## M-C Vohl

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5754207/publications.pdf

Version: 2024-02-01

318 papers 20,585 citations

54 h-index 128 g-index

334 all docs

334 docs citations

times ranked

334

27622 citing authors

#	Article	IF	CITATIONS
1	Genetic studies of body mass index yield new insights for obesity biology. Nature, 2015, 518, 197-206.	13.7	3,823
2	Defining the role of common variation in the genomic and biological architecture of adult human height. Nature Genetics, 2014, 46, 1173-1186.	9.4	1,818
3	The common PPAR $\hat{1}^3$ Pro12Ala polymorphism is associated with decreased risk of type 2 diabetes. Nature Genetics, 2000, 26, 76-80.	9.4	1,672
4	New genetic loci link adipose and insulin biology to body fat distribution. Nature, 2015, 518, 187-196.	13.7	1,328
5	A genome-wide approach accounting for body mass index identifies genetic variants influencing fasting glycemic traits and insulin resistance. Nature Genetics, 2012, 44, 659-669.	9.4	762
6	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
7	Precision Nutrition: A Review of Personalized Nutritional Approaches for the Prevention and Management of Metabolic Syndrome. Nutrients, 2017, 9, 913.	1.7	292
8	A Survey of Genes Differentially Expressed in Subcutaneous and Visceral Adipose Tissue in Men*. Obesity, 2004, 12, 1217-1222.	4.0	282
9	Single-cell analysis of human adipose tissue identifies depot- and disease-specific cell types. Nature Metabolism, 2020, 2, 97-109.	5.1	272
10	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. Nature Communications, 2016, 7, 10495.	5.8	245
11	A survey of genetic and epigenetic variation affecting human gene expression. Physiological Genomics, 2004, 16, 184-193.	1.0	228
12	Genetic Variants of <i>FTO</i> Influence Adiposity, Insulin Sensitivity, Leptin Levels, and Resting Metabolic Rate in the Quebec Family Study. Diabetes, 2008, 57, 1147-1150.	0.3	206
13	Differential methylation in glucoregulatory genes of offspring born before vs. after maternal gastrointestinal bypass surgery. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 11439-11444.	3.3	197
14	Differential epigenomic and transcriptomic responses in subcutaneous adipose tissue between low and high responders to caloric restriction. American Journal of Clinical Nutrition, 2010, 91, 309-320.	2.2	193
15	The PPAR-gamma P12A polymorphism modulates the relationship between dietary fat intake and components of the metabolic syndrome: results from the Québec Family Study. Clinical Genetics, 2003, 63, 109-116.	1.0	170
16	Abdominal Visceral Fat is Associated with a <i>Bcl</i> I Restriction Fragment Length Polymorphism at the Glucocorticoid Receptor Gene Locus. Obesity, 1997, 5, 186-192.	4.0	169
17	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. Nature Communications, 2017, 8, 14977.	5 <b>.</b> 8	169
18	5' Flanking Variants of Resistin Are Associated With Obesity. Diabetes, 2002, 51, 1629-1634.	0.3	158

#	Article	IF	Citations
19	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
20	Molecular scanning of the human PPARα gene: association of the L162V mutation with hyperapobetalipoproteinemia. Journal of Lipid Research, 2000, 41, 945-952.	2.0	155
21	Genome-wide meta-analysis uncovers novel loci influencing circulating leptin levels. Nature Communications, 2016, 7, 10494.	5.8	153
22	Glycerol: a neglected variable in metabolic processes?. BioEssays, 2001, 23, 534-542.	1.2	139
23	Molecular scanning of the human PPARa gene: association of the L162v mutation with hyperapobetalipoproteinemia. Journal of Lipid Research, 2000, 41, 945-52.	2.0	124
24	Contribution of abdominal obesity and hypertriglyceridemia to impaired fasting glucose and coronary artery disease. American Journal of Cardiology, 2002, 90, 15-18.	0.7	114
25	Novel loci associated with usual sleep duration: the CHARGE Consortium Genome-Wide Association Study. Molecular Psychiatry, 2015, 20, 1232-1239.	4.1	112
26	Associations between dietary patterns and obesity phenotypes. International Journal of Obesity, 2009, 33, 1419-1426.	1.6	108
27	Association between the PPARÎ $\pm$ -L162V polymorphism and components of the metabolic syndrome. Journal of Human Genetics, 2004, 49, 482-489.	1.1	105
28	Plasma <i>n</i> -3 fatty acid response to an <i>n</i> -3 fatty acid supplement is modulated by apoE É>4 but not by the common PPAR-α L162V polymorphism in men. British Journal of Nutrition, 2009, 102, 1121-1124.	1.2	98
29	Leptin and adiponectin DNA methylation levels in adipose tissues and blood cells are associated with BMI, waist girth and LDL-cholesterol levels in severely obese men and women. BMC Medical Genetics, 2015, 16, 29.	2.1	96
30	Relationships of Abdominal Obesity and Hyperinsulinemia to Angiographically Assessed Coronary Artery Disease in Men With Known Mutations in the LDL Receptor Gene. Circulation, 1998, 97, 871-877.	1.6	91
31	Association between insulin secretion, insulin sensitivity and type 2 diabetes susceptibility variants identified in genome-wide association studies. Acta Diabetologica, 2009, 46, 217-226.	1.2	91
32	A Novel Lecithin-Cholesterol Acyltransferase Antioxidant Activity Prevents the Formation of Oxidized Lipids during Lipoprotein Oxidationâ€. Biochemistry, 1999, 38, 5976-5981.	1.2	87
33	The interleukin 6 a^174G/C Polymorphism is associated with indices of obesity in men. Journal of Human Genetics, 2003, 48, 0014-0019.	1.1	84
34	Relation of the "Hypertriglyceridemic Waist―Phenotype to Earlier Manifestations of Coronary Artery Disease in Patients With Glucose Intolerance and Type 2 Diabetes Mellitus. American Journal of Cardiology, 2007, 99, 369-373.	0.7	84
35	Characterization of functional methylomes by next-generation capture sequencing identifies novel disease-associated variants. Nature Communications, 2015, 6, 7211.	5.8	84
36	Neuromedin $\hat{l}^2$ : a strong candidate gene linking eating behaviors and susceptibility to obesity. American Journal of Clinical Nutrition, 2004, 80, 1478-1486.	2.2	83

