated target identification through coding variant fine-mapping in type 2 diabetes. Nature Genetics, 2018, 50, 559-571.	9.4	356
281	Regulatory variants at KLF14 influence type 2 diabetes risk via a female-specific effect on adipocyte size and body composition. Nature Genetics, 2018, 50, 572-580.	9.4	143
282	The Extending Spectrum of NPC1-Related Human Disorders: From Niemann–Pick C1 Disease to Obesity. Endocrine Reviews, 2018, 39, 192-220.	8.9	32
283	A Common Allele in FGF21 Associated with Sugar Intake Is Associated with Body Shape, Lower Total Body-Fat Percentage, and Higher Blood Pressure. Cell Reports, 2018, 23, 327-336.	2.9	76
284	Causal Pathways from Blood Pressure to Larger QRS Amplitudes: a Mendelian Randomization Study. Scientific Reports, 2018, 8, 5817.	1.6	1
285	Differential effects of PCSK9 variants on risk of coronary disease and ischaemic stroke. European Heart Journal, 2018, 39, 354-359.	1.0	43
286	Genome-wide analysis of self-reported risk-taking behaviour and cross-disorder genetic correlations in the UK Biobank cohort. Translational Psychiatry, 2018, 8, 39.	2.4	57
287	The current state of <scp>GPCR</scp> â€based drug discovery to treat metabolic disease. British Journal of Pharmacology, 2018, 175, 4060-4071.	2.7	40
288	Challenges in Sex- and Gender-Centered Prevention and Management of Cardiovascular Disease: Implications of Genetic, Metabolic, and Environmental Paths. Angiology, 2018, 69, 843-853.	0.8	20
289	Prospective Studies Exploring the Possible Impact of an ID3 Polymorphism on Changes in Obesity Measures. Obesity, 2018, 26, 747-754.	1.5	1
290	A systematic analysis highlights multiple long non-coding RNAs associated with cardiometabolic disorders. Journal of Human Genetics, 2018, 63, 431-446.	1.1	17
291	Identification of Novel Potentially Pleiotropic Variants Associated With Osteoporosis and Obesity Using the cFDR Method. Journal of Clinical Endocrinology and Metabolism, 2018, 103, 125-138.	1.8	39
292	Family with sequence similarity 13, member A modulates adipocyte insulin signaling and preserves systemic metabolic homeostasis. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 1529-1534.	3.3	24
293	Causal associations between risk factors and common diseases inferred from GWAS summary data. Nature Communications, 2018, 9, 224.	5.8	629
294	Detecting Polygenic Adaptation in Admixture Graphs. Genetics, 2018, 208, 1565-1584.	1.2	101
295	Identification of genetic elements in metabolism by high-throughput mouse phenotyping. Nature Communications, 2018, 9, 288.	5.8	59

#	Article	IF	CITATIONS
296	Dysregulated lipid storage and its relationship with insulin resistance and cardiovascular risk factors in non-obese Asian patients with type 2 diabetes. Adipocyte, 2018, 7, 1-10.	1.3	28
297	Polygenic Contribution in Individuals With Early-Onset Coronary Artery Disease. Circulation Genomic and Precision Medicine, 2018, 11, e001849.	1.6	41
298	The effect of age and gender on the genetic regulation of serum 25-hydroxyvitamin D - the FIN-D2D population-based study. Journal of Steroid Biochemistry and Molecular Biology, 2018, 178, 229-233.	1.2	10
299	Multi-trait analysis of genome-wide association summary statistics using MTAG. Nature Genetics, 2018, 50, 229-237.	9.4	700
300	Toppar: an interactive browser for viewing association study results. Bioinformatics, 2018, 34, 1922-1924.	1.8	9
301	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. Nature Genetics, 2018, 50, 668-681.	9.4	2,224
302	Influence of obesity-related risk factors in the aetiology of glioma. British Journal of Cancer, 2018, 118, 1020-1027.	2.9	32
303	Role of Blood Lipids in the Development of Ischemic Stroke and its Subtypes. Stroke, 2018, 49, 820-827.	1.0	132
304	Statin-induced myopathy SLCO1B1 521T‬> C is associated with prediabetes, high body mass index and normal lipid profile in Emirati population. Diabetes Research and Clinical Practice, 2018, 139, 272-277.	1.1	5
305	Epigenome-wide association in adipose tissue from the METSIM cohort. Human Molecular Genetics, 2018, 27, 1830-1846.	1.4	38
306	Multiple genetic variations confer risks for obesity and type 2 diabetes mellitus in arab descendants from UAE. International Journal of Obesity, 2018, 42, 1345-1353.	1.6	26
307	Comprehensive review and annotation of susceptibility SNPs associated with obesityâ€related traits. Obesity Reviews, 2018, 19, 917-930.	3.1	31
308	Genetic variants and clinical relevance associated with gestational diabetes mellitus in Chinese women: a case-control study. Journal of Maternal-Fetal and Neonatal Medicine, 2018, 31, 2115-2121.	0.7	7
309	CLOCK 3111T/C genetic variant influences the daily rhythm of autonomic nervous function: relevance to body weight control. International Journal of Obesity, 2018, 42, 190-197.	1.6	8
310	Loss of BIM increases mitochondrial oxygen consumption and lipid oxidation, reduces adiposity and improves insulin sensitivity in mice. Cell Death and Differentiation, 2018, 25, 217-225.	5.0	18
311	Cohort Profile: The Oxford Biobank. International Journal of Epidemiology, 2018, 47, 21-21g.	0.9	39
312	Genome-Wide Meta-Analysis Unravels Interactions between Magnesium Homeostasis and Metabolic Phenotypes. Journal of the American Society of Nephrology: JASN, 2018, 29, 335-348.	3.0	34
313	Multiethnic Meta-Analysis Identifies <i>RAI1</i> as a Possible Obstructive Sleep Apnea–related Quantitative Trait Locus in Men. American Journal of Respiratory Cell and Molecular Biology, 2018, 58, 391-401.	1.4	65

#	Article	IF	CITATIONS
314	Biochemical and cellular properties of insulin receptor signalling. Nature Reviews Molecular Cell Biology, 2018, 19, 31-44.	16.1	486
315	Phenotypic Consequences of a Genetic Predisposition to Enhanced Nitric Oxide Signaling. Circulation, 2018, 137, 222-232.	1.6	87
316	Genetics of obesity: what genetic association studies have taught us about the biology of obesity and its complications. Lancet Diabetes and Endocrinology,the, 2018, 6, 223-236.	5 . 5	335
317	Genomeâ€Wide Study of Subcutaneous and Visceral Adipose Tissue Reveals Novel Sexâ€Specific Adiposity Loci in Mexican Americans. Obesity, 2018, 26, 202-212.	1.5	16
318	Associations of adult genetic risk scores for adiposity with childhood abdominal, liver and pericardial fat assessed by magnetic resonance imaging. International Journal of Obesity, 2018, 42, 897-904.	1.6	7
319	Big Data and medicine: a big deal?. Journal of Internal Medicine, 2018, 283, 418-429.	2.7	48
320	The genetics of obstructive sleep apnoea. Respirology, 2018, 23, 18-27.	1.3	63
321	On the origin of obesity: identifying the biological, environmental and cultural drivers of genetic risk among human populations. Obesity Reviews, 2018, 19, 121-149.	3.1	158
322	Genetic risk scores for body fat distribution attenuate weight loss in women during dietary intervention. International Journal of Obesity, 2018, 42, 370-375.	1.6	14
323	Deciphering adipose tissue heterogeneity. Annals of the New York Academy of Sciences, 2018, 1411, 5-20.	1.8	77
324	Application of the distance-based F test in an mGWAS investigating \hat{I}^2 diversity of intestinal microbiota identifies variants in <i>SLC9A8</i> (NHE8) and 3 other loci. Gut Microbes, 2018, 9, 68-75.	4.3	27
325	The impact of abdominal fat on abdominal aorta calcification measured on non-enhanced CT. Medicine (United States), 2018, 97, e13233.	0.4	13
326	Bayesian multiple logistic regression for case-control GWAS. PLoS Genetics, 2018, 14, e1007856.	1.5	28
327	Shared genetic risk contributes to type 1 and type 2 diabetes etiology. Human Molecular Genetics, 2018 ,	1.4	45
328	Exploring Coronary Artery Disease GWAs Targets With Functional Links to Immunometabolism. Frontiers in Cardiovascular Medicine, 2018, 5, 148.	1.1	10
329	A comprehensive evaluation of the genetic architecture of sudden cardiac arrest. European Heart Journal, 2018, 39, 3961-3969.	1.0	59
330	Genetics of Obesity in Diverse Populations. Current Diabetes Reports, 2018, 18, 145.	1.7	27
331	Epigenome-wide methylation differences in a group of lean and obese women – A HUNT Study. Scientific Reports, 2018, 8, 16330.	1.6	27

#	Article	IF	CITATIONS
332	GWAS identifies 14 loci for device-measured physical activity and sleep duration. Nature Communications, 2018, 9, 5257.	5.8	241
333	Association of Genetic Variants Related to Gluteofemoral vs Abdominal Fat Distribution With Type 2 Diabetes, Coronary Disease, and Cardiovascular Risk Factors. JAMA - Journal of the American Medical Association, 2018, 320, 2553.	3.8	152
334	An Enrichment Analysis for Cardiometabolic Traits Suggests Non-Random Assignment of Genes to microRNAs. International Journal of Molecular Sciences, 2018, 19, 3666.	1.8	4
335	Genetics of self-reported risk-taking behaviour, trans-ethnic consistency and relevance to brain gene expression. Translational Psychiatry, 2018, 8, 178.	2.4	29
336	Fine-mapping type 2 diabetes loci to single-variant resolution using high-density imputation and islet-specific epigenome maps. Nature Genetics, 2018, 50, 1505-1513.	9.4	1,331
337	Evaluation of the causal effects between subjective wellbeing and cardiometabolic health: mendelian randomisation study. BMJ: British Medical Journal, 2018, 362, k3788.	2.4	59
338	Epigenetic prediction of complex traits and death. Genome Biology, 2018, 19, 136.	3.8	146
339	Type 2 diabetes genetic loci informed by multi-trait associations point to disease mechanisms and subtypes: A soft clustering analysis. PLoS Medicine, 2018, 15, e1002654.	3.9	373
340	Phenotype-Specific Enrichment of Mendelian Disorder Genes near GWAS Regions across 62 Complex Traits. American Journal of Human Genetics, 2018, 103, 535-552.	2.6	90
341	Opposite Genetic Effects of CMIP Polymorphisms on the Risk of Type 2 Diabetes and Obesity: A Family-Based Study in China. International Journal of Molecular Sciences, 2018, 19, 1011.	1.8	12
342	Adiposity Is a Key Correlate of Circulating Fibroblast Growth Factor-21 Levels in African Males with or without Type 2 Diabetes Mellitus. Journal of Obesity, 2018, 2018, 1-8.	1.1	9
343	Birthweight, Type 2 Diabetes Mellitus, and Cardiovascular Disease. Circulation Genomic and Precision Medicine, 2018, 11, e002054.	1.6	96
344	Body composition and insulin resistance in children. European Journal of Clinical Nutrition, 2018, 72, 1239-1245.	1.3	62
345	Genetics of obesity and its measures in India. Journal of Genetics, 2018, 97, 1047-1071.	0.4	6
346	Association of Genetically Enhanced Lipoprotein Lipase–Mediated Lipolysis and Low-Density Lipoprotein Cholesterol–Lowering Alleles With Risk of Coronary Disease and Type 2 Diabetes. JAMA Cardiology, 2018, 3, 957.	3.0	55
347	Smoking modifies the effect of two independent SNPs rs5063 and rs198358 of NPPA on central obesity in the Chinese Han population. Journal of Genetics, 2018, 97, 987-994.	0.4	3
348	The role of eating behavior traits in mediating genetic susceptibility to obesity. American Journal of Clinical Nutrition, 2018, 108, 445-452.	2.2	39
349	Genome-wide association study of response to tumour necrosis factor inhibitor therapy in rheumatoid arthritis. Pharmacogenomics Journal, 2018, 18, 657-664.	0.9	41

#	Article	IF	CITATIONS
350	PhenoSpD: an integrated toolkit for phenotypic correlation estimation and multiple testing correction using GWAS summary statistics. GigaScience, 2018, 7, .	3.3	46
351	Identification of Novel Candidate Markers of Type 2 Diabetes and Obesity in Russia by Exome Sequencing with a Limited Sample Size. Genes, 2018, 9, 415.	1.0	22
352	Perceived neighborhood social cohesion and cardiometabolic risk: a geneÂ×Âenvironment study. Biodemography and Social Biology, 2018, 64, 173-186.	0.4	1
353	Deep phenotyping in zebrafish reveals genetic and diet-induced adiposity changes that may inform disease risk. Journal of Lipid Research, 2018, 59, 1536-1545.	2.0	13
354	Sex differences in lipid and lipoprotein metabolism. Molecular Metabolism, 2018, 15, 45-55.	3.0	286
355	Role of obesity in smoking behaviour: Mendelian randomisation study in UK Biobank. BMJ: British Medical Journal, 2018, 361, k1767.	2.4	122
356	Physical activity modifies genetic susceptibility to obesity in postmenopausal women. Menopause, 2018, 25, 1131-1137.	0.8	7
357	A joint view on genetic variants for adiposity differentiates subtypes with distinct metabolic implications. Nature Communications, 2018, 9, 1946.	5.8	33
358	Method to estimate the approximate samples size that yield a certain number of significant GWAS signals in polygenic traits. Genetic Epidemiology, 2018, 42, 488-496.	0.6	2
359	The Carbohydrate-Insulin Model of Obesity Is Difficult to Reconcile With Current Evidence. JAMA Internal Medicine, 2018, 178, 1103.	2.6	44
360	The transcription factor Rfx7 limits metabolism of NK cells and promotes their maintenance and immunity. Nature Immunology, 2018, 19, 809-820.	7.0	42
361	Causal Inference in Cancer Epidemiology: What Is the Role of Mendelian Randomization?. Cancer Epidemiology Biomarkers and Prevention, 2018, 27, 995-1010.	1.1	109
362	Genetics of Bone Fat and Energy Regulation. , 2018, , 301-315.		0
363	Identification of novel high-impact recessively inherited type 2 diabetes risk variants in the Greenlandic population. Diabetologia, 2018, 61, 2005-2015.	2.9	14
364	Mendelian randomisation in type 2 diabetes and coronary artery disease. Current Opinion in Genetics and Development, 2018, 50, 111-120.	1.5	13
365	Impaired Adipogenesis and Dysfunctional Adipose Tissue in Human Hypertrophic Obesity. Physiological Reviews, 2018, 98, 1911-1941.	13.1	285
366	Maternal central obesity and birth size: a Mendelian randomization analysis. Lipids in Health and Disease, 2018, 17, 181.	1.2	13
367	DNA methylation of <i>SSPN</i> is linked to adipose tissue distribution and glucose metabolism. FASEB Journal, 2018, 32, 6898-6910.	0.2	6

#	Article	IF	Citations
368	FKBP5 expression in human adipose tissue: potential role in glucose and lipid metabolism, adipogenesis and type 2 diabetes. Endocrine, 2018, 62, 116-128.	1.1	63
369	Serum magnesium levels and risk of coronary artery disease: Mendelian randomisation study. BMC Medicine, 2018, 16, 68.	2.3	36
370	Genome-Wide Association Studies and Risk Scores for Coronary Artery Disease: Sex Biases. Advances in Experimental Medicine and Biology, 2018, 1065, 627-642.	0.8	13
371	Adiponectin and coronary artery disease risk: A bi-directional Mendelian randomization study. International Journal of Cardiology, 2018, 268, 222-226.	0.8	24
372	Empirical Bayesian approach to testing multiple hypotheses with separate priors for left and right alternatives. Statistical Applications in Genetics and Molecular Biology, $2018,17,$.	0.2	0
373	Genetics of Obesity Traits: A Bivariate Genome-Wide Association Analysis. Frontiers in Genetics, 2018, 9, 179.	1.1	29
374	Reading Mendelian randomisation studies: a guide, glossary, and checklist for clinicians. BMJ: British Medical Journal, 2018, 362, k601.	2.4	1,880
375	A coding and non-coding transcriptomic perspective on the genomics of human metabolic disease. Nucleic Acids Research, 2018, 46, 7772-7792.	6.5	41
376	Identification and functional analysis of glycemic trait loci in the China Health and Nutrition Survey. PLoS Genetics, 2018, 14, e1007275.	1.5	30
377	A fine-mapping study of central obesity loci incorporating functional annotation and imputation. European Journal of Human Genetics, 2018, 26, 1369-1377.	1.4	4
378	Admixture mapping and fine-mapping of birth weight loci in the Black Women's Health Study. Human Genetics, 2018, 137, 535-542.	1.8	6
379	Trends in Obesity and Risk of Cardiovascular Disease. Current Epidemiology Reports, 2018, 5, 243-251.	1.1	36
380	Addendum: A joint view on genetic variants for adiposity differentiates subtypes with distinct metabolic implications. Nature Communications, 2018, 9, 2861.	5.8	16
381	Biological clocks and physical functioning in monozygotic female twins. BMC Geriatrics, 2018, 18, 83.	1.1	22
382	Waist-hip ratio related genetic loci are associated with risk of impaired fasting glucose in Chinese children: a case control study. Nutrition and Metabolism, 2018, 15, 34.	1.3	6
383	Assessing causal estimates of the association of obesity-related traits with coronary artery disease using a Mendelian randomization approach. Scientific Reports, 2018, 8, 7146.	1.6	11
384	Circulating vitamin E levels and Alzheimer's disease: a Mendelian randomization study. Neurobiology of Aging, 2018, 72, 189.e1-189.e9.	1.5	53
385	Identification of nine new susceptibility loci for endometrial cancer. Nature Communications, 2018, 9, 3166.	5.8	178

#	Article	IF	CITATIONS
386	Genes that make you fat, but keep you healthy. Journal of Internal Medicine, 2018, 284, 450-463.	2.7	48
387	Identification of susceptibility pathways for the role of chromosome 15q25.1 in modifying lung cancer risk. Nature Communications, 2018, 9, 3221.	5.8	60
388	A Large Multiethnic Genome-Wide Association Study of Adult Body Mass Index Identifies Novel Loci. Genetics, 2018, 210, 499-515.	1,2	131
389	Genetic association between adiposity and gout: a Mendelian randomization study. Rheumatology, 2018, 57, 2145-2148.	0.9	56
390	Are serum concentrations of vitamin B-12 causally related to cardiometabolic risk factors and disease? A Mendelian randomization study. American Journal of Clinical Nutrition, 2018, 108, 398-404.	2.2	22
391	AMPK activation negatively regulates GDAP1, which influences metabolic processes and circadian gene expression in skeletal muscle. Molecular Metabolism, 2018, 16, 12-23.	3.0	17
392	Zebrafish as a Model for Obesity and Diabetes. Frontiers in Cell and Developmental Biology, 2018, 6, 91.	1.8	175
393	<i>BDNF</i> : mRNA expression in urine cells of patients with chronic kidney disease and its role in kidney function. Journal of Cellular and Molecular Medicine, 2018, 22, 5265-5277.	1.6	30
394	Cohort Profile: Genetics of Diabetes Audit and Research in Tayside Scotland (GoDARTS). International Journal of Epidemiology, 2018, 47, 380-381j.	0.9	59
395	Genomic insights into the causes of type 2 diabetes. Lancet, The, 2018, 391, 2463-2474.	6. 3	110
396	Genetically driven adiposity traits increase the risk of coronary artery disease independent of blood pressure, dyslipidaemia, glycaemic traits. European Journal of Human Genetics, 2018, 26, 1547-1553.	1.4	8
397	Human Genetics of Obesity and Type 2 Diabetes Mellitus. Circulation Genomic and Precision Medicine, 2018, 11, e002090.	1.6	58
398	Genome-wide association study of habitual physical activity in over 377,000 UK Biobank participants identifies multiple variants including CADM2 and APOE. International Journal of Obesity, 2018, 42, 1161-1176.	1.6	249
399	Determinants of body fat distribution in humans may provide insight about obesity-related health risks. Journal of Lipid Research, 2019, 60, 1710-1719.	2.0	132
400	A Geometric Perspective on the Power of Principal Component Association Tests in Multiple Phenotype Studies. Journal of the American Statistical Association, 2019, 114, 975-990.	1.8	26
401	Identification of Dietary Patterns Associated with Incidence of Hyperglycemia in Middle-Aged and Older Korean Adults. Nutrients, 2019, 11, 1801.	1.7	16
403	Expression analysis of CEBPA and its antisense RNA revealed their dysregulation in peripheral blood of coronary artery disease patients. Gene Reports, 2019, 16, 100466.	0.4	0
404	Lipidomics, Atrial Conduction, and Body Mass Index. Circulation Genomic and Precision Medicine, 2019, 12, e002384.	1.6	9

#	Article	IF	Citations
405	Aldehyde Dehydrogenases Genetic Polymorphism and Obesity: From Genomics to Behavior and Health. Advances in Experimental Medicine and Biology, 2019, 1193, 135-154.	0.8	1
406	Race–ethnic differences in the associations of maternal lipid trait genetic risk scores with longitudinal fetal growth. Journal of Clinical Lipidology, 2019, 13, 821-831.	0.6	8
407	Conservation, acquisition, and functional impact of sex-biased gene expression in mammals. Science, 2019, 365, .	6.0	152
408	Genetic Association Study of Eight Steroid Hormones and Implications for Sexual Dimorphism of Coronary Artery Disease. Journal of Clinical Endocrinology and Metabolism, 2019, 104, 5008-5023.	1.8	37
409	Genetic variants influencing obesity-related traits in Japanese population. Annals of Human Biology, 2019, 46, 298-304.	0.4	7
410	Genetic mapping of cell type specificity for complex traits. Nature Communications, 2019, 10, 3222.	5.8	212
411	Genetic predisposition to higher body fat yet lower cardiometabolic risk in children and adolescents. International Journal of Obesity, 2019, 43, 2007-2016.	1.6	5
412	Bone morphogenetic protein 2 is a depot-specific regulator of human adipogenesis. International Journal of Obesity, 2019, 43, 2458-2468.	1.6	21
413	Novel Genetic Locus of Visceral Fat and Systemic Inflammation. Journal of Clinical Endocrinology and Metabolism, 2019, 104, 3735-3742.	1.8	11
414	Gene–Environment Interactions on Body Fat Distribution. International Journal of Molecular Sciences, 2019, 20, 3690.	1.8	29
416	A Phenome-Wide Mendelian Randomization Study of Pancreatic Cancer Using Summary Genetic Data. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 2070-2078.	1.1	24
417	Methylome and transcriptome maps of human visceral and subcutaneous adipocytes reveal key epigenetic differences at developmental genes. Scientific Reports, 2019, 9, 9511.	1.6	24
418	Body Shape and Alzheimer's Disease: A Mendelian Randomization Analysis. Frontiers in Neuroscience, 2019, 13, 1084.	1.4	13
419	Genome-Wide Assessment for RestingÂHeart Rate and Shared Genetics WithÂCardiometabolic Traits and Type 2 Diabetes. Journal of the American College of Cardiology, 2019, 74, 2162-2174.	1.2	28
420	Circulating vitamin E and cardiometabolic measures: a Mendelian randomization analysis. Journal of Clinical Biochemistry and Nutrition, 2019, 65, 160-169.	0.6	3
421	Bayesian multivariate reanalysis of large genetic studies identifies many new associations. PLoS Genetics, 2019, 15, e1008431.	1.5	14
422	The genetic underpinnings of obesity. Current Opinion in Physiology, 2019, 12, 57-64.	0.9	0
423	Sex differences in cardiometabolic disorders. Nature Medicine, 2019, 25, 1657-1666.	15.2	244

#	Article	IF	Citations
424	Colocalization of GWAS and eQTL signals at loci with multiple signals identifies additional candidate genes for body fat distribution. Human Molecular Genetics, 2019, 28, 4161-4172.	1.4	41
425	Disparity in Adiposity among Adults with Normal Body Mass Index and Waist-to-Height Ratio. IScience, 2019, 21, 612-623.	1.9	8
426	Multi-ancestry sleep-by-SNP interaction analysis in 126,926 individuals reveals lipid loci stratified by sleep duration. Nature Communications, 2019, 10, 5121.	5.8	62
427	Causal relationships between obesity and the leading causes of death in women and men. PLoS Genetics, 2019, 15, e1008405.	1.5	113
428	HOPS: a quantitative score reveals pervasive horizontal pleiotropy in human genetic variation is driven by extreme polygenicity of human traits and diseases. Genome Biology, 2019, 20, 222.	3.8	47
429	Genetic risk score predicts risk for overweight and obesity in Finnish preadolescents. Clinical Obesity, 2019, 9, e12342.	1.1	10
430	Abdominal adiposity and cardiometabolic risk factors in children and adolescents: a Mendelian randomization analysis. American Journal of Clinical Nutrition, 2019, 110, 1079-1087.	2.2	22
431	Genome-wide association analysis of 95 549 individuals identifies novel loci and genes influencing optic disc morphology. Human Molecular Genetics, 2019, 28, 3680-3690.	1.4	19
432	Whole Genome Analyses of Chinese Population and De Novo Assembly of A Northern Han Genome. Genomics, Proteomics and Bioinformatics, 2019, 17, 229-247.	3.0	42
433	Metabolic and Endocrine Consequences of Bariatric Surgery. Frontiers in Endocrinology, 2019, 10, 626.	1.5	62
434	Detailed Functional Characterization of a Waist-Hip Ratio Locus in 7p15.2 Defines an Enhancer Controlling Adipocyte Differentiation. IScience, 2019, 20, 42-59.	1.9	6
435	Genetic Basis of Obesity and Type 2 Diabetes in Africans: Impact on Precision Medicine. Current Diabetes Reports, 2019, 19, 105.	1.7	9
436	The Many Faces of Obesity and Its Influence on Breast Cancer Risk. Frontiers in Oncology, 2019, 9, 765.	1.3	56
437	Adipose Tissue Gene Expression Associations Reveal Hundreds of Candidate Genes for Cardiometabolic Traits. American Journal of Human Genetics, 2019, 105, 773-787.	2.6	45
438	Large-Scale "OMICS―Studies to Explore the Physiopatholgy of HIV-1 Infection. Frontiers in Genetics, 2019, 10, 799.	1.1	8
439	A Survey of Gene Prioritization Tools for Mendelian and Complex Human Diseases. Journal of Integrative Bioinformatics, 2019, 16, .	1.0	25
440	Genome-wide association study of body fat distribution identifies adiposity loci and sex-specific genetic effects. Nature Communications, 2019, 10, 339.	5.8	163
441	Sex-specific moderation by lifestyle and psychosocial factors on the genetic contributions to adiposity in 112,151 individuals from UK Biobank. Scientific Reports, 2019, 9, 363.	1.6	6

#	Article	IF	Citations
442	Identification of 12 genetic loci associated with human healthspan. Communications Biology, 2019, 2, 41.	2.0	89
443	Overarching Key Issues for Feeding, Eating, and Weight Disorders. , 2019, , 1-7.		0
444	A fully adjusted twoâ€stage procedure for rankâ€normalization in genetic association studies. Genetic Epidemiology, 2019, 43, 263-275.	0.6	60
445	The Association Between Adiposity and Inpatient Hospital Costs in the UK Biobank Cohort. Applied Health Economics and Health Policy, 2019, 17, 359-370.	1.0	16
446	Genetic architecture of human thinness compared to severe obesity. PLoS Genetics, 2019, 15, e1007603.	1.5	98
447	Genetics of Central Obesity and Body Fat. , 2019, , 153-174.		2
448	Comprehensive and Systematic Analysis of Gene Expression Patterns Associated with Body Mass Index. Scientific Reports, 2019, 9, 7447.	1.6	13
449	Open Chromatin Profiling in Adipose Tissue Marks Genomic Regions with Functional Roles in Cardiometabolic Traits. G3: Genes, Genomes, Genetics, 2019, 9, 2521-2533.	0.8	19
450	An evaluation of noncoding genome annotation tools through enrichment analysis of 15 genome-wide association studies. Briefings in Bioinformatics, 2019, 20, 995-1003.	3.2	3
451	Regional fat depot masses are influenced by protein-coding gene variants. PLoS ONE, 2019, 14, e0217644.	1.1	9
452	Role of MEIS1 in restless legs syndrome: From GWAS to functional studies in mice. Advances in Pharmacology, 2019, 84, 175-184.	1.2	21
453	Associations Between Measures of Sarcopenic Obesity and Risk of Cardiovascular Disease and Mortality: A Cohort Study and Mendelian Randomization Analysis Using the UK Biobank. Journal of the American Heart Association, 2019, 8, e011638.	1.6	75
454	Genome-wide analysis of dental caries and periodontitis combining clinical and self-reported data. Nature Communications, 2019, 10, 2773.	5.8	183
455	Sexual Dimorphism of a Genetic Risk Score for Obesity and Related Traits among Chinese Patients with Type 2 Diabetes. Obesity Facts, 2019, 12, 328-343.	1.6	7
456	Authors' response: Associations of obesity and circulating insulin and glucose with breast cancer risk. International Journal of Epidemiology, 2019, 48, 1016-1017.	0.9	1
457	Geographic Variation and Bias in the Polygenic Scores of Complex Diseases and Traits in Finland. American Journal of Human Genetics, 2019, 104, 1169-1181.	2.6	90
458	Bivariate causal mixture model quantifies polygenic overlap between complex traits beyond genetic correlation. Nature Communications, 2019, 10, 2417.	5.8	190
459	MicroRNA-196a links human body fat distribution to adipose tissue extracellular matrix composition. EBioMedicine, 2019, 44, 467-475.	2.7	22