#	Article	IF	CITATIONS
37	Epigenetic changes in blood leukocytes following an omega-3 fatty acid supplementation. Clinical Epigenetics, 2017, 9, 43.	1.8	82
38	Common Polymorphisms in the Promoter of the Visfatin Gene (PBEF1) Influence Plasma Insulin Levels in a French-Canadian Population. Diabetes, 2006, 55, 2896-2902.	0.3	76
39	The metabolic signature associated with the Western dietary pattern: a cross-sectional study. Nutrition Journal, 2013, 12, 158.	1.5	76
40	Effect of n-3 fatty acids on the expression of inflammatory genes in THP-1 macrophages. Lipids in Health and Disease, 2016, 15, 69.	1.2	75
41	Disturbance in uniformly <sup>13</sup> C-labelled DHA metabolism in elderly human subjects carrying the apoE ε4 allele. British Journal of Nutrition, 2013, 110, 1751-1759.	1.2	74
42	A principal component meta-analysis on multiple anthropometric traits identifies novel loci for body shape. Nature Communications, 2016, 7, 13357.	5.8	74
43	Effect of apolipoprotein E, peroxisome proliferator-activated receptor alpha and lipoprotein lipase gene mutations on the ability of fenofibrate to improve lipid profiles and reach clinical guideline targets among hypertriglyceridemic patients. Pharmacogenetics and Genomics, 2002, 12, 313-320.	5.7	72
44	Functional variation in allelic methylomes underscores a strong genetic contribution and reveals novel epigenetic alterations in the human epigenome. Genome Biology, 2017, 18, 50.	3.8	71
45	Effects of 6-month vitamin D supplementation on insulin sensitivity and secretion: a randomised, placebo-controlled trial. European Journal of Endocrinology, 2019, 181, 287-299.	1.9	64
46	Transcriptomic and metabolomic signatures of an n-3 polyunsaturated fatty acids supplementation in a normolipidemic/normocholesterolemic Caucasian population. Journal of Nutritional Biochemistry, 2013, 24, 54-61.	1.9	63
47	Effect of liver fatty acid binding protein (FABP) T94A missense mutation on plasma lipoprotein responsiveness to treatment with fenofibrate. Journal of Human Genetics, 2004, 49, 424-432.	1.1	62
48	LINE-1 methylation in visceral adipose tissue of severely obese individuals is associated with metabolic syndrome status and related phenotypes. Clinical Epigenetics, 2012, 4, 10.	1.8	62
49	DNA variation in the genes of the Na,K-adenosine triphosphatase and its relation with resting metabolic rate, respiratory quotient, and body fat Journal of Clinical Investigation, 1994, 93, 838-843.	3.9	62
50	Influence of LDL receptor gene mutation and apo E polymorphism on lipoprotein response to simvastatin treatment among adolescents with heterozygous familial hypercholesterolemia. Atherosclerosis, 2002, 160, 361-368.	0.4	61
51	Comparison of the effect of two lowâ€density lipoprotein receptor class mutations on coronary heart disease among French anadian patients heterozygous for familial hypercholesterolaemia. European Journal of Clinical Investigation, 1997, 27, 366-373.	1.7	60
52	Genome-wide linkage scan reveals multiple susceptibility loci influencing lipid and lipoprotein levels in the Québec Family Study. Journal of Lipid Research, 2004, 45, 419-426.	2.0	60
53	Associations between dietary patterns and gene expression profiles of healthy men and women: a cross-sectional study. Nutrition Journal, 2013, 12, 24.	1.5	60
54	ZFP36: a Promising Candidate Gene for Obesity-Related Metabolic Complications Identified by Converging Genomics. Obesity Surgery, 2007, 17, 372-382.	1.1	57