#	Article	IF	Citations
460	Assessment of Causal Direction Between Gut Microbiota–Dependent Metabolites and Cardiometabolic Health: A Bidirectional Mendelian Randomization Analysis. Diabetes, 2019, 68, 1747-1755.	0.3	114
461	Transcultural Diabetes Care in The United States – A Position Statement by the American Association of Clinical Endocrinologists. Endocrine Practice, 2019, 25, 729-765.	1.1	19
462	Adipocyte Metabolism and Insulin Signaling Perturbations: Insights from Genetics. Trends in Endocrinology and Metabolism, 2019, 30, 396-406.	3.1	17
463	Benefits and limitations of genome-wide association studies. Nature Reviews Genetics, 2019, 20, 467-484.	7.7	1,226
464	Genetic variation in CADM2 as a link between psychological traits and obesity. Scientific Reports, 2019, 9, 7339.	1.6	45
465	Mendelian randomization evaluation of causal effects of fibrinogen on incident coronary heart disease. PLoS ONE, 2019, 14, e0216222.	1.1	17
466	Associations of variants In the hexokinase 1 and interleukin 18 receptor regions with oxyhemoglobin saturation during sleep. PLoS Genetics, 2019, 15, e1007739.	1.5	28
467	Genetic Determinants of Weight Loss After Bariatric Surgery. Obesity Surgery, 2019, 29, 2554-2561.	1.1	17
468	GWAS of bone size yields twelve loci that also affect height, BMD, osteoarthritis or fractures. Nature Communications, 2019, 10, 2054.	5.8	74
469	Genetic Predisposition Impacts Clinical Changes in a Lifestyle Coaching Program. Scientific Reports, 2019, 9, 6805.	1.6	48
470	Nuclear Receptors. Methods in Molecular Biology, 2019, , .	0.4	1
471	High-Dimensional Data Approaches to Understanding Nuclear Hormone Receptor Signaling. Methods in Molecular Biology, 2019, 1966, 291-311.	0.4	1
472	Shared genetic underpinnings of childhood obesity and adult cardiometabolic diseases. Human Genomics, 2019, 13, 17.	1.4	17
473	Low-Carb and Ketogenic Diets in Type 1 and Type 2 Diabetes. Nutrients, 2019, 11, 962.	1.7	129
474	Gene Lifestyle Interactions With Relation to Obesity, Cardiometabolic, and Cardiovascular Traits Among South Asians. Frontiers in Endocrinology, 2019, 10, 221.	1.5	15
475	GWAS of smoking behaviour in 165,436 Japanese people reveals seven new loci and shared genetic architecture. Nature Human Behaviour, 2019, 3, 471-477.	6.2	54
476	Metabolomic correlates of central adiposity and earlier-life body mass index. Journal of Lipid Research, 2019, 60, 1136-1143.	2.0	2
477	Increased Diet Quality is Associated with Long†Term Reduction of Abdominal and Pericardial Fat. Obesity, 2019, 27, 670-677.	1.5	13

#	Article	IF	CITATIONS
478	<i>deTS</i> : tissue-specific enrichment analysis to decode tissue specificity. Bioinformatics, 2019, 35, 3842-3845.	1.8	51
479	The Genetic Basis of Metabolic Disease. Cell, 2019, 177, 146-161.	13.5	104
480	Primers on nutrigenetics and nutri(epi)genomics: Origins and development of precision nutrition. Biochimie, 2019, 160, 156-171.	1.3	58
481	Genetic overlap between birthweight and adult cardiometabolic diseases has implications for genomic medicine. Scientific Reports, 2019, 9, 4076.	1.6	5
482	Discovering metabolic disease gene interactions by correlated effects on cellular morphology. Molecular Metabolism, 2019, 24, 108-119.	3.0	13
483	Influence of obesity on surgical complications of patients with ovarian tumors. Oncology Letters, 2019, 17, 4590-4594.	0.8	3
484	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.	9.4	112
485	Identification of a 1p21 independent functional variant for abdominal obesity. International Journal of Obesity, 2019, 43, 2480-2490.	1.6	5
486	Genome-wide association study reveals sex-specific genetic architecture of facial attractiveness. PLoS Genetics, 2019, 15, e1007973.	1.5	5
487	Dairy Product Intake and Risk of Type 2 Diabetes in EPIC-InterAct: A Mendelian Randomization Study. Diabetes Care, 2019, 42, 568-575.	4.3	29
488	Evaluation of GDF15 as a therapeutic target of cardiometabolic diseases in human: A Mendelian randomization study. EBioMedicine, 2019, 41, 85-90.	2.7	33
489	Genetic Architecture of Human Obesity Traits in the Rhesus Macaque. Obesity, 2019, 27, 479-488.	1.5	1
490	Genetic basis of motoric cognitive risk syndrome in the Health and Retirement Study. Neurology, 2019, 92, e1427-e1434.	1.5	23
491	Shared genetic architecture between metabolic traits and Alzheimer's disease: a large-scale genome-wide cross-trait analysis. Human Genetics, 2019, 138, 271-285.	1.8	52
492	Obesity and Bone Health Revisited: A Mendelian Randomization Study for Koreans. Journal of Bone and Mineral Research, 2019, 34, 1058-1067.	3.1	26
493	Protein-coding variants implicate novel genes related to lipid homeostasis contributing to body-fat distribution. Nature Genetics, 2019, 51, 452-469.	9.4	89
494	Causal relationships among the gut microbiome, short-chain fatty acids and metabolic diseases. Nature Genetics, 2019, 51, 600-605.	9.4	854
495	The effect of liver enzymes on adiposity: a Mendelian randomization study. Scientific Reports, 2019, 9, 16792.	1.6	4

#	Article	IF	CITATIONS
496	Childhood obesity leads to adult type 2 diabetes and coronary artery diseases. Medicine (United) Tj ETQq0 0 0 rg	BT/Qverlo	ock 10 Tf 50 7
497	Epigenome-Wide Association Study of Incident Type 2 Diabetes in a British Population: EPIC-Norfolk Study. Diabetes, 2019, 68, 2315-2326.	0.3	77
498	The integrative biology of type 2 diabetes. Nature, 2019, 576, 51-60.	13.7	621
499	Recent developments in lipodystrophy. Current Opinion in Lipidology, 2019, 30, 284-290.	1.2	14
500	PCA-based GRS analysis enhances the effectiveness for genetic correlation detection. Briefings in Bioinformatics, 2019, 20, 2291-2298.	3.2	6
501	Diagnosis of obesity and use of obesity biomarkers in science and clinical medicine. Metabolism: Clinical and Experimental, 2019, 92, 61-70.	1.5	170
502	Impact of Genes and Environment on Obesity and Cardiovascular Disease. Endocrinology, 2019, 160, 81-100.	1.4	31
503	Differential expression of miRNAs related to angiogenesis and adipogenesis in subcutaneous fat of obese and nonobese women. Molecular Biology Reports, 2019, 46, 965-973.	1.0	6
504	Associations of Mitochondrial and Nuclear Mitochondrial Variants and Genes with Seven Metabolic Traits. American Journal of Human Genetics, 2019, 104, 112-138.	2.6	106
505	The influence of obesity-related factors in the etiology of renal cell carcinomaâ€"A mendelian randomization study. PLoS Medicine, 2019, 16, e1002724.	3.9	59
506	DNA Sequence Variation in <i>ACVR1C</i> Encoding the Activin Receptor-Like Kinase 7 Influences Body Fat Distribution and Protects Against Type 2 Diabetes. Diabetes, 2019, 68, 226-234.	0.3	31
507	Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder. Nature Genetics, 2019, 51, 63-75.	9.4	1,594
508	A linear mixed-model approach to study multivariate gene–environment interactions. Nature Genetics, 2019, 51, 180-186.	9.4	112
509	A genetic variant in LINGO2 contributes to the risk of gestational diabetes mellitus in a Chinese population. Journal of Cellular Physiology, 2019, 234, 7012-7018.	2.0	6
510	Association of Methylation Signatures at Hepatocellular Carcinoma Pathway Genes with Adiposity and Insulin Resistance Phenotypes. Nutrition and Cancer, 2019, 71, 840-851.	0.9	10
511	Furthering Precision Medicine Genomics With Healthy Living Medicine. Progress in Cardiovascular Diseases, 2019, 62, 60-67.	1.6	7
512	Measured maternal prepregnancy anthropometry and newborn DNA methylation. Epigenomics, 2019, 11, 187-198.	1.0	14
513	Epigenetic Regulation of Fat Deposition: A Focus on Krüppel-Like Factor 14 (Klf14). , 2019, , 351-367.		0

#	Article	IF	CITATIONS
514	Functionally oriented analysis of cardiometabolic traits in a trans-ethnic sample. Human Molecular Genetics, 2019, 28, 1212-1224.	1.4	12
515	A low visceral fat proportion, independent of total body fat mass, protects obese adolescent girls against fatty liver and glucose dysregulation: a longitudinal study. International Journal of Obesity, 2019, 43, 673-682.	1.6	30
516	Genome-Wide Meta-analysis of Gene–Environmental Interaction for Insulin Resistance Phenotypes and Breast Cancer Risk in Postmenopausal Women. Cancer Prevention Research, 2019, 12, 31-42.	0.7	15
517	Generalizing polygenic risk scores from Europeans to Hispanics/Latinos. Genetic Epidemiology, 2019, 43, 50-62.	0.6	89
518	Established and emerging strategies to crack the genetic code of obesity. Obesity Reviews, 2019, 20, 212-240.	3.1	21
519	Genome-Wide and Abdominal MRI Data Provide Evidence That a Genetically Determined Favorable Adiposity Phenotype Is Characterized by Lower Ectopic Liver Fat and Lower Risk of Type 2 Diabetes, Heart Disease, and Hypertension. Diabetes, 2019, 68, 207-219.	0.3	72
520	Genetics and epigenetics in obesity. Metabolism: Clinical and Experimental, 2019, 92, 37-50.	1.5	230
521	Identification of Novel Loci Associated With Hip Shape: A Meta-Analysis of Genomewide Association Studies. Journal of Bone and Mineral Research, 2019, 34, 241-251.	3.1	47
522	Obesity, mitochondrial dysfunction, and obstructive lung disease., 2019, , 143-167.		2
523	Role of metabolic syndrome and its components as mediators of the genetic effect on type 2 diabetes: A familyâ€based study in China. Journal of Diabetes, 2019, 11, 552-562.	0.8	3
524	SumHer better estimates the SNP heritability of complex traits from summary statistics. Nature Genetics, 2019, 51, 277-284.	9.4	181
525	Obesity-associated family with sequence similarity 13, member A (FAM13A) is dispensable for adipose development and insulin sensitivity. International Journal of Obesity, 2019, 43, 1269-1280.	1.6	12
526	Meta-analysis of genome-wide association studies for body fat distribution in 694Â649 individuals of European ancestry. Human Molecular Genetics, 2019, 28, 166-174.	1.4	752
527	Associations of obesity and circulating insulin and glucose with breast cancer risk: a Mendelian randomization analysis. International Journal of Epidemiology, 2019, 48, 795-806.	0.9	81
528	Big data management challenges in health researchâ€"a literature review. Briefings in Bioinformatics, 2019, 20, 156-167.	3.2	56
529	Genetic contribution to waist-to-hip ratio in Mexican children and adolescents based on 12 loci validated in European adults. International Journal of Obesity, 2019, 43, 13-22.	1.6	8
530	Large-Scale Epidemiologic Studies of Cardiovascular Diseases in China: Need for Improved Data Collection, Methods, Transparency, and Documentation. Global Heart, 2019, 13, 3.	0.9	2
531	Muscle transcriptome signature and gene regulatory network analysis in two divergent lines of a hilly bovine species Mithun (Bos frontalis). Genomics, 2020, 112, 252-262.	1.3	12

#	Article	IF	CITATIONS
532	Diet–Gene Interactions. , 2020, , 371-376.		0
533	Adiposity and asthma in adults: a bidirectional Mendelian randomisation analysis of The HUNT Study. Thorax, 2020, 75, 202-208.	2.7	22
534	Genetic contributions to NAFLD: leveraging shared genetics to uncover systems biology. Nature Reviews Gastroenterology and Hepatology, 2020, 17, 40-52.	8.2	203
535	Schizophrenia, Bipolar Disorder, and Alzheimer's Disease are not Causal Factors of Bone Mineral Density: A Mendelian Randomization Analysis. Calcified Tissue International, 2020, 106, 131-146.	1.5	8
536	Life Course Adiposity and Amyotrophic Lateral Sclerosis: A Mendelian Randomization Study. Annals of Neurology, 2020, 87, 434-441.	2.8	30
537	<i>CTNNB1/</i> \hat{l}^2 <i>-catenin</i> dysfunction contributes to adiposity by regulating the cross-talk of mature adipocytes and preadipocytes. Science Advances, 2020, 6, eaax9605.	4.7	50
538	Precision medicine at the academic-industry interface. , 2020, , 545-560.		1
539	Genetic variants that associate with cirrhosis have pleiotropic effects on human traits. Liver International, 2020, 40, 405-415.	1.9	38
540	Changes in Proteomic Profiles are Related to Changes in BMI and Fat Distribution During 10 Years of Aging. Obesity, 2020, 28, 178-186.	1.5	13
541	Sex Differences in Genomic Drivers of Adipose Distribution and Related Cardiometabolic Disorders. Arteriosclerosis, Thrombosis, and Vascular Biology, 2020, 40, 45-60.	1.1	55
542	The causal role of elevated uric acid and waist circumference on the risk of metabolic syndrome components. International Journal of Obesity, 2020, 44, 865-874.	1.6	18
543	Causal associations between urinary sodium with body mass, shape and composition: a Mendelian randomization study. Scientific Reports, 2020, 10, 17475.	1.6	10
544	Sex differences in human adipose tissue gene expression and genetic regulation involve adipogenesis. Genome Research, 2020, 30, 1379-1392.	2.4	35
545	Genetics of height and risk of atrial fibrillation: A Mendelian randomization study. PLoS Medicine, 2020, 17, e1003288.	3.9	51
546	Hormonal Contraception and Depression: Updated Evidence and Implications in Clinical Practice. Clinical Drug Investigation, 2020, 40, 1097-1106.	1.1	30
547	Wnt/ \hat{l}^2 -catenin signaling regulates adipose tissue lipogenesis and adipocyte-specific loss is rigorously defended by neighboring stromal-vascular cells. Molecular Metabolism, 2020, 42, 101078.	3.0	53
548	Giant Island Mice Exhibit Widespread Gene Expression Changes in Key Metabolic Organs. Genome Biology and Evolution, 2020, 12, 1277-1301.	1.1	1
549	The genetic architecture of the association between eating behaviors and obesity: combining genetic twin modeling and polygenic risk scores. American Journal of Clinical Nutrition, 2020, 112, 956-966.	2.2	11

#	Article	IF	CITATIONS
550	Obesity and insulin resistance in children. Current Opinion in Pediatrics, 2020, 32, 582-588.	1.0	22
551	Non-random sampling leads to biased estimates of transcriptome association. Scientific Reports, 2020, 10, 6193.	1.6	1
552	Multiple-Tissue Integrative Transcriptome-Wide Association Studies Discovered New Genes Associated With Amyotrophic Lateral Sclerosis. Frontiers in Genetics, 2020, 11, 587243.	1,1	15
553	The Molecular Mechanisms Underlying Mitochondria-Associated Endoplasmic Reticulum Membrane-Induced Insulin Resistance. Frontiers in Endocrinology, 2020, 11, 592129.	1.5	39
554	Causal Inference for Genetic Obesity, Cardiometabolic Profile and COVID-19 Susceptibility: A Mendelian Randomization Study. Frontiers in Genetics, 2020, 11, 586308.	1.1	56
555	Exploring the Role of Contactins across Psychological, Psychiatric and Cardiometabolic Traits within UK Biobank. Genes, 2020, 11, 1326.	1.0	6
556	Distinct infrastructure of lipid networks in visceral and subcutaneous adipose tissues in overweight humans. American Journal of Clinical Nutrition, 2020, 112, 979-990.	2.2	7
557	Pediatric obesity: prevention is better than care. Italian Journal of Pediatrics, 2020, 46, 103.	1.0	21
558	A rank-based normalization method with the fully adjusted full-stage procedure in genetic association studies. PLoS ONE, 2020, 15, e0233847.	1.1	8
559	The contribution of tissue-grouped BMI-associated gene sets to cardiometabolic-disease risk: a Mendelian randomization study. International Journal of Epidemiology, 2020, 49, 1246-1256.	0.9	8
560	Genetics of Obesity in East Asians. Frontiers in Genetics, 2020, 11, 575049.	1.1	19
561	The genetic architecture of appendicular lean mass characterized by association analysis in the UK Biobank study. Communications Biology, 2020, 3, 608.	2.0	83
562	Novel aspects on the role of white adipose tissue in type 2 diabetes. Current Opinion in Pharmacology, 2020, 55, 47-52.	1.7	8
563	Genetic Influences on Disease Subtypes. Annual Review of Genomics and Human Genetics, 2020, 21, 413-435.	2.5	23
564	Canalization and Robustness in Human Genetics and Disease. Annual Review of Genetics, 2020, 54, 189-211.	3.2	22
565	Association of obesity and its genetic predisposition with the risk of severe COVID-19: Analysis of population-based cohort data. Metabolism: Clinical and Experimental, 2020, 112, 154345.	1.5	63
566	Methylome-wide association study of central adiposity implicates genes involved in immune and endocrine systems. Epigenomics, 2020, 12, 1483-1499.	1.0	6
567	A Body Shape Index (ABSI) achieves better mortality risk stratification than alternative indices of abdominal obesity: results from a large European cohort. Scientific Reports, 2020, 10, 14541.	1.6	84

#	Article	IF	CITATIONS
568	The impact of sex on gene expression across human tissues. Science, 2020, 369, .	6.0	329
569	Genetic Studies of Leptin Concentrations Implicate Leptin in the Regulation of Early Adiposity. Diabetes, 2020, 69, 2806-2818.	0.3	26
570	Machine Learning based histology phenotyping to investigate the epidemiologic and genetic basis of adipocyte morphology and cardiometabolic traits. PLoS Computational Biology, 2020, 16, e1008044.	1.5	16
571	Genetic Predisposition to Coronary Artery Disease in Type 2 Diabetes Mellitus. Circulation Genomic and Precision Medicine, 2020, 13, e002769.	1.6	5
572	Adiposity, metabolites, and colorectal cancer risk: Mendelian randomization study. BMC Medicine, 2020, 18, 396.	2.3	76
573	Precision medicine in the era of artificial intelligence: implications in chronic disease management. Journal of Translational Medicine, 2020, 18, 472.	1.8	99
574	A lead candidate functional single nucleotide polymorphism within the WARS2 gene associated with waist-hip-ratio does not alter RNA stability. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2020, 1863, 194640.	0.9	1
575	Identification of type 2 diabetes loci in 433,540 East Asian individuals. Nature, 2020, 582, 240-245.	13.7	282
576	Mendelian randomization analysis does not support causal associations of birth weight with hypertension risk and blood pressure in adulthood. European Journal of Epidemiology, 2020, 35, 685-697.	2.5	9
577	Deciphering Sex-Specific Genetic Architectures Using Local Bayesian Regressions. Genetics, 2020, 215, 231-241.	1.2	7
578	Metabolomic Links between Sugar-Sweetened Beverage Intake and Obesity. Journal of Obesity, 2020, 2020, 1-10.	1,1	11
579	RSPO3 impacts body fat distribution and regulates adipose cell biology in vitro. Nature Communications, 2020, 11, 2797.	5. 8	34
580	The polygenic architecture of left ventricular mass mirrors the clinical epidemiology. Scientific Reports, 2020, 10, 7561.	1.6	13
581	Shared Genetic Etiology of Obesity-Related Traits and Barrett's Esophagus/Adenocarcinoma: Insights from Genome-Wide Association Studies. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 427-433.	1.1	7
582	Genome-wide associations of human gut microbiome variation and implications for causal inference analyses. Nature Microbiology, 2020, 5, 1079-1087.	5.9	144
583	Starch Digestion–Related Amylase Genetic Variants, Diet, and Changes in Adiposity: Analyses in Prospective Cohort Studies and a Randomized Dietary Intervention. Diabetes, 2020, 69, 1917-1926.	0.3	8
584	Relationship of prenatal maternal obesity and diabetes to offspring neurodevelopmental and psychiatric disorders: a narrative review. International Journal of Obesity, 2020, 44, 1981-2000.	1.6	47
585	IGF-1 and cardiometabolic diseases: a Mendelian randomisation study. Diabetologia, 2020, 63, 1775-1782.	2.9	44

#	Article	IF	CITATIONS
586	Phenotype-specific differences in polygenicity and effect size distribution across functional annotation categories revealed by Al-MiXeR. Bioinformatics, 2020, 36, 4749-4756.	1.8	6
587	Estimation of non-null SNP effect size distributions enables the detection of enriched genes underlying complex traits. PLoS Genetics, 2020, 16, e1008855.	1.5	9
588	Discovery of small-molecule enzyme activators by activity-based protein profiling. Nature Chemical Biology, 2020, 16, 997-1005.	3.9	31
589	Transcription Factor KLF14 and Metabolic Syndrome. Frontiers in Cardiovascular Medicine, 2020, 7, 91.	1.1	23
590	Semaphorin 3 C is a Novel Adipokine Representing Exercise-Induced Improvements of Metabolism in Metabolically Healthy Obese Young Males. Scientific Reports, 2020, 10, 10005.	1.6	10
591	Multi-Omics Analysis Reveals MicroRNAs Associated With Cardiometabolic Traits. Frontiers in Genetics, 2020, 11, 110.	1.1	17
592	Childhood obesity and the associated rise in cardiometabolic complications. Nature Metabolism, 2020, 2, 223-232.	5.1	92
593	Sex differences on adipose tissue remodeling: from molecular mechanisms to therapeutic interventions. Journal of Molecular Medicine, 2020, 98, 483-493.	1.7	24
594	FAM13A Represses AMPK Activity and Regulates Hepatic Glucose and Lipid Metabolism. IScience, 2020, 23, 100928.	1.9	16
595	The influence of rare variants in circulating metabolic biomarkers. PLoS Genetics, 2020, 16, e1008605.	1.5	9
596	Perceived neighborhood social cohesion and cardiometabolic risk: a gene $\tilde{A}-$ environment study. Biodemography and Social Biology, 2020, 65, 1-15.	0.4	7
597	Genome-wide association identifies seven loci for pelvic organ prolapse in Iceland and the UK Biobank. Communications Biology, 2020, 3, 129.	2.0	20
598	FAM13A affects body fat distribution and adipocyte function. Nature Communications, 2020, 11, 1465.	5.8	36
599	Capturing functional epigenomes for insight into metabolic diseases. Molecular Metabolism, 2020, 38, 100936.	3.0	9
600	Insights into the multifactorial causation of obesity by integrated genetic and epigenetic analysis. Obesity Reviews, 2020, 21, e13019.	3.1	24
601	Genetic colocalization atlas points to common regulatory sites and genes for hematopoietic traits and hematopoietic contributions to disease phenotypes. BMC Medical Genomics, 2020, 13, 89.	0.7	10
602	A Complex Systems Model of Breast Cancer Etiology: The Paradigm II Conceptual Model. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1720-1730.	1.1	11
603	Evaluating the cardiovascular safety of sclerostin inhibition using evidence from meta-analysis of clinical trials and human genetics. Science Translational Medicine, 2020, 12, .	5.8	68

#	Article	IF	CITATIONS
604	Genetic evidence that higher central adiposity causes gastro-oesophageal reflux disease: a Mendelian randomization study. International Journal of Epidemiology, 2020, 49, 1270-1281.	0.9	20
605	Regulation of inflammation in diabetes: From genetics to epigenomics evidence. Molecular Metabolism, 2020, 41, 101041.	3.0	23
606	Genome-wide association study of adipocyte lipolysis in the GENetics of adipocyte lipolysis (GENiAL) cohort. Molecular Metabolism, 2020, 34, 85-96.	3.0	11
607	The LCORL Locus Is under Selection in Large-Sized Pakistani Goat Breeds. Genes, 2020, 11, 168.	1.0	25
608	Waist circumference increases risk of coronary heart disease: Evidence from a Mendelian randomization study. Molecular Genetics & Enomic Medicine, 2020, 8, e1186.	0.6	18
609	Genome-wide association study of dietary intake in the UK biobank study and its associations with schizophrenia and other traits. Translational Psychiatry, 2020, 10, 51.	2.4	33
610	MAFLD: A Consensus-Driven Proposed Nomenclature for Metabolic Associated Fatty Liver Disease. Gastroenterology, 2020, 158, 1999-2014.e1.	0.6	1,840
611	Insights into the aetiology of snoring from observational and genetic investigations in the UK Biobank. Nature Communications, 2020, 11, 817.	5.8	74
612	Osteoporosis- and obesity-risk interrelationships: an epigenetic analysis of GWAS-derived SNPs at the developmental gene <i>TBX15</i> . Epigenetics, 2020, 15, 728-749.	1.3	11
613	Exploring genetic targets of psoriasis using genome wide association studies (GWAS) for drug repurposing. 3 Biotech, 2020, 10, 43.	1.1	8
614	Impact of Genetic and Nongenetic Factors on Body Mass Index and Waist-Hip Ratio Change in HIV-Infected Individuals Initiating Antiretroviral Therapy. Open Forum Infectious Diseases, 2020, 7, ofz464.	0.4	7
615	Exploring the causal pathway from ischemic stroke to atrial fibrillation: a network Mendelian randomization study. Molecular Medicine, 2020, 26, 7.	1.9	38
616	Heterogeneity in Obesity: Genetic Basis and Metabolic Consequences. Current Diabetes Reports, 2020, 20, 1.	1.7	25
617	Genetic Determinants of Clustering of Cardiometabolic Risk Factors in U.K. Biobank. Metabolic Syndrome and Related Disorders, 2020, 18, 121-127.	0.5	14
618	Risk factors for insulin resistance in midlife Singaporean women. Maturitas, 2020, 137, 50-56.	1.0	6
619	Promoter-anchored chromatin interactions predicted from genetic analysis of epigenomic data. Nature Communications, 2020, 11, 2061.	5.8	8
620	An examination of potential mediators of the relationship between polygenic scores of BMI and waist circumference and phenotypic adiposity. Psychology and Health, 2020, 35, 1151-1161.	1.2	5
621	A combination of genetics and microbiota influences the severity of the obesity phenotype in diet-induced obesity. Scientific Reports, 2020, 10, 6118.	1.6	9

#	Article	IF	CITATIONS
622	Could vitamin D reduce obesity-associated inflammation? Observational and Mendelian randomization study. American Journal of Clinical Nutrition, 2020, 111, 1036-1047.	2.2	28
623	The Role of Genetic Variation of BMI, Body Composition, and Fat Distribution for Mental Traits and Disorders: A Look-Up and Mendelian Randomization Study. Frontiers in Genetics, 2020, 11, 373.	1.1	20
624	Wntless regulates lipogenic gene expression in adipocytes and protects against diet-induced metabolic dysfunction. Molecular Metabolism, 2020, 39, 100992.	3.0	19
625	Genetic identification of cell types underlying brain complex traits yields insights into the etiology of Parkinson's disease. Nature Genetics, 2020, 52, 482-493.	9.4	216
626	A missense variant in Mitochondrial Amidoxime Reducing Component 1 gene and protection against liver disease. PLoS Genetics, 2020, 16, e1008629.	1.5	101
627	Genome-wide association studies and Mendelian randomization analyses for leisure sedentary behaviours. Nature Communications, 2020, 11, 1770.	5.8	66
628	<p>Converging Relationships of Obesity and Hyperuricemia with Special Reference to Metabolic Disorders and Plausible Therapeutic Implications</p> . Diabetes, Metabolic Syndrome and Obesity: Targets and Therapy, 2020, Volume 13, 943-962.	1.1	38
629	Developmentally Driven Changes in Adipogenesis in Different Fat Depots Are Related to Obesity. Frontiers in Endocrinology, 2020, 11, 138.	1.5	12
630	Distinct genetic subtypes of adiposity and glycemic changes in response to weight-loss diet intervention: the POUNDS Lost trial. European Journal of Nutrition, 2021, 60, 249-258.	1.8	6
631	Dysmetabolic adipose tissue in obesity: morphological and functional characteristics of adipose stem cells and mature adipocytes in healthy and unhealthy obese subjects. Journal of Endocrinological Investigation, 2021, 44, 921-941.	1.8	32
632	N-glycans as functional effectors of genetic and epigenetic disease risk. Molecular Aspects of Medicine, 2021, 79, 100891.	2.7	32
633	A genome-wide association study of severe asthma exacerbations in Latino children and adolescents. European Respiratory Journal, 2021, 57, 2002693.	3.1	15
634	Genetic variation in the body mass index of adult survivors of childhood acute lymphoblastic leukemia: A report from the Childhood Cancer Survivor Study and the St. Jude Lifetime Cohort. Cancer, 2021, 127, 310-318.	2.0	6
635	No Evidence for Passive Gene-Environment Correlation or the Influence of Genetic Risk for Psychiatric Disorders on Adult Body Composition via the Adoption Design. Behavior Genetics, 2021, 51, 58-67.	1.4	2
636	Overall and abdominal obesity in relation to venous thromboembolism. Journal of Thrombosis and Haemostasis, 2021, 19, 460-469.	1.9	33
637	Genome-wide Association Studies Reveal Novel Locus With Sex-/Therapy-Specific Fracture Risk Effects in Childhood Cancer Survivors. Journal of Bone and Mineral Research, 2020, 36, 685-695.	3.1	7
638	Conventional and Genetic Evidence on the Association between Adiposity and CKD. Journal of the American Society of Nephrology: JASN, 2021, 32, 127-137.	3.0	39
639	Sex-specific genetic effects across biomarkers. European Journal of Human Genetics, 2021, 29, 154-163.	1.4	48

#	Article	IF	CITATIONS
640	Detecting Clustered Independent Rare Variant Associations Using Genetic Algorithms. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2021, 18, 932-939.	1.9	6
641	Interactions of Carbohydrate Intake and Physical Activity with Regulatory Genes Affecting Glycaemia: A Food4Me Study Analysis. Lifestyle Genomics, 2021, 14, 63-72.	0.6	2
642	tRNA Biology in the Pathogenesis of Diabetes: Role of Genetic and Environmental Factors. International Journal of Molecular Sciences, 2021, 22, 496.	1.8	9
643	Sex-dimorphic genetic effects and novel loci for fasting glucose and insulin variability. Nature Communications, 2021, 12, 24.	5.8	87
644	BDNF Gene as a Precision Skill of Obesity Management. Advances in Experimental Medicine and Biology, 2021, 1331, 233-248.	0.8	7
645	Genetically predicted physical activity levels are associated with lower colorectal cancer risk: a Mendelian randomisation study. British Journal of Cancer, 2021, 124, 1330-1338.	2.9	17
646	Sex Differences in the Incidence of Obesity-Related Gastrointestinal Cancer. International Journal of Molecular Sciences, 2021, 22, 1253.	1.8	14
647	Genomic risk score provides predictive performance for type 2 diabetes in the UK biobank. Acta Diabetologica, 2021, 58, 467-474.	1.2	11
648	A Genomeâ€Wide Association Study of Childhood Body Fatness. Obesity, 2021, 29, 446-453.	1.5	8
649	Genome-wide association study of circulating levels of glucagon during an oral glucose tolerance test. BMC Medical Genomics, 2021, 14, 3.	0.7	3
651	Nontrivial Replication of Loci Detected by Multi-Trait Methods. Frontiers in Genetics, 2021, 12, 627989.	1.1	4
653	GWAS of peptic ulcer disease implicates Helicobacter pylori infection, other gastrointestinal disorders and depression. Nature Communications, 2021, 12, 1146.	5.8	93
654	Causal associations of waist circumference and waist-to-hip ratio with type II diabetes mellitus: new evidence from Mendelian randomization. Molecular Genetics and Genomics, 2021, 296, 605-613.	1.0	13
655	Polygenic risk modeling with latent trait-related genetic components. European Journal of Human Genetics, 2021, 29, 1071-1081.	1.4	14
656	First genome-wide association study of 99 body measures derived from 3-dimensional body scans. Genes and Diseases, 2022, 9, 777-788.	1.5	2
657	Genome-wide discovery of genetic loci that uncouple excess adiposity from its comorbidities. Nature Metabolism, 2021, 3, 228-243.	5.1	70
658	Longitudinal associations of total and trunk fat in childhood and adolescence and risk of hepatic steatosis at 24 years. Pediatric Obesity, 2021, 16, e12773.	1.4	2
659	Who will benefit from bariatric surgery for diabetes? A protocol for an observational cohort study. BMJ Open, 2021, 11, e042355.	0.8	1