#	Article	IF	CITATIONS
55	Biological plausibility for interactions between dietary fat, resveratrol, <i>ACE2</i> , and SARS-CoV illness severity. American Journal of Physiology - Endocrinology and Metabolism, 2020, 318, E830-E833.	1.8	57
56	Electronic health record-based genome-wide meta-analysis provides insights on the genetic architecture of non-alcoholic fatty liver disease. Cell Reports Medicine, 2021, 2, 100437.	3.3	56
57	ApoB-100 gene EcoRI polymorphism. Relations to plasma lipoprotein changes associated with abdominal visceral obesity Arteriosclerosis and Thrombosis: A Journal of Vascular Biology, 1994, 14, 527-533.	3.8	55
58	Methylation and Expression of Immune and Inflammatory Genes in the Offspring of Bariatric Bypass Surgery Patients. Journal of Obesity, 2013, 2013, 1-9.	1.1	55
59	Moderators of the intention-behaviour and perceived behavioural control-behaviour relationships for leisure-time physical activity. International Journal of Behavioral Nutrition and Physical Activity, 2008, 5, 7.	2.0	54
60	Gene–diet interactions on plasma lipid levels in the Inuit population. British Journal of Nutrition, 2013, 109, 953-961.	1.2	54
61	Glycerol as a Correlate of Impaired Glucose Tolerance: Dissection of a Complex System by Use of a Simple Genetic Trait. American Journal of Human Genetics, 2000, 66, 1558-1568.	2.6	53
62	Contribution of the cholesteryl ester transfer protein gene TaqIB polymorphism to the reduced plasma HDL-cholesterol levels found in abdominal obese men with the features of the insulin resistance syndrome. International Journal of Obesity, 1999, 23, 918-925.	1.6	52
63	The peroxisome proliferator-activated receptor $\hat{l}$ ± Leu $162$ Val polymorphism influences the metabolic response to a dietary intervention altering fatty acid proportions in healthy men. American Journal of Clinical Nutrition, 2005, 81, 523-530.	2.2	52
64	<i>DPP4</i> Gene DNA Methylation in the Omentum is Associated With Its Gene Expression and Plasma Lipid Profile in Severe Obesity. Obesity, 2011, 19, 388-395.	1.5	52
65	Association between Polymorphisms in the Fatty Acid Desaturase Gene Cluster and the Plasma Triacylglycerol Response to an n-3 PUFA Supplementation. Nutrients, 2012, 4, 1026-1041.	1.7	52
66	The effect of mere-measurement of cognitions on physical activity behavior: a randomized controlled trial among overweight and obese individuals. International Journal of Behavioral Nutrition and Physical Activity, 2011, 8, 2.	2.0	51
67	Association between yogurt consumption, dietary patterns, and cardio-metabolic risk factors. European Journal of Nutrition, 2016, 55, 577-587.	1.8	51
68	Contribution of receptor negative versus receptor defective mutations in the LDL-receptor gene to angiographically assessed coronary artery disease among young (25–49 years) versus middle-aged (50–64 years) men. Atherosclerosis, 1999, 143, 153-161.	0.4	50
69	Features of the metabolic syndrome are modulated by an interaction between the peroxisome proliferator-activated receptor-delta â`87T>C polymorphism and dietary fat in French-Canadians. International Journal of Obesity, 2007, 31, 411-417.	1.6	50
70	The T111I mutation in the EL gene modulates the impact of dietary fat on the HDL profile in women. Journal of Lipid Research, 2003, 44, 1902-1908.	2.0	49
71	Comprehensive genetic analysis of the dipeptidyl peptidase-4 gene and cardiovascular disease risk factors in obese individuals. Acta Diabetologica, 2009, 46, 13-21.	1.2	49
72	DNA methylation variations at CETP and LPL gene promoter loci: New molecular biomarkers associated with blood lipid profile variability. Atherosclerosis, 2013, 228, 413-420.	0.4	49