#	Article	IF	Citations
660	Habitual consumption of alcohol with meals and lung cancer: a Mendelian randomization study. Annals of Translational Medicine, 2021, 9, 263-263.	0.7	6
661	Bias in two-sample Mendelian randomization when using heritable covariable-adjusted summary associations. International Journal of Epidemiology, 2021, 50, 1639-1650.	0.9	65
662	Cardiometabolic risk factors for COVID-19 susceptibility and severity: A Mendelian randomization analysis. PLoS Medicine, 2021, 18, e1003553.	3.9	105
663	Selection into shift work is influenced by educational attainment and body mass index: a Mendelian randomization study in the UK Biobank. International Journal of Epidemiology, 2021, 50, 1229-1240.	0.9	9
666	Chronic tissue inflammation and metabolic disease. Genes and Development, 2021, 35, 307-328.	2.7	122
667	Birthweight DNA methylation signatures in infant saliva. Clinical Epigenetics, 2021, 13, 57.	1.8	3
668	Associations and limited shared genetic aetiology between bipolar disorder and cardiometabolic traits in the UK Biobank. Psychological Medicine, 2022, 52, 4039-4048.	2.7	10
669	Causal effect of education on type 2 diabetes: A network Mendelian randomization study. World Journal of Diabetes, 2021, 12, 261-277.	1.3	3
671	Genetics of Obesity: What We Have Learned Over Decades of Research. Obesity, 2021, 29, 802-820.	1.5	71
672	Genome-wide association study of neck circumference identifies sex-specific loci independent of generalized adiposity. International Journal of Obesity, 2021, 45, 1532-1541.	1.6	8
673	Genetic Thyrotropin Regulation of Atrial Fibrillation Risk Is Mediated Through an Effect on Height. Journal of Clinical Endocrinology and Metabolism, 2021, 106, 2124-2132.	1.8	8
674	Causal Associations of Anthropometric Measurements With Fracture Risk and Bone Mineral Density: A Mendelian Randomization Study. Journal of Bone and Mineral Research, 2020, 36, 1281-1287.	3.1	13
676	Plasma Cortisol and Risk of Atrial Fibrillation: A Mendelian Randomization Study. Journal of Clinical Endocrinology and Metabolism, 2021, 106, e2521-e2526.	1.8	9
677	The impact of late-career job loss and genetic risk on body mass index: Evidence from variance polygenic scores. Scientific Reports, 2021, 11, 7647.	1.6	11
678	On Bayes Decision Rule for Testing Multiple Hypotheses with Non-symmetric Alternatives. Journal of Statistical Theory and Practice, 2021, 15, 1.	0.3	0
679	Allele-specific variation at <i>APOE</i> i>increases nonalcoholic fatty liver disease and obesity but decreases risk of Alzheimer's disease and myocardial infarction. Human Molecular Genetics, 2021, 30, 1443-1456.	1.4	20
680	Synergizing Mouse and Human Studies to Understand the Heterogeneity of Obesity. Advances in Nutrition, 2021, 12, 2023-2034.	2.9	13
681	A genome-wide methylation study of body fat traits in the Norfolk Island isolate. Nutrition, Metabolism and Cardiovascular Diseases, 2021, 31, 1556-1563.	1.1	4

#	Article	IF	CITATIONS
682	Impact of serum calcium levels on total body bone mineral density: A mendelian randomization study in five age strata. Clinical Nutrition, 2021, 40, 2726-2733.	2.3	16
683	Interpreting type 1 diabetes risk with genetics and single-cell epigenomics. Nature, 2021, 594, 398-402.	13.7	170
684	Pharmacogenetic association of diabetes-associated genetic risk score with rapid progression of coronary artery calcification following treatment with HMG-CoA-reductase inhibitors â€"results of the Heinz Nixdorf Recall Study. Naunyn-Schmiedeberg's Archives of Pharmacology, 2021, 394, 1713-1725.	1.4	4
686	Genetically predicted serum vitamin D and COVID-19: a Mendelian randomisation study. BMJ Nutrition, Prevention and Health, 2021, 4, 213-225.	1.9	25
687	Nutritional Regulation of Human Brown Adipose Tissue. Nutrients, 2021, 13, 1748.	1.7	20
688	GWAS of allometric body-shape indices in UK Biobank identifies loci suggesting associations with morphogenesis, organogenesis, adrenal cell renewal and cancer. Scientific Reports, 2021, 11, 10688.	1.6	68
689	Impact of body composition on COVID-19 susceptibility and severity: A two-sample multivariable Mendelian randomization study. Metabolism: Clinical and Experimental, 2021, 118, 154732.	1.5	59
690	Genomics of body fat distribution. Journal of Genetics, 2021, 100, 1.	0.4	0
691	Methylation and Expression of FTO and PLAG1 Genes in Childhood Obesity: Insight into Anthropometric Parameters and Glucose–Lipid Metabolism. Nutrients, 2021, 13, 1683.	1.7	13
692	Transcriptome-wide association analysis of brain structures yields insights into pleiotropy with complex neuropsychiatric traits. Nature Communications, 2021, 12, 2878.	5.8	25
693	Leveraging eQTLs to identify individual-level tissue of interest for a complex trait. PLoS Computational Biology, 2021, 17, e1008915.	1.5	3
694	Genetic markers of abdominal obesity and weight loss after gastric bypass surgery. PLoS ONE, 2021, 16, e0252525.	1.1	3
695	Genetic Evidence for Different Adiposity Phenotypes and Their Opposing Influences on Ectopic Fat and Risk of Cardiometabolic Disease. Diabetes, 2021, 70, 1843-1856.	0.3	42
696	Genetics of Body Fat Distribution: Comparative Analyses in Populations with European, Asian and African Ancestries. Genes, 2021, 12, 841.	1.0	21
698	An Intronic Risk SNP rs12454712 for Central Obesity Acts As an Allele-Specific Enhancer To Regulate <i>BCL2</i> Expression. Diabetes, 2021, 70, 1679-1688.	0.3	10
700	AKR1C2 and AKR1C3 expression in adipose tissue: Association with body fat distribution and regulatory variants. Molecular and Cellular Endocrinology, 2021, 527, 111220.	1.6	11
701	Association of Genetic Variants in IGF2-Related Genes With Risk of Metabolic Syndrome in the Chinese Han Population. Frontiers in Endocrinology, 2021, 12, 654747.	1.5	2
702	The heritability of body composition. BMC Pediatrics, 2021, 21, 225.	0.7	10

#	Article	IF	CITATIONS
703	Genome-wide association studies identify 137 genetic loci for DNA methylation biomarkers of aging. Genome Biology, 2021, 22, 194.	3.8	90
704	The geometry of clinical labs and wellness states from deeply phenotyped humans. Nature Communications, 2021, 12, 3578.	5.8	19
706	Unlocking the causal link of metabolically different adiposity subtypes with brain volumes and the risks of dementia and stroke: A Mendelian randomization study. Neurobiology of Aging, 2021, 102, 161-169.	1.5	18
707	A Novel Approach Integrating Hierarchical Clustering and Weighted Combination for Association Study of Multiple Phenotypes and a Genetic Variant. Frontiers in Genetics, 2021, 12, 654804.	1.1	7
708	Polycystic Ovary Syndrome Susceptibility Loci Inform Disease Etiological Heterogeneity. Journal of Clinical Medicine, 2021, 10, 2688.	1.0	10
709	Shared genetic architecture between neuroticism, coronary artery disease and cardiovascular risk factors. Translational Psychiatry, 2021, 11, 368.	2.4	10
710	Genome-wide association study of body fat distribution traits in Hispanics/Latinos from the HCHS/SOL. Human Molecular Genetics, 2021, 30, 2190-2204.	1.4	8
711	Identification of a Novel Locus for Gait Speed Decline With Aging: The Long Life Family Study. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2021, 76, e307-e313.	1.7	0
712	Epigenome-Wide Association Study Reveals Methylation Loci Associated With Offspring Gestational Diabetes Mellitus Exposure and Maternal Methylome. Diabetes Care, 2021, 44, 1992-1999.	4.3	17
713	Genetic correlation and causal relationships between cardio-metabolic traits and lung function impairment. Genome Medicine, 2021, 13, 104.	3.6	11
714	Instrumental Heterogeneity in Sex-Specific Two-Sample Mendelian Randomization: Empirical Results From the Relationship Between Anthropometric Traits and Breast/Prostate Cancer. Frontiers in Genetics, 2021, 12, 651332.	1.1	6
715	Wnt Signaling: From Mesenchymal Cell Fate to Lipogenesis and Other Mature Adipocyte Functions. Diabetes, 2021, 70, 1419-1430.	0.3	19
717	Discovery Genome-Wide Association Study of Body Composition in 4,386 Adults From the UK Biobank's Pilot Imaging Enhancement Study. Frontiers in Endocrinology, 2021, 12, 692677.	1.5	0
718	Telomere length and metabolic syndrome traits: A Mendelian randomisation study. Aging Cell, 2021, 20, e13445.	3.0	12
720	Genetic association of the PERIOD3 (PER3) Clock gene with extreme obesity. Obesity Research and Clinical Practice, 2021, 15, 334-338.	0.8	3
721	Joint disease-specificity at the regulatory base-pair level. Nature Communications, 2021, 12, 4161.	5.8	18
722	A Body Shape Index (ABSI), hip index, and risk of cancer in the UK Biobank cohort. Cancer Medicine, 2021, 10, 5614-5628.	1.3	28
723	Life Course Adiposity and Alzheimer's Disease: A Mendelian Randomization Study. Journal of Alzheimer's Disease, 2021, 82, 503-512.	1.2	10

#	Article	IF	CITATIONS
724	Abdominal and gluteofemoral fat depots show opposing associations with postprandial lipemia. American Journal of Clinical Nutrition, 2021, 114, 1467-1475.	2.2	9
725	Birth weight and adult income: An examination of mediation through adult height and body mass. Health Economics (United Kingdom), 2021, 30, 2383-2398.	0.8	2
726	HMGB2 orchestrates mitotic clonal expansion by binding to the promoter of C/EBP \hat{l}^2 to facilitate adipogenesis. Cell Death and Disease, 2021, 12, 666.	2.7	16
727	Transcription factor RFX7 governs a tumor suppressor network in response to p53 and stress. Nucleic Acids Research, 2021, 49, 7437-7456.	6.5	17
728	Genetic Variation in the ASTN2 Locus in Cardiovascular, Metabolic and Psychiatric Traits: Evidence for Pleiotropy Rather Than Shared Biology. Genes, 2021, 12, 1194.	1.0	4
729	Discriminating Heterogeneous Trajectories of Resilience and Depression After Major Life Stressors Using Polygenic Scores. JAMA Psychiatry, 2021, 78, 744.	6.0	33
730	Fat biology and metabolic balance: On the significance of sex. Molecular and Cellular Endocrinology, 2021, 533, 111336.	1.6	10
731	Low Plasma Adiponectin in Risk of Type 2 Diabetes: Observational Analysis and One- and Two-Sample Mendelian Randomization Analyses in 756,219 Individuals. Diabetes, 2021, 70, 2694-2705.	0.3	17
734	Genetically Predicted Insomnia in Relation to 14 Cardiovascular Conditions and 17 Cardiometabolic Risk Factors: A Mendelian Randomization Study. Journal of the American Heart Association, 2021, 10, e020187.	1.6	21
735	Trefoil Factor Family Member 2: From a High-Fat-Induced Gene to a Potential Obesity Therapy Target. Metabolites, 2021, 11, 536.	1.3	5
736	A targeted multi-omics approach reveals paraoxonase-1 as a determinant of obesity-associated fatty liver disease. Clinical Epigenetics, 2021, 13, 158.	1.8	9
737	Alcohol use and cardiometabolic risk in the UK Biobank: A Mendelian randomization study. PLoS ONE, 2021, 16, e0255801.	1.1	24
738	Identification of TBX15 as an adipose master trans regulator of abdominal obesity genes. Genome Medicine, 2021, 13, 123.	3.6	23
740	Multifactorial Basis and Therapeutic Strategies in Metabolism-Related Diseases. Nutrients, 2021, 13, 2830.	1.7	27
741	The genetic architecture of plasma kynurenine includes cardiometabolic disease mechanisms associated with the SH2B3 gene. Scientific Reports, 2021, 11, 15652.	1.6	4
742	Mediation role of body fat distribution (FD) on the relationship between CAV1 rs3807992 polymorphism and metabolic syndrome in overweight and obese women. BMC Medical Genomics, 2021, 14, 202.	0.7	6
743	Waist circumference and waistâ€toâ€height ratio in 7â€yearâ€old children—WHO Childhood Obesity Surveillance Initiative. Obesity Reviews, 2021, 22, e13208.	3.1	13
744	Composite trait Mendelian randomization reveals distinct metabolic and lifestyle consequences of differences in body shape. Communications Biology, 2021, 4, 1064.	2.0	13

#	Article	IF	Citations
745	The link between attention deficit hyperactivity disorder (ADHD) symptoms and obesity-related traits: genetic and prenatal explanations. Translational Psychiatry, 2021, 11, 455.	2.4	19
746	A murine model of the human CREBRFR457Q obesity-risk variant does not influence energy or glucose homeostasis in response to nutritional stress. PLoS ONE, 2021, 16, e0251895.	1.1	3
748	Cannabis use and obesity-traits: A Mendelian randomization study. Drug and Alcohol Dependence, 2021, 226, 108863.	1.6	4
750	Obesity cardiomyopathy: evidence, mechanisms, and therapeutic implications. Physiological Reviews, 2021, 101, 1745-1807.	13.1	150
751	Disentangling sex differences in the shared genetic architecture of posttraumatic stress disorder, traumatic experiences, and social support with body size and composition. Neurobiology of Stress, 2021, 15, 100400.	1.9	3
752	Epigenetic rewiring of skeletal muscle enhancers after exercise training supports a role in whole-body function and human health. Molecular Metabolism, 2021, 53, 101290.	3.0	13
753	Insights into the adipose stem cell niche in health and disease. , 2022, , 57-80.		2
754	Obesity, Type 2 Diabetes, Lifestyle Factors, and Risk of Gallstone Disease: A Mendelian Randomization Investigation. Clinical Gastroenterology and Hepatology, 2022, 20, e529-e537.	2.4	53
755	A Systematic Two-Sample Mendelian Randomization Analysis Identifies Shared Genetic Origin of Endometriosis and Associated Phenotypes. Life, 2021, 11, 24.	1.1	10
756	Biological constraints on GWAS SNPs at suggestive significance thresholds reveal additional BMI loci. ELife, 2021, 10, .	2.8	27
758	Pharmacogenomics and circadian rhythms as mediators of cardiovascular drug-drug interactions. Current Research in Pharmacology and Drug Discovery, 2021, 2, 100025.	1.7	2
759	Genetics of Smoking and Risk of Atherosclerotic Cardiovascular Diseases. JAMA Network Open, 2021, 4, e2034461.	2.8	42
760	Gene-Environment Interaction and Individual Susceptibility to Metabolic Disorders., 2020,, 81-94.		1
761	Precision behavioral medicine: Implications of genetic and genomic discoveries for behavioral weight loss treatment American Psychologist, 2018, 73, 1045-1055.	3.8	9
762	Identification of new susceptibility loci for type 2 diabetes and shared etiological pathways with coronary heart disease. Nature Genetics, 2017, 49, 1450-1457.	9.4	218
763	Effects of Obesity Related Genetic Variations on Visceral and Subcutaneous Fat Distribution in a Chinese Population. Scientific Reports, 2016, 6, 20691.	1.6	47
764	UCP1-independent thermogenesis. Biochemical Journal, 2020, 477, 709-725.	1.7	85
822	Nonconserved Long Intergenic Noncoding RNAs Associate With Complex Cardiometabolic Disease Traits. Arteriosclerosis, Thrombosis, and Vascular Biology, 2021, 41, 501-511.	1.1	8

#	Article	lF	Citations
823	What lipodystrophies teach us about the metabolic syndrome. Journal of Clinical Investigation, 2019, 129, 4009-4021.	3.9	96
824	Deciphering the cellular interplays underlying obesity-induced adipose tissue fibrosis. Journal of Clinical Investigation, 2019, 129, 4032-4040.	3.9	157
825	Adiponectin and human eating behaviour: a Mendelian randomization study. Egyptian Journal of Medical Human Genetics, $2019, 20, .$	0.5	4
826	Evaluation of human dermal fibroblasts directly reprogrammed to adipocyte-like cells as a metabolic disease model. DMM Disease Models and Mechanisms, 2017, 10, 1411-1420.	1.2	11
827	How difficult is the validation of clinical biomarkers?. F1000Research, 2015, 4, 101.	0.8	12
828	Six Novel Loci Associated with Circulating VEGF Levels Identified by a Meta-analysis of Genome-Wide Association Studies. PLoS Genetics, 2016, 12, e1005874.	1.5	56
829	Genome-wide physical activity interactions in adiposity $\hat{a} \in A$ meta-analysis of 200,452 adults. PLoS Genetics, 2017, 13, e1006528.	1.5	158
830	Discovery and fine-mapping of adiposity loci using high density imputation of genome-wide association studies in individuals of African ancestry: African Ancestry Anthropometry Genetics Consortium. PLoS Genetics, 2017, 13, e1006719.	1.5	98
831	Common, low-frequency, and rare genetic variants associated with lipoprotein subclasses and triglyceride measures in Finnish men from the METSIM study. PLoS Genetics, 2017, 13, e1007079.	1.5	49
832	Genetic Predisposition to an Impaired Metabolism of the Branched-Chain Amino Acids and Risk of Type 2 Diabetes: A Mendelian Randomisation Analysis. PLoS Medicine, 2016, 13, e1002179.	3.9	324
833	Multivariate Analysis of Anthropometric Traits Using Summary Statistics of Genome-Wide Association Studies from GIANT Consortium. PLoS ONE, 2016, 11, e0163912.	1,1	19
834	Comparison of HapMap and 1000 Genomes Reference Panels in a Large-Scale Genome-Wide Association Study. PLoS ONE, 2017, 12, e0167742.	1.1	29
835	Obesity, metabolic factors and risk of different histological types of lung cancer: A Mendelian randomization study. PLoS ONE, 2017, 12, e0177875.	1.1	79
836	Approaches to detect genetic effects that differ between two strata in genome-wide meta-analyses: Recommendations based on a systematic evaluation. PLoS ONE, 2017, 12, e0181038.	1.1	27
837	Intestinal Ralstonia pickettii augments glucose intolerance in obesity. PLoS ONE, 2017, 12, e0181693.	1.1	53
838	Nutrigenetics—Personalized Nutrition in the Genetic Age. Exploratory Research and Hypothesis in Medicine, 2017, 2, 1-8.	0.1	8
839	Life style, Perfusion deficits and Co-morbidities Precipitate Inflammation and Cerebrovascular Disorders in Aged Subjects. Discoveries, 2015, 3, e39.	1.5	5
840	The role of epigenetics in hypothalamic energy balance control: implications for obesity. Cell Stress, 2019, 3, 208-220.	1.4	20

#	ARTICLE	IF	CITATIONS
841	Epigenetics of obesity. Cardiovascular Therapy and Prevention (Russian Federation), 2020, 19, 2632.	0.4	6
842	Identification of rs7350481 at chromosome 11q23.3 as a novel susceptibility locus for metabolic syndrome in Japanese individuals by an exome-wide association study. Oncotarget, 2017, 8, 39296-39308.	0.8	13
843	The Landscape of Pervasive Horizontal Pleiotropy in Human Genetic Variation is Driven by Extreme Polygenicity of Human Traits and Diseases. SSRN Electronic Journal, 0, , .	0.4	3
844	Hypoalbuminemia as a marker of protein metabolism disarrangement in patients with stable chronic heart failure. Minerva Medica, 2020, 111, 226-238.	0.3	4
845	Red blood cell components: time to revisit the sources of variability. Blood Transfusion, 2017, 15, 116-125.	0.3	28
846	Sex and Gender Differences in Heart Failure. International Journal of Heart Failure, 2020, 2, 157.	0.9	43
847	Genetic and Environmental Factors Contributing to Visceral Adiposity in Asian Populations. Endocrinology and Metabolism, 2020, 35, 681-695.	1.3	30
848	Neuroinflammation and comorbidities are frequently ignored factors in CNS pathology. Neural Regeneration Research, 2015, 10, 1349.	1.6	90
849	Dietary habits, lifestyle factors and neurodegenerative diseases. Neural Regeneration Research, 2020, 15, 394.	1.6	102
850	A big-data approach to understanding metabolic rate and response to obesity in laboratory mice. ELife, 2020, 9, .	2.8	54
851	Differentially expressed genes reflect disease-induced rather than disease-causing changes in the transcriptome. Nature Communications, 2021, 12, 5647.	5.8	61
852	Leptin Receptor (rs1137101) and Brain-Derived Neurotrophic Factor (rs925946) Gene Variants Are Associated with Obesity in the Early- but Not in the Late-Onset Population of Hungarian Psoriatic Patients. Life, 2021, 11, 1086.	1.1	3
853	Histone H4 lysine 16 acetylation controls central carbon metabolism and diet-induced obesity in mice. Nature Communications, 2021, 12, 6212.	5.8	16
854	Metabolic Messengers: tumour necrosis factor. Nature Metabolism, 2021, 3, 1302-1312.	5.1	155
855	Chromatin accessibility and gene expression during adipocyte differentiation identify context-dependent effects at cardiometabolic GWAS loci. PLoS Genetics, 2021, 17, e1009865.	1.5	9
856	Insights into modifiable risk factors of cholelithiasis: AÂMendelian randomization study. Hepatology, 2022, 75, 785-796.	3.6	102
857	FTO and PLAG1 Genes Expression and FTO Methylation Predict Changes in Circulating Levels of Adipokines and Gastrointestinal Peptides in Children. Nutrients, 2021, 13, 3585.	1.7	6
858	Developmental mechanisms of sex differences: from cells to organisms. Development (Cambridge), 2021, 148, .	1.2	21

#	ARTICLE	IF	CITATIONS
859	Identifying causality, genetic correlation, priority and pathways of large-scale complex exposures of breast and ovarian cancers. British Journal of Cancer, 2021, 125, 1570-1581.	2.9	11
861	Appearance and Internal Aging. , 2015, , 1-10.		0
862	Electronic Health Records in the Genomics Era: Impact of Clinical Genomics. MOJ Proteomics $\&$ Bioinformatics, 2015, 2, .	0.1	0
864	Obesity Kills: Can Genetics Help in the Targeting of Obesity Prevention?. Journal of Clinical & Medical Genomics, 2016, 04, .	0.1	0
867	Diabetes in the Caribbean., 2017,, 127-149.		1
869	Advancing Islet Transplantation: From Donor to Engraftment. , 2017, , 189-211.		0
871	The Genetic Determinants of Common Obesity-Susceptibility., 2017,, 383-425.		0
872	Epigenetic Regulation of Fat Deposition: A Focus on Kr $ ilde{A}$ 4ppel-Like Factor 14 (Klf14). , 2017, , 1-17.		0
881	Obésité de l'adulte. , 2018, , 199-218.		0
882	Nutrigenética e sÃndrome metabólica. , 0, , 185-216.		0
883	SÃndrome metabólica: conceitos, epidemiologia e etiopatogênese. , 0, , 19-38.		0
895	Metabolic syndrome: what changed during last 30 years?. Reproductive Endocrinology, 2018, .	0.0	2
897	The Multifaceted Legacy of the Human Genome Program for Evolutionary Biology: An Epistemological Perspective. Perspectives on Science, 2019, 27, 117-152.	0.3	3
913	Causal associations of body mass index and waist-to-hip ratio with cardiometabolic traits among Chinese children: A Mendelian randomization study. Nutrition, Metabolism and Cardiovascular Diseases, 2020, 30, 1554-1563.	1.1	6
918	Coding Variants are Relevant to the Expression of Obesityâ€Related Genes for Pediatric Adiposity. Obesity, 2021, 29, 194-203.	1.5	3
919	Genetics of obstructive sleep apnea. , 2023, , 55-64.		0
921	Sex as a modulator of lipid metabolism and metabolic disease. , 2020, , 45-61.		0
933	Protein Glycosylation in Diabetes. Advances in Experimental Medicine and Biology, 2021, 1325, 285-305.	0.8	6

#	Article	IF	Citations
934	Insomnia and Coronary Artery Diseases: A Mendelian Randomisation Study. Cardiology Discovery, 2021, 1, 154-162.	0.6	1
935	Birth weight, adult weight, and cardiovascular biomarkers: Evidence from the Cardiovascular Young Finns Study. Preventive Medicine, 2022, 154, 106894.	1.6	5
937	Adipose Tissue Fibrosis in Obesity: Etiology and Challenges. Annual Review of Physiology, 2022, 84, 135-155.	5.6	49
938	The interactions between dietary fats intake and Caveolin 1 rs 3807992 polymorphism with fat distribution in overweight and obese women: a cross-sectional study. BMC Medical Genomics, 2021, 14, 265.	0.7	3
940	Blood Levels of the SMOC1 Hepatokine Are Not Causally Linked with Type 2 Diabetes: A Bidirectional Mendelian Randomization Study. Nutrients, 2021, 13, 4208.	1.7	4
941	Transmission distortion and genetic incompatibilities between alleles in a multigenerational mouse advanced intercross line. Genetics, 2022, 220, .	1.2	5
942	Inferred expression regulator activities suggest genes mediating cardiometabolic genetic signals. PLoS Computational Biology, 2021, 17, e1009563.	1.5	3
943	Effect of General Adiposity and Central Body Fat Distribution on the Circulating Metabolome: A Multicohort Nontargeted Metabolomics Observational and Mendelian Randomization Study. Diabetes, 2022, 71, 329-339.	0.3	14
944	Mendelian randomization analysis of the causal impact of body mass index and waist-hip ratio on rates of hospital admission. Economics and Human Biology, 2022, 44, 101088.	0.7	6
945	Integration of Realâ€World Data and Genetics to Support Target Identification and Validation. Clinical Pharmacology and Therapeutics, 2022, 111, 63-76.	2.3	1
946	APOL1 renal risk variants are associated with obesity and body composition in African ancestry adults. Medicine (United States), 2021, 100, e27785.	0.4	6
947	Targeting the cytoplasmic polyadenylation element-binding protein CPEB4 protects against diet-induced obesity and microbiome dysbiosis. Molecular Metabolism, 2021, 54, 101388.	3.0	8
948	Lipids, Anthropometric Measures, Smoking and Physical Activity Mediate the Causal Pathway From Education to Breast Cancer in Women: A Mendelian Randomization Study. Journal of Breast Cancer, 2021, 24, 504.	0.8	1
949	Polygenic architecture and cardiovascular risk of familial combined hyperlipidemia. Atherosclerosis, 2022, 340, 35-43.	0.4	10
950	Comparison of Anti-factor Xa Levels in Female and Male Patients with Obesity After Enoxaparin Application for Thromboprophylaxis. Obesity Surgery, 2022, 32, 861-867.	1,1	5
951	Network Mendelian randomization study: exploring the causal pathway from insomnia to type 2 diabetes. BMJ Open Diabetes Research and Care, 2022, 10, e002510.	1.2	7
953	Obesity genetics: current state of the problem. Profilakticheskaya Meditsina, 2021, 24, 89.	0.2	1
954	The Physiology of Bodyweight Regulation. , 2022, , 1808-1814.		0

#	ARTICLE	IF	CITATIONS
956	Phenome-wide association study of the major histocompatibility complex region in the Korean population identifies novel association signals. Human Molecular Genetics, 2022, , .	1.4	1
957	Mendelian randomisation analyses of UK Biobank and published data suggest that increased adiposity lowers risk of breast and prostate cancer. Scientific Reports, 2022, 12, 909.	1.6	2
958	Genome-wide association study identifies genetic risk loci for adiposity in a Taiwanese population. PLoS Genetics, 2022, 18, e1009952.	1.5	6
959	Identification of genetic loci simultaneously associated with multiple cardiometabolic traits. Nutrition, Metabolism and Cardiovascular Diseases, 2022, 32, 1027-1034.	1.1	4
960	Genetic Predictors of Change in Waist Circumference and Waist-to-Hip Ratio With Lifestyle Intervention: The Trans-NIH Consortium for Genetics of Weight Loss Response to Lifestyle Intervention. Diabetes, 2022, 71, 669-676.	0.3	5
961	Rare Variant Analysis of Obesity-Associated Genes in Young Adults With Severe Obesity From a Consanguineous Population of Pakistan. Diabetes, 2022, 71, 694-705.	0.3	7
962	Sex hormones, adiposity, and metabolic traits in men and women: a Mendelian randomisation study. European Journal of Endocrinology, 2022, 186, 407-416.	1.9	17
963	Pathological and therapeutic roles of bioactive peptide trefoil factor 3 in diverse diseases: recent progress and perspective. Cell Death and Disease, 2022, 13, 62.	2.7	25
964	iPSCs in insulin resistance, type 2 diabetes, and the metabolic syndrome., 2022,, 275-302.		0
965	Distinct properties of adipose stem cell subpopulations determine fat depot-specific characteristics. Cell Metabolism, 2022, 34, 458-472.e6.	7.2	56
966	Identification of the genetic mechanism that associates <i>L3MBTL3</i> to multiple sclerosis. Human Molecular Genetics, 2022, 31, 2155-2163.	1.4	4
967	Comprehensive Statistical and Bioinformatics Analysis in the Deciphering of Putative Mechanisms by Which Lipid-Associated GWAS Loci Contribute to Coronary Artery Disease. Biomedicines, 2022, 10, 259.	1.4	7
968	Harnessing tissue-specific genetic variation to dissect putative causal pathways between body mass index and cardiometabolic phenotypes. American Journal of Human Genetics, 2022, 109, 240-252.	2.6	15
969	Obesity and risk of female reproductive conditions: A Mendelian randomisation study. PLoS Medicine, 2022, 19, e1003679.	3.9	50
970	Assessment of causal effects of visceral adipose tissue on risk of cancers: a Mendelian randomization study. International Journal of Epidemiology, 2022, 51, 1204-1218.	0.9	15
971	PGS-server: accuracy, robustness and transferability of polygenic score methods for biobank scale studies. Briefings in Bioinformatics, 2022, 23, .	3.2	12
972	Smoking behavior might affect allergic rhinitis and vasomotor rhinitis differently: A mendelian randomization appraisal. World Allergy Organization Journal, 2022, 15, 100630.	1.6	5
973	Adiposity, diabetes, lifestyle factors and risk of gastroesophageal reflux disease: a Mendelian randomization study. European Journal of Epidemiology, 2022, 37, 747-754.	2.5	29