#	Article	IF	CITATIONS
73	Carotenoids as biomarkers of fruit and vegetable intake in men and women. British Journal of Nutrition, 2016, 116, 1206-1215.	1.2	48
74	The Lipoprotein Lipase Hin dlll Polymorphism Modulates Plasma Triglyceride Levels in Visceral Obesity. Arteriosclerosis, Thrombosis, and Vascular Biology, 1995, 15, 714-720.	1.1	48
75	Validation of the Use of Peripheral Blood Mononuclear Cells as Surrogate Model for Skeletal Muscle Tissue in Nutrigenomic Studies. OMICS A Journal of Integrative Biology, 2011, 15, 1-7.	1.0	47
76	Effects of FADS and ELOVL polymorphisms on indexes of desaturase and elongase activities: results from a pre-post fish oil supplementation. Genes and Nutrition, 2014, 9, 437.	1.2	47
77	Associations Between Dietary Protein Sources, Plasma BCAA and Short-Chain Acylcarnitine Levels in Adults. Nutrients, 2019, 11, 173.	1.7	47
78	Genome-wide association study of the plasma triglyceride response to an n-3 polyunsaturated fatty acid supplementation. Journal of Lipid Research, 2014, 55, 1245-1253.	2.0	44
79	Genome-wide meta-analysis of macronutrient intake of 91,114 European ancestry participants from the cohorts for heart and aging research in genomic epidemiology consortium. Molecular Psychiatry, 2019, 24, 1920-1932.	4.1	44
80	Plasma concentrations of apolipoprotein B are modulated by a gene–diet interaction effect between the LFABP T94A polymorphism and dietary fat intake in French-Canadian men. Molecular Genetics and Metabolism, 2004, 82, 296-303.	0.5	43
81	Differential methylation in visceral adipose tissue of obese men discordant for metabolic disturbances. Physiological Genomics, 2014, 46, 216-222.	1.0	43
82	Impact of adiponectin gene polymorphisms on plasma lipoprotein and adiponectin concentrations of viscerally obese men. Journal of Lipid Research, 2005, 46, 237-244.	2.0	42
83	Chylomicron retention disease: A long term study of two cohorts. Molecular Genetics and Metabolism, 2009, 97, 136-142.	0.5	42
84	Human resistin gene polymorphism is associated with visceral obesity and fasting and oral glucose stimulated C-peptide in the Québec Family Study. Journal of Endocrinological Investigation, 2004, 27, 1003-1009.	1.8	41
85	Association between olfactory receptor genes, eating behavior traits and adiposity: Results from the Quebec Family Study. Physiology and Behavior, 2012, 105, 772-776.	1.0	41
86	<i>ADRB3</i> gene promoter DNA methylation in blood and visceral adipose tissue isÂassociated with metabolic disturbances in men. Epigenomics, 2014, 6, 33-43.	1.0	41
87	Kinetics of 13C-DHA before and during fish-oil supplementation in healthy older individuals. American Journal of Clinical Nutrition, 2014, 100, 105-112.	2.2	40
88	Visceral obesity attenuates the effect of the hepatic lipase â^'514C>T polymorphism on plasma HDL-cholesterol levels in French-Canadian men. Molecular Genetics and Metabolism, 2003, 78, 31-36.	0.5	39
89	Fish nutrients decrease expression levels of tumor necrosis factor-α in cultured human macrophages. Physiological Genomics, 2010, 40, 189-194.	1.0	38
90	Differences in metabolomic and transcriptomic profiles between responders and non-responders to an n-3 polyunsaturated fatty acids (PUFAs) supplementation. Genes and Nutrition, 2013, 8, 411-423.	1.2	38

#	Article	IF	CITATIONS
91	Evidence for a Major Quantitative Trait Locus on Chromosome 17q21 Affecting Low-Density Lipoprotein Peak Particle Diameter. Circulation, 2003, 107, 2361-2368.	1.6	37
92	Genetics of LDL particle heterogeneity. Journal of Lipid Research, 2004, 45, 1008-1026.	2.0	37
93	Effects of Age, Sex, Body Mass Index and APOE Genotype on Cardiovascular Biomarker Response to an n-3 Polyunsaturated Fatty Acid Supplementation. Journal of Nutrigenetics and Nutrigenomics, 2013, 6, 73-82.	1.8	37
94	Hyperinsulinemia and Abdominal Obesity Affect the Expression of Hypertriglyceridemia in Heterozygous Familial Lipoprotein Lipase Deficiency. Diabetes, 1997, 46, 2063-2068.	0.3	36
95	Relative contribution of low-density lipoprotein receptor and lipoprotein lipase gene mutations to angiographically assessed coronary artery disease among French Canadians. American Journal of Cardiology, 1998, 82, 299-305.	0.7	36
96	Influences of the PPARÎ $\pm$ -L162V polymorphism on plasma HDL2-cholesterol response of abdominally obese men treated with gemfibrozil. Genetics in Medicine, 2002, 4, 311-315.	1.1	36
97	Geographic distribution of Frenchâ€Canadian lowâ€density lipoprotein receptor gene mutations in the Province of Quebec. Clinical Genetics, 1997, 52, 1-6.	1.0	36
98	Natural Rumenâ€Derived <i>trans</i> Fatty Acids Are Associated with Metabolic Markers of Cardiac Health. Lipids, 2015, 50, 873-882.	0.7	36
99	Development and validation of a nutrition knowledge questionnaire for a Canadian population. Public Health Nutrition, 2017, 20, 1184-1192.	1.1	36
100	Visceral obesity and hyperinsulinemia modulate the impact of the microsomal triglyceride transfer protein â~493G/T polymorphism on plasma lipoprotein levels in men. Atherosclerosis, 2002, 160, 317-324.	0.4	35
101	Profiling Serum Bile Acid Glucuronides in Humans: Gender Divergences, Genetic Determinants, and Response to Fenofibrate. Clinical Pharmacology and Therapeutics, 2013, 94, 533-543.	2.3	35
102	Nutrigenomics–Âperspectives from registered dietitians: a report from the Quebecâ€wide eâ€consultation on nutrigenomics among registered dietitians. Journal of Human Nutrition and Dietetics, 2014, 27, 391-400.	1.3	35
103	Dairy Product Consumption Has No Impact on Biomarkers of Inflammation among Men and Women with Low-Grade Systemic Inflammation. Journal of Nutrition, 2014, 144, 1760-1767.	1.3	34
104	Detection of a novel mutation (stop 468) in exon 10 of the low-density lipoprotein receptor gene causing familial hypercholesterolemia among French Canadians. Human Molecular Genetics, 1994, 3, 1689-1691.	1.4	33
105	Effect of an Oat Bran-Rich Supplement on the Metabolic Profile of Overweight Premenopausal Women. Annals of Nutrition and Metabolism, 2005, 49, 141-148.	1.0	33
106	Variants within the muscle and liver isoforms of the carnitine palmitoyltransferase I (CPT1) gene interact with fat intake to modulate indices of obesity in French-Canadians. Journal of Molecular Medicine, 2007, 85, 129-137.	1.7	33
107	Association between Metabolite Profiles, Metabolic Syndrome and Obesity Status. Nutrients, 2016, 8, 324.	1.7	33
108	Omega-3 fatty acids status in human subjects estimated using a food frequency questionnaire and plasma phospholipids levels. Nutrition Journal, 2012, 11, 46.	1.5	32

#	Article	IF	Citations
109	A Study of the Differential Effects of Eicosapentaenoic Acid (EPA) and Docosahexaenoic Acid (DHA) on Gene Expression Profiles of Stimulated Thp-1 Macrophages. Nutrients, 2017, 9, 424.	1.7	32
110	Association of <i>OSBPL11</i> Gene Polymorphisms With Cardiovascular Disease Risk Factors in Obesity. Obesity, 2009, 17, 1466-1472.	1.5	31
111	Association between polymorphisms in phospholipase A2 genes and the plasma triglyceride response to an n-3 PUFA supplementation: a clinical trial. Lipids in Health and Disease, 2015, 14, 12.	1.2	31
112	Cross-tissue comparisons of leptin and adiponectin. Adipocyte, 2014, 3, 132-140.	1.3	30
113	Supplementation with Resveratrol and Curcumin Does Not Affect the Inflammatory Response to a High-Fat Meal in Older Adults with Abdominal Obesity: A Randomized, Placebo-Controlled Crossover Trial. Journal of Nutrition, 2018, 148, 379-388.	1.3	30
114	Body mass index is associated with epigenetic age acceleration in the visceral adipose tissue of subjects with severe obesity. Clinical Epigenetics, 2019, 11, 172.	1.8	30
115	Polygenic risk score for predicting weight loss after bariatric surgery. JCI Insight, 2018, 3, .	2.3	30
116	Molecular Screening of the 11βâ€HSD1 Gene in Men Characterized by the Metabolic Syndrome. Obesity, 2004, 12, 1570-1575.	4.0	29
117	Effects of the peroxisome proliferator-activated receptor- $\hat{I}^3$ co-activator-1 Gly482Ser variant on features of the metabolic syndrome. Molecular Genetics and Metabolism, 2005, 86, 300-306.	0.5	29
118	Quantitative Trait Locus on 15q for a Metabolic Syndrome Variable Derived from Factor Analysis*. Obesity, 2007, 15, 544-550.	1.5	29
119	<i>PPARα</i> L162V polymorphism alters the potential of nâ€3 fatty acids to increase lipoprotein lipase activity. Molecular Nutrition and Food Research, 2010, 54, 543-550.	1.5	29
120	Effect of Implementation Intentions to Change Behaviour: Moderation by Intention Stability. Psychological Reports, 2010, 106, 147-159.	0.9	29
121	A variant in the <i>LRRFIP1</i> gene is associated with adiposity and inflammation. Obesity, 2013, 21, 185-192.	1.5	29
122	Poor Adherence to Dietary Guidelines Among French-Speaking Adults in the Province of Quebec, Canada: The PREDISE Study. Canadian Journal of Cardiology, 2018, 34, 1665-1673.	0.8	29
123	Circulating glutamate level as a potential biomarker for abdominal obesity and metabolic risk. Nutrition, Metabolism and Cardiovascular Diseases, 2019, 29, 1353-1360.	1.1	29
124	The lipoprotein/lipid profile is modulated by a geneâ€"diet interaction effect between polymorphisms in the liver X receptor-α and dietary cholesterol intake in French-Canadians. British Journal of Nutrition, 2007, 97, 11-18.	1.2	28
125	Prediction of daily fruit and vegetable consumption among overweight and obese individuals. Appetite, 2010, 54, 480-484.	1.8	28
126	The Peroxisome Proliferatorâ€Activated Receptor α L162V Mutation Is Associated with Reduced Adiposity. Obesity, 2003, 11, 809-816.	4.0	27