#	Article	IF	CITATIONS
974	Separating the direct effects of traits on atherosclerotic cardiovascular disease from those mediated by type 2 diabetes. Diabetologia, 2022, 65, 790-799.	2.9	9
976	Identification of Scd5 as a functional regulator of visceral fat deposition and distribution. IScience, 2022, 25, 103916.	1.9	3
977	The interactions between interleukin-1 family genes: IL1A, IL1B, IL1RN, and obesity parameters. BMC Genomics, 2022, 23, 112.	1.2	10
978	Sex differences in white adipose tissue expansion: emerging molecular mechanisms. Clinical Science, 2021, 135, 2691-2708.	1.8	10
979	Smoking modifies the effect of two independent SNPs rs5063 and rs198358 of NPPA on central obesity in the Chinese Han population. Journal of Genetics, 2018, 97, 987-994.	0.4	1
980	Genetics of obesity and its measures in India. Journal of Genetics, 2018, 97, 1047-1071.	0.4	1
981	Exploring Obesity as a Gendered Contagion: Impact on Lifestyle Interventions to Improve Cardiovascular Health. Clinical Therapeutics, 2022, 44, 23-32.	1.1	3
982	Adipose-Derived Extracellular Vesicles: Systemic Messengers and Metabolic Regulators in Health and Disease. Frontiers in Physiology, 2022, 13, 837001.	1.3	17
983	Metabolic and Epigenetic Regulation by Estrogen in Adipocytes. Frontiers in Endocrinology, 2022, 13, 828780.	1.5	23
985	Roles of Cardiometabolic Factors in Mediating the Causal Effect of Type 2 Diabetes on Cardiovascular Diseases: A Two-Step, Two-Sample Multivariable Mendelian Randomization Study. Frontiers in Cardiovascular Medicine, 2022, 9, 813208.	1.1	11
986	Reassessing Human Adipose Tissue. New England Journal of Medicine, 2022, 386, 768-779.	13.9	170
988	Increased Serum VEGF-B Level Is Associated With Renal Function Impairment in Patients With Type 2 Diabetes. Frontiers in Endocrinology, 2022, 13, 862545.	1.5	5
990	Shared genetic loci for body fat storage and adipocyte lipolysis in humans. Scientific Reports, 2022, 12, 3666.	1.6	3
991	A single-cell atlas of human and mouse white adipose tissue. Nature, 2022, 603, 926-933.	13.7	277
992	Management of BMI Is a Potential New Approach for the Prevention of Idiopathic Pulmonary Fibrosis. Frontiers in Genetics, 2022, 13, 821029.	1.1	6
993	Genetic and phenotypic links between obesity and extracellular vesicles. Human Molecular Genetics, 2022, 31, 3643-3651.	1.4	2
995	Aberrant overexpression of HOTAIR inhibits abdominal adipogenesis through remodelling of genome-wide DNA methylation and transcription. Molecular Metabolism, 2022, 60, 101473.	3.0	11
996	Genome-Wide Association Study Identifies Genetic Loci Associated With Fat Cell Number and Overlap With Genetic Risk Loci for Type 2 Diabetes. Diabetes, 2022, 71, 1350-1362.	0.3	3

#	Article	IF	Citations
997	Changes in adiposity over the life course and gene expression in postmenopausal women. Cancer Medicine, 2022 , , .	1.3	1
998	Proteomic Profiles of Body Mass Index and Waist-to-Hip Ratio and Their Role in Incidence of Diabetes. Journal of Clinical Endocrinology and Metabolism, 2022, 107, e2982-e2990.	1.8	8
999	A Novel Hierarchical Clustering Approach for Joint Analysis of Multiple Phenotypes Uncovers Obesity Variants Based on ARIC. Frontiers in Genetics, 2022, 13, 791920.	1.1	5
1001	Plasma Growth Factor Gene Expression and Mammographic Breast Density in Postmenopausal Women. Cancer Prevention Research, 2022, 15, 391-398.	0.7	1
1002	Impact of nonrandom selection mechanisms on the causal effect estimation for two-sample Mendelian randomization methods. PLoS Genetics, 2022, 18, e1010107.	1.5	2
1004	Immune Cell Regulation of White Adipose Progenitor Cell Fate. Frontiers in Endocrinology, 2022, 13, 859044.	1.5	5
1005	Identification of genetic effects underlying type 2 diabetes in South Asian and European populations. Communications Biology, 2022, 5, 329.	2.0	21
1006	Genetically predicted frailty index and risk of stroke and Alzheimer's disease. European Journal of Neurology, 2022, 29, 1913-1921.	1.7	9
1007	Ancestral diversity improves discovery and fine-mapping of genetic loci for anthropometric traitsâ€"The Hispanic/Latino Anthropometry Consortium. Human Genetics and Genomics Advances, 2022, 3, 100099.	1.0	3
1009	Inflammation subtypes in psychosis and their relationships with genetic risk for psychiatric and cardiometabolic disorders. Brain, Behavior, & Immunity - Health, 2022, 22, 100459.	1.3	8
1010	Contributions of obesity to kidney health and disease: insights from Mendelian randomization and the human kidney transcriptomics. Cardiovascular Research, 2022, 118, 3151-3161.	1.8	17
1012	Identification of Rare Loss-of-Function Genetic Variation Regulating Body Fat Distribution. Journal of Clinical Endocrinology and Metabolism, 2022, 107, 1065-1077.	1.8	12
1013	The association between depression and metabolic syndrome and its components: a bidirectional two-sample Mendelian randomization study. Translational Psychiatry, 2021, 11, 633.	2.4	51
1014	Aberrant Ca ²⁺ signaling by IP ₃ Rs in adipocytes links inflammation to metabolic dysregulation in obesity. Science Signaling, 2021, 14, eabf2059.	1.6	5
1015	Abdominal volume index trajectories and risk of diabetes mellitus: Results from the China Health and Nutrition Survey. Journal of Diabetes Investigation, 2022, 13, 868-877.	1.1	9
1018	Causative Mechanisms of Childhood and Adolescent Obesity Leading to Adult Cardiometabolic Disease: A Literature Review. Applied Sciences (Switzerland), 2021, 11, 11565.	1.3	7
1019	Genetic liability for prescription opioid use and risk of cardiovascular diseases: a multivariable Mendelian randomization study. Addiction, 2022, 117, 1382-1391.	1.7	33
1020	Impact of microRNA Regulated Macrophage Actions on Adipose Tissue Function in Obesity. Cells, 2022, 11, 1336.	1.8	7

#	Article	IF	Citations
1021	Distinct causal effects of body fat distribution on cardiometabolic traits among children: findings from the BCAMS study. Nutrition, Metabolism and Cardiovascular Diseases, 2022, , .	1.1	7
1032	Transcriptome-wide analyses of adipose tissue in outbred rats reveal genetic regulatory mechanisms relevant for human obesity. Physiological Genomics, 2022, 54, 206-219.	1.0	9
1033	Genetically Determined Lifestyle and Cardiometabolic Risk Factors Mediate the Association of Genetically Predicted Age at Menarche With Genetic Predisposition to Myocardial Infarction: A Two-Step, Two-Sample Mendelian Randomization Study. Frontiers in Cardiovascular Medicine, 2022, 9, 821068.	1.1	3
1034	The Causal Association Between Obesity and Primary Open-Angle Glaucoma: A Two-Sample Mendelian Randomization Study. Frontiers in Genetics, 2022, 13, 835524.	1.1	13
1036	Genetic analysis of the PCSK9 locus in psychological, psychiatric, metabolic and cardiovascular traits in UK Biobank. European Journal of Human Genetics, 2022, 30, 1380-1390.	1.4	8
1037	Increased Adipose Tissue Indices of Androgen Catabolism and Aromatization in Women With Metabolic Dysfunction. Journal of Clinical Endocrinology and Metabolism, 2022, 107, e3330-e3342.	1.8	8
1038	Can adult polygenic scores improve prediction of body mass index in childhood?. International Journal of Obesity, 2022, 46, 1375-1383.	1.6	7
1039	The relationship between central obesity and bone mineral density: a Mendelian randomization study. Diabetology and Metabolic Syndrome, 2022, 14, 63.	1.2	8
1040	Causality of abdominal obesity on cognition: a trans-ethnic Mendelian randomization study. International Journal of Obesity, 2022, 46, 1487-1492.	1.6	10
1041	Mendelian randomization highlights the causal association of obesity with periodontal diseases. Journal of Clinical Periodontology, 2022, 49, 662-671.	2.3	11
1043	General and abdominal obesity operate differently as influencing factors of fracture risk in old adults. IScience, 2022, 25, 104466.	1.9	6
1045	Association of genetic variants related to combined exposure to higher BMI and waist-to-hip ratio on lifelong cardiovascular risk in UK Biobank. Public Health Nutrition, 2023, 26, 416-424.	1.1	0
1047	Systematic analysis of nutrigenomic effects of polyphenols related to cardiometabolic health in humans – Evidence from untargeted mRNA and miRNA studies. Ageing Research Reviews, 2022, 79, 101649.	5.0	11
1048	Mendelian Randomization Study of Age at Diagnosis of Diabetes with Cardiovascular Diseases and Cardiometabolic Traits. SSRN Electronic Journal, 0, , .	0.4	0
1049	Observational and Genetic Associations of Modifiable Risk Factors with Aortic Valve Stenosis: A Prospective Cohort Study of 0.5 Million Participants. Nutrients, 2022, 14, 2273.	1.7	7
1050	Genetic Architecture of Plasma Alphaâ€Aminoadipic Acid Reveals a Relationship With Highâ€Density Lipoprotein Cholesterol. Journal of the American Heart Association, 2022, 11, .	1.6	6
1053	Adiposity and grip strength: a Mendelian randomisation study in UK Biobank. BMC Medicine, 2022, 20, .	2.3	6
1054	Genetic Variation in Targets of Antidiabetic Drugs and Alzheimer Disease Risk. Neurology, 2022, 99, .	1.5	18

#	Article	IF	CITATIONS
1055	Pinpointing the genetic and cellular links between sleep and metabolism. Sleep, 0, , .	0.6	1
1058	Prediction of type 2 diabetes mellitus onset using logistic regression-based scorecards. ELife, 0, 11, .	2.8	7
1059	Causal Associations of Obesity With Achilles Tendinopathy: A Two-Sample Mendelian Randomization Study. Frontiers in Endocrinology, 0, 13 , .	1.5	2
1060	Metabolic (dysfunction)-associated fatty liver disease in individuals of normal weight. Nature Reviews Gastroenterology and Hepatology, 2022, 19, 638-651.	8.2	69
1061	CLSTN3 gene variant associates with obesity risk and contributes to dysfunction in white adipose tissue. Molecular Metabolism, 2022, 63, 101531.	3.0	7
1062	PTBP2 \hat{a} e" a gene with relevance for both Anorexia nervosa and body weight regulation. Translational Psychiatry, 2022, 12, .	2.4	4
1063	The association of obesity-related traits on COVID-19 severity and hospitalization is affected by socio-economic status: a multivariable Mendelian randomization study. International Journal of Epidemiology, 2022, 51, 1371-1383.	0.9	4
1065	A Novel Insight into Controversial Risk Factors of Intrahepatic Cholangiocarcinoma: A Mendelian Randomization Study. SSRN Electronic Journal, 0, , .	0.4	0
1067	A genome-wide cross-trait analysis identifies shared loci and causal relationships of type 2 diabetes and glycaemic traits with polycystic ovary syndrome. Diabetologia, 2022, 65, 1483-1494.	2.9	13
1068	Identifying Causes of Fracture Beyond Bone Mineral Density: Evidence From Human Genetics. Journal of Bone and Mineral Research, 2020, 37, 1592-1602.	3.1	5
1069	Inherited basis of visceral, abdominal subcutaneous and gluteofemoral fat depots. Nature Communications, 2022, 13, .	5.8	43
1070	Causality of anthropometric markers associated with polycystic ovarian syndrome: Findings of a Mendelian randomization study. PLoS ONE, 2022, 17, e0269191.	1.1	4
1073	Comparison of Methods Utilizing Sex-Specific PRSs Derived From GWAS Summary Statistics. Frontiers in Genetics, 0, 13, .	1.1	4
1075	Nutrition for precision health: The time is now. Obesity, 2022, 30, 1335-1344.	1.5	6
1076	The consequences of a weightâ€eentric approach to healthcare: A case for a paradigm shift in how clinicians address body weight. Nutrition in Clinical Practice, 2022, 37, 1291-1306.	1.1	16
1077	Bidirectional two-sample Mendelian randomization analysis identifies causal associations between relative carbohydrate intake and depression. Nature Human Behaviour, 2022, 6, 1569-1576.	6.2	30
1078	DNA methylation and waist-to-hip ratio: an epigenome-wide association study in Chinese monozygotic twins. Journal of Endocrinological Investigation, 2022, 45, 2365-2376.	1.8	7
1079	Rare loss of function variants in the hepatokine gene INHBE protect from abdominal obesity. Nature Communications, 2022, 13, .	5.8	15