#	Article	IF	CITATIONS
127	Visceral adipose tissue zinc finger protein 36 mRNA levels are correlated with insulin, insulin resistance index, and adiponectinemia in women. European Journal of Endocrinology, 2007, 157, 451-457.	1.9	27
128	Evidence of Interaction between Type 2 Diabetes Susceptibility Genes and Dietary Fat Intake for Adiposity and Glucose Homeostasis-Related Phenotypes. Journal of Nutrigenetics and Nutrigenomics, 2009, 2, 225-234.	1.8	27
129	Compendium of genome-wide scans of lipid-related phenotypes. Journal of Lipid Research, 2004, 45, 2174-2184.	2.0	26
130	Validity of a self-reported measure of familial history of obesity. Nutrition Journal, 2008, 7, 27.	1.5	26
131	An explained varianceâ€based genetic risk score associated with gestational diabetes antecedent and with progression to preâ€diabetes and type 2 diabetes: a cohort study. BJOG: an International Journal of Obstetrics and Gynaecology, 2015, 122, 411-419.	1.1	26
132	Rapid restriction fragment analysis for screening four point mutations of the Low-density lipoprotein receptor gene in French Canadians. Human Mutation, 1995, 6, 243-246.	1.1	25
133	Heterozygous familial hypercholesterolemia in children: low-density lipoprotein receptor mutational analysis and variation in the expression of plasma lipoprotein-lipid concentrations. Atherosclerosis, 1996, 126, 163-171.	0.4	25
134	Identification of three mutations in the low-density lipoprotein receptor gene causing familial hypercholesterolemia among French Canadians. Human Mutation, 1998, 11, S226-S231.	1.1	25
135	Plasminogen-activator inhibitor-1 polymorphisms are associated with obesity and fat distribution in the Qu??bec Family Study: evidence of interactions with menopause. Menopause, 2005, 12, 136-143.	0.8	25
136	Risks of nutrigenomics and nutrigenetics? What the scientists say. Genes and Nutrition, 2014, 9, 370.	1.2	25
137	Docosahexaenoic acid-enriched canola oil increases adiponectin concentrations: A randomized crossover controlled intervention trial. Nutrition, Metabolism and Cardiovascular Diseases, 2015, 25, 52-59.	1.1	25
138	Phosphoinositide cycle gene polymorphisms affect the plasma lipid profile in the Quebec Family Study. Molecular Genetics and Metabolism, 2009, 97, 149-154.	0.5	24
139	GAD2 gene sequence variations are associated with eating behaviors and weight gain in women from the Quebec family study. Physiology and Behavior, 2009, 98, 505-510.	1.0	24
140	Effects of Peroxisome Proliferator-Activated Receptors, Dietary Fat Intakes and Gene–Diet Interactions on Peak Particle Diameters of Low-Density Lipoproteins. Journal of Nutrigenetics and Nutrigenomics, 2011, 4, 36-48.	1.8	24
141	Interaction between Common Genetic Variants and Total Fat Intake on Low-Density Lipoprotein Peak Particle Diameter: A Genome-Wide Association Study. Journal of Nutrigenetics and Nutrigenomics, 2015, 8, 44-53.	1.8	24
142	Fine mapping of genome-wide association study signals to identify genetic markers of the plasma triglyceride response to an omega-3 fatty acid supplementation. American Journal of Clinical Nutrition, 2019, 109, 176-185.	2.2	24
143	Effects of the <i>FABP2</i> A54T Mutation on Triglyceride Metabolism of Viscerally Obese Men. Obesity, 2001, 9, 668-675.	4.0	23
144	Associations between glucose tolerance, insulin sensitivity and insulin secretion phenotypes and polymorphisms in adiponectin and adiponectin receptor genes in the Quebec Family Study. Diabetic Medicine, 2008, 25, 400-406.	1.2	23

#	Article	IF	Citations
145	Prediction of Leisureâ€time Physical Activity Among Obese Individuals. Obesity, 2009, 17, 706-712.	1.5	23
146	Association of <i>LIPA</i> Gene Polymorphisms With Obesityâ€Related Metabolic Complications Among Severely Obese Patients. Obesity, 2012, 20, 2075-2082.	1.5	23
147	Cardiometabolic risk factors are influenced by <i><scp>S</scp>tearoylâ€<scp>C</scp>o<scp>A D</scp>esaturase</i> ( <i><scp>SCD</scp></i> ) â°' <i>&lt;1</i> ) gene polymorphisms and <i><i><i><i><i>&lt;5</i></i></i></i></i>	1.5	23
148	The Mspl polymorphism of the apolipoprotein A-II gene as a modulator of the dyslipidemic state found in visceral obesity. Atherosclerosis, 1997, 128, 183-190.	0.4	22
149	Fine mapping of low-density lipoprotein receptor gene by genetic linkage on chromosome 19p13.1-p13.3 and study of the founder effect of four French Canadian low-density lipoprotein receptor gene mutations. Atherosclerosis, 1999, 143, 145-151.	0.4	22
150	The c.419-420insA in the MTP gene is associated with abetalipoproteinemia among French-Canadians. Molecular Genetics and Metabolism, 2004, 81, 140-143.	0.5	22
151	A Simple Method to Assess Fruit and Vegetable Intake among Obese and Non-obese Individuals. Canadian Journal of Public Health, 2008, 99, 494-498.	1.1	22
152	Genes, Fat Intake, and Cardiovascular Disease Risk Factors in the Quebec Family Study. Obesity, 2007, 15, 2336-2347.	1.5	21
153	Contribution of Genetic and Metabolic Syndrome to Omental Adipose Tissue PAI-1 Gene mRNA and Plasma Levels in Obesity. Obesity Surgery, 2010, 20, 492-499.	1.1	21
154	Effects of a Supplementation of n-3 Polyunsaturated Fatty Acids with or without Fish Gelatin on Gene Expression in Peripheral Blood Mononuclear Cells in Obese, Insulin-Resistant Subjects. Journal of Nutrigenetics and Nutrigenomics, 2011, 4, 192-202.	1.8	21
155	Effect of different concentrations of omega-3 fatty acids on stimulated THP-1 macrophages. Genes and Nutrition, 2017, 12, 7.	1.2	21
156	Yogurt consumption, body composition, and metabolic health in the Québec Family Study. European Journal of Nutrition, 2018, 57, 1591-1603.	1.8	21
157	Relation Between <i>BgllI</i> Polymorphism in 3βâ€Hydroxysteroid Dehydrogenase Gene and Adipose Tissue Distribution in Humans. Obesity, 1994, 2, 444-449.	4.0	20
158	Population prevalence of APOE, APOC3 and PPAR-α mutations associated to hypertriglyceridemia in French Canadians. Journal of Human Genetics, 2004, 49, 691-700.	1.1	20
159	Dietary patterns and associated lifestyles in individuals with and without familial history of obesity: a cross-sectional study. International Journal of Behavioral Nutrition and Physical Activity, 2006, 3, 38.	2.0	20
160	Evidence of a quantitative trait locus for energy and macronutrient intakes on chromosome 3q27.3: the QuÃ@bec Family Study. American Journal of Clinical Nutrition, 2008, 88, 1142-1148.	2.2	20
161	Genetic contribution to C-reactive protein levels in severe obesity. Molecular Genetics and Metabolism, 2012, 105, 494-501.	0.5	20
162	Polymorphisms, de novo lipogenesis, and plasma triglyceride response following fish oil supplementation. Journal of Lipid Research, 2013, 54, 2866-2873.	2.0	20