#	Article	IF	CITATIONS
1080	HOTAIR interacts with PRC2 complex regulating the regional preadipocyte transcriptome and human fat distribution. Cell Reports, 2022, 40, 111136.	2.9	17
1082	Strategies to identify causal common genetic variants and corresponding effector genes for paediatric obesity. Pediatric Obesity, 2022, 17, .	1.4	1
1083	Sex Differences in Adiposity and Cardiovascular Diseases. International Journal of Molecular Sciences, 2022, 23, 9338.	1.8	16
1085	Examination on the risk factors of cholangiocarcinoma: A Mendelian randomization study. Frontiers in Pharmacology, 0, 13 , .	1.6	5
1086	Genetically predicted cortisol levels and risk of venous thromboembolism. PLoS ONE, 2022, 17, e0272807.	1.1	3
1087	Integrated analyses of growth differentiation factor-15 concentration and cardiometabolic diseases in humans. ELife, $0,11,.$	2.8	6
1089	Metabolic Effects of the Waist-To-Hip Ratio Associated Locus GRB14/COBLL1 Are Related to GRB14 Expression in Adipose Tissue. International Journal of Molecular Sciences, 2022, 23, 8558.	1.8	3
1090	Identification of novel genes whose expression in adipose tissue affects body fat mass and distribution: an RNA-Seq and Mendelian Randomization study. European Journal of Human Genetics, 0, ,	1.4	O
1091	Causal relationships of obesity on musculoskeletal chronic pain: A two-sample Mendelian randomization study. Frontiers in Endocrinology, 0, 13, .	1.5	4
1092	Multiancestry exome sequencing reveals INHBE mutations associated with favorable fat distribution and protection from diabetes. Nature Communications, 2022, 13 , .	5.8	18
1093	Phenotype Harmonization in the GLIDE2 Oral Health Genomics Consortium. Journal of Dental Research, 0, , 002203452211097.	2.5	2
1094	A Wars2 mutant mouse shows a sex and diet specific change in fat distribution, reduced food intake and depot-specific upregulation of WAT browning. Frontiers in Physiology, 0, 13 , .	1.3	5
1096	Large-scale genome-wide association study of coronary artery disease in genetically diverse populations. Nature Medicine, 2022, 28, 1679-1692.	15.2	106
1097	The role of obesity, type 2 diabetes, and metabolic factors in gout: A Mendelian randomization study. Frontiers in Endocrinology, 0, 13 , .	1.5	11
1098	The weight of childhood adversity: evidence that childhood adversity moderates the impact of genetic risk on waist circumference in adulthood. International Journal of Obesity, 0, , .	1.6	1
1099	Genetic insights into therapeutic targets for aortic aneurysms: A Mendelian randomization study. EBioMedicine, 2022, 83, 104199.	2.7	13
1100	Polygenic Scores of Diabetes Related Traits Highlight Differences in Subgroups of Type 2 Diabetes in India. SSRN Electronic Journal, 0, , .	0.4	0
1101	Phenotypeâ€informed polygenic risk scores are associated with worse outcome in individuals at risk of Alzheimer's disease. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2022, 14,	1.2	1

#	Article	IF	CITATIONS
1102	The association of measures of body shape and adiposity with incidence of cardiometabolic disease from an ageing perspective. GeroScience, 0 , , .	2.1	0
1103	Causal associations of obesity related anthropometric indicators and body compositions with knee and hip arthritis: A large-scale genetic correlation study. Frontiers in Endocrinology, 0, 13 , .	1.5	7
1104	Genome-wide association analyses of physical activity and sedentary behavior provide insights into underlying mechanisms and roles in disease prevention. Nature Genetics, 2022, 54, 1332-1344.	9.4	64
1105	Keratinocyte-associated protein 3 plays a role in body weight and adiposity with differential effects in males and females. Frontiers in Genetics, 0, 13, .	1.1	7
1106	Modelling metabolic diseases and drug response using stem cells and organoids. Nature Reviews Endocrinology, 2022, 18, 744-759.	4.3	30
1107	Age at Menarche Mediating Visceral Adipose Tissue's Influence on Pre-eclampsia: A Mendelian Randomization Study. Journal of Clinical Endocrinology and Metabolism, 2023, 108, 405-413.	1.8	2
1108	Leveraging omics data to boost the power of genome-wide association studies. Human Genetics and Genomics Advances, 2022, 3, 100144.	1.0	0
1109	Genetic liability to obesity and peptic ulcer disease: a Mendelian randomization study. BMC Medical Genomics, 2022, 15 , .	0.7	4
1110	Adiposity and the risk of dementia: mediating effects from inflammation and lipid levels. European Journal of Epidemiology, 2022, 37, 1261-1271.	2.5	5
1111	The causal relationship between abdominal obesity and lower bone mineral density: A two-sample mendelian randomization study. Frontiers in Genetics, $0,13,.$	1.1	4
1112	Mendelian randomization prioritizes abdominal adiposity as anÂindependent causal factor for liver fat accumulation and cardiometabolic diseases. Communications Medicine, 2022, 2, .	1,9	8
1113	Causal relationship among obesity and body fat distribution and epilepsy subtypes. Frontiers in Neurology, 0, 13, .	1.1	1
1114	The Sexual Dimorphism of Human Adipose Depots. Biomedicines, 2022, 10, 2615.	1.4	7
1116	scGWAS: landscape of trait-cell type associations by integrating single-cell transcriptomics-wide and genome-wide association studies. Genome Biology, 2022, 23, .	3.8	11
1117	Polymorphism of LYPLAL1 and TGFA genes associated with progression of knee osteoarthritis in residents Central Chernozem Region of Russia. Travmatologi \tilde{A}^{ξ} I Ortopedi \tilde{A}^{ξ} Rossii, 0, , .	0.1	0
1118	Causal relationship between obesity, lifestyle factors and risk of benign prostatic hyperplasia: a univariable and multivariable Mendelian randomization study. Journal of Translational Medicine, 2022, 20, .	1.8	12
1119	Sex-specific genetic association of brain and muscle Arnt-like protein-1 (BMAL1) and obesity in Chinese youth. Obesity Research and Clinical Practice, 2022, 16, 464-469.	0.8	2
1120	Role of long non-coding RNAs in adipose tissue metabolism and associated pathologies. Biochemical Pharmacology, 2022, 206, 115305.	2.0	2

#	Article	IF	CITATIONS
1121	Phenotype expansion and neurological manifestations of neurobehavioural disease caused by a variant in RFX7. European Journal of Medical Genetics, 2023, 66, 104657.	0.7	2
1123	Genome-wide association study of a lipedema phenotype among women in the UK Biobank identifies multiple genetic risk factors. European Journal of Human Genetics, 2023, 31, 338-344.	1.4	3
1124	Causal Associations Between Age at Diagnosis of Diabetes and Cardiovascular Outcomes: A Mendelian Randomization Study. Journal of Clinical Endocrinology and Metabolism, 2023, 108, 1202-1214.	1.8	3
1126	Investigating modifiable pathways in psoriasis: A Mendelian randomization study. Journal of the American Academy of Dermatology, 2023, 88, 593-601.	0.6	3
1127	Genetic investigation of the contribution of body composition to anorexia nervosa in an electronic health record setting. Translational Psychiatry, 2022, 12, .	2.4	0
1128	Whole-exome sequence analysis of anthropometric traits illustrates challenges in identifying effects of rare genetic variants. Human Genetics and Genomics Advances, 2023, 4, 100163.	1.0	2
1129	Semi-Automated Image Segmentation of Peri-Prostatic Tissue on MRI and Radiomics Features Stability: A Feasibility Study for Locally Advanced Prostate Cancer Detection. , 2022, , .		0
1132	Prenatal ozone exposure programs a sexually dimorphic susceptibility to highâ€fat diet in adolescent <scp>Long Evans</scp> rats. FASEB Journal, 2022, 36, .	0.2	0
1133	Insight into genetic, biological, and environmental determinants of sexual-dimorphism in type 2 diabetes and glucose-related traits. Frontiers in Cardiovascular Medicine, 0, 9, .	1.1	0
1135	A fast and powerful linear mixed model approach for genotype-environment interaction tests in large-scale GWAS. Briefings in Bioinformatics, 2023, 24, .	3.2	3
1136	Effects of Pre-Pregnancy Overweight/Obesity on the Pattern of Association of Hypertension Susceptibility Genes with Preeclampsia. Life, 2022, 12, 2018.	1.1	6
1138	The Role of Molecular and Hormonal Factors in Obesity and the Effects of Physical Activity in Children. International Journal of Molecular Sciences, 2022, 23, 15413.	1.8	2
1140	Fragile Effects of Climatic Variation on Goat Protein and its Products: A Review. Current Research in Nutrition and Food Science, 2022, 10, 884-894.	0.3	5
1141	BMI-adjusted adipose tissue volumes exhibit depot-specific and divergent associations with cardiometabolic diseases. Nature Communications, 2023, 14, .	5.8	24
1142	Identification of Novel Genetic Risk Factors for Focal Segmental Glomerulosclerosis in Children: Results From the Chronic Kidney Disease in Children (CKiD) Cohort. American Journal of Kidney Diseases, 2023, 81, 635-646.e1.	2.1	6
1143	Diabetic retinopathy risk in patients with unhealthy lifestyle: A Mendelian randomization study. Frontiers in Endocrinology, 0, 13, .	1.5	6
1144	Causal effect of central obesity on left ventricular structure and function in preserved EF population: A Mendelian randomization study. Frontiers in Cardiovascular Medicine, 0, 9, .	1.1	2
1145	Causal Relationships of General and Abdominal Adiposity on Osteoarthritis: A Two-Sample Mendelian Randomization Study. Journal of Clinical Medicine, 2023, 12, 320.	1.0	9

#	Article	IF	CITATIONS
1146	The Polymorphic Locus rs167479 of the RGL3 Gene Is Associated with the Risk of Severe Preeclampsia. Russian Journal of Genetics, 2022, 58, 1543-1550.	0.2	0
1147	Illuminating the †healthy obese' phenotype. Nature Metabolism, 2023, 5, 193-194.	5.1	4
1149	A phenome-wide comparative analysis of genetic discordance between obesity and type 2 diabetes. Nature Metabolism, 2023, 5, 237-247.	5.1	18
1150	Weak and pleiotropy robust sexâ€stratified Mendelian randomization in the one sample and two sample settings. Genetic Epidemiology, 2023, 47, 135-151.	0.6	2
1151	Causal effect of lower birthweight on nonâ€alcoholic fatty liver disease and mediating roles of insulin resistance and metabolites. Liver International, 2023, 43, 829-839.	1.9	6
1153	A Mendelian Randomization Analysis Investigates Causal Associations between Inflammatory Bowel Diseases and Variable Risk Factors. Nutrients, 2023, 15, 1202.	1.7	9
1154	GWAS-Identified Variants for Obesity Do Not Influence the Risk of Developing Multiple Myeloma: A Population-Based Study and Meta-Analysis. International Journal of Molecular Sciences, 2023, 24, 6029.	1.8	2
1155	Schizophrenia and cardiometabolic abnormalities: A Mendelian randomization study. Frontiers in Genetics, $0,14,.$	1.1	0
1156	Evaluating the effect of metabolic traits on oral and oropharyngeal cancer risk using Mendelian randomization. ELife, 0, 12, .	2.8	7
1157	Adiposity impacts cognitive function in Asian populations: an epidemiological and Mendelian Randomization study. The Lancet Regional Health - Western Pacific, 2023, 33, 100710.	1.3	10
1158	Multi-omics approaches for precision obesity management. Wiener Klinische Wochenschrift, 0, , .	1.0	1
1159	Does educational attainment modify the causal relationship between adiposity and cardiovascular disease? A Mendelian randomization study. SSM - Population Health, 2023, 21, 101351.	1.3	0
1160	Obesity and risk of gestational diabetes mellitus: A two-sample Mendelian randomization study. Diabetes Research and Clinical Practice, 2023, 197, 110561.	1.1	7
1161	Consolidation of metabolomic, proteomic, and GWAS data in connective model of schizophrenia. Scientific Reports, 2023, 13, .	1.6	3
1162	A sex-specific genome-wide association study of depression phenotypes in UK Biobank. Molecular Psychiatry, 2023, 28, 2469-2479.	4.1	12
1163	Genetically predicted adipose tissue distribution influences the risk of atherosclerosis. European Journal of Preventive Cardiology, 0, , .	0.8	0
1164	Human RSPO1 Mutation Represses Beige Adipocyte Thermogenesis and Contributes to Dietâ€Induced Adiposity. Advanced Science, 2023, 10, .	5.6	5
1165	Iron status and obesity-related traits: A two-sample bidirectional Mendelian randomization study. Frontiers in Endocrinology, 0, 14, .	1.5	0

#	Article	IF	CITATIONS
1166	The interplay between inflammatory cytokines and cardiometabolic disease: bi-directional mendelian randomisation study., 2023, 2, e000157.		4
1167	Gene–Nutrient Interactions in Obesity: COBLL1 Genetic Variants Interact with Dietary Fat Intake to Modulate the Incidence of Obesity. International Journal of Molecular Sciences, 2023, 24, 3758.	1.8	O
1168	LGR4: A New Receptor Member in Endocrine and Metabolic Diseases. Endocrine Reviews, 0, , .	8.9	1
1169	Establishing causal relationships between sleep and adiposity traits using Mendelian randomization. Obesity, 2023, 31, 861-870.	1.5	11
1170	Capturing the multifaceted function of adipose tissue macrophages. Frontiers in Immunology, 0, 14, .	2.2	2
1171	Genetics of sexually dimorphic adipose distribution in humans. Nature Genetics, 2023, 55, 461-470.	9.4	15
1173	Genetic Evidence for Causal Effects of Socioeconomic, Lifestyle, and Cardiometabolic Factors on Epigenetic-Age Acceleration. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2023, 78, 1083-1091.	1.7	11
1174	Multi-omics analysis identifies RFX7 targets involved in tumor suppression and neuronal processes. Cell Death Discovery, 2023, 9, .	2.0	3
1176	Technologies, strategies, and cautions when deconvoluting genomeâ€wide association signals: ⟨i⟩FTO⟨ i⟩ in focus. Obesity Reviews, 2023, 24, .	3.1	3
1177	Causal associations between modifiable risk factors and pancreatitis: A comprehensive Mendelian randomization study. Frontiers in Immunology, 0, 14 , .	2.2	6
1178	The Genetic Basis of Childhood Obesity: A Systematic Review. Nutrients, 2023, 15, 1416.	1.7	8
1179	Obesity and the risk of cardiometabolic diseases. Nature Reviews Cardiology, 2023, 20, 475-494.	6.1	48
1181	Gene-by-Sex Interactions: Genome-Wide Association Study Reveals Five SNPs Associated with Obesity and Overweight in a Male Population. Genes, 2023, 14, 799.	1.0	2
1182	Identifying the potential causal role of insomnia symptoms on 11,409 health-related outcomes: a phenome-wide Mendelian randomisation analysis in UK Biobank. BMC Medicine, 2023, 21, .	2.3	6
1183	The impact of obesity on lung function measurements and respiratory disease: A Mendelian randomization study. Annals of Human Genetics, 0, , .	0.3	0
1184	The relationship between adiposity and cognitive function: a bidirectional Mendelian randomization study in UK Biobank. International Journal of Epidemiology, 0, , .	0.9	0
1185	Genetics and epigenetics in the obesity phenotyping scenario. Reviews in Endocrine and Metabolic Disorders, 2023, 24, 775-793.	2.6	4
1187	Repurposing antidiabetic drugs for rheumatoid arthritis: results from a two-sample Mendelian randomization study. European Journal of Epidemiology, 2023, 38, 809-819.	2.5	6

#	Article	IF	CITATIONS
1188	The Influence of FAM13A and PPAR-Î ³ 2 Gene Polymorphisms on the Metabolic State of Postmenopausal Women. Genes, 2023, 14, 914.	1.0	0
1190	Pathophysiology of the Cardiometabolic Alterations in Obesity. , 2023, , 69-83.		0
1237	Genetics and Epigenetics in Obesity: What Do We Know so Far?. Current Obesity Reports, 2023, 12, 482-501.	3.5	1
1263	Sex differences in energy metabolism: natural selection, mechanisms and consequences. Nature Reviews Nephrology, 2024, 20, 56-69.	4.1	3
1291	White adipocyte dysfunction and obesity-associated pathologies in humans. Nature Reviews Molecular Cell Biology, 0 , , .	16.1	3
1293	Precision Nutrition and Obesity. , 2024, , 317-332.		O