#	Article	IF	CITATIONS
163	Genetic regulation of differentially methylated genes in visceral adipose tissue of severely obese men discordant for the metabolic syndrome. Translational Research, 2017, 184, 1-11.e2.	2.2	20
164	Effects of Daily Raspberry Consumption on Immune-Metabolic Health in Subjects at Risk of Metabolic Syndrome: A Randomized Controlled Trial. Nutrients, 2020, 12, 3858.	1.7	20
165	Omega-3 fatty acids, polymorphisms and lipid related cardiovascular disease risk factors in the Inuit population. Nutrition and Metabolism, 2013, 10, 26.	1.3	19
166	Polymorphisms in Fatty Acid Desaturase (FADS) Gene Cluster: Effects on Glycemic Controls Following an Omega-3 Polyunsaturated Fatty Acids (PUFA) Supplementation. Genes, 2013, 4, 485-498.	1.0	19
167	Polymorphisms in Genes Involved in Fatty Acid $\hat{l}^2$ -Oxidation Interact with Dietary Fat Intakes to Modulate the Plasma TG Response to a Fish Oil Supplementation. Nutrients, 2014, 6, 1145-1163.	1.7	19
168	Novel Genetic Loci Associated with the Plasma Triglyceride Response to an Omega-3 Fatty Acid Supplementation. Journal of Nutrigenetics and Nutrigenomics, 2016, 9, 1-11.	1.8	19
169	Correlates of the difference in plasma carotenoid concentrations between men and women. British Journal of Nutrition, 2019, 121, 172-181.	1.2	19
170	Associations Between Nutrition Knowledge and Overall Diet Quality: The Moderating Role of Sociodemographic Characteristics—Results From the PREDISE Study. American Journal of Health Promotion, 2021, 35, 38-47.	0.9	19
171	The ApoB-100 Gene <i>Eco</i> RI Polymorphism Influences the Relationship Between Features of the Insulin Resistance Syndrome and the Hyper-ApoB and Dense LDL Phenotype in Men. Diabetes, 1996, 45, 1405-1411.	0.3	18
172	Heritability of LDL peak particle diameter in the Quebec Family Study. Genetic Epidemiology, 2003, 25, 375-381.	0.6	18
173	Low plasma adiponectin exacerbates the risk of premature coronary artery disease in familial hypercholesterolemia. Atherosclerosis, 2008, 196, 262-269.	0.4	18
174	Associations between dairy intake and metabolic risk parameters in a healthy French-Canadian population. Applied Physiology, Nutrition and Metabolism, 2014, 39, 1323-1331.	0.9	18
175	Expression and Sequence Variants of Inflammatory Genes; Effects on Plasma Inflammation Biomarkers Following a 6-Week Supplementation with Fish Oil. International Journal of Molecular Sciences, 2016, 17, 375.	1.8	18
176	Familial resemblances in human plasma metabolites are attributable to both genetic and common environmental effects. Nutrition Research, 2019, 61, 22-30.	1.3	18
177	Guiding Global Best Practice in Personalized Nutrition Based on Genetics: The Development of a Nutrigenomics Care Map. Journal of the Academy of Nutrition and Dietetics, 2022, 122, 259-269.	0.4	18
178	Effect of the PPAR-Alpha L162V Polymorphism on the Cardiovascular Disease Risk Factor in Response to n–3 Polyunsaturated Fatty Acids. Journal of Nutrigenetics and Nutrigenomics, 2008, 1, 205-212.	1.8	17
179	Network Analysis of the Potential Role of DNA Methylation in the Relationship between Plasma Carotenoids and Lipid Profile. Nutrients, 2019, 11, 1265.	1.7	17
180	Evidence for Interaction betweenPPARGPro12Ala andPPARGC1AGly482Ser Polymorphisms in Determining Type 2 Diabetes Intermediate Phenotypes in Overweight Subjects. Experimental and Clinical Endocrinology and Diabetes, 2009, 117, 455-459.	0.6	16

#	Article	IF	Citations
181	Thymic stromal lymphopoietin: an immune cytokine gene associated with the metabolic syndrome and blood pressure in severe obesity. Clinical Science, 2012, 123, 99-109.	1.8	16
182	Genome-Wide Association Study of Dietary Pattern Scores. Nutrients, 2017, 9, 649.	1.7	16
183	Dissecting features of epigenetic variants underlying cardiometabolic risk using full-resolution epigenome profiling in regulatory elements. Nature Communications, 2019, 10, 1209.	5.8	16
184	Altered branched-chain $\hat{l}_{\pm}$ -keto acid metabolism is a feature of NAFLD in individuals with severe obesity. JCI Insight, 2022, 7, .	2.3	16
185	The pleiotropic expression of the myotonic dystrophy protein kinase gene illustrates the complex relationships between genetic, biological and clinical covariates of male aging. Aging Male, 2002, 5, 223-232.	0.9	15
186	Effect of apoC-III gene polymorphisms on the lipoprotein-lipid profile of viscerally obese men. Journal of Lipid Research, 2003, 44, 986-993.	2.0	15
187	Molecular screening of the microsomal triglyceride transfer protein: association between polymorphisms and both abdominal obesity and plasma apolipoprotein B concentration. Journal of Human Genetics, 2004, 49, 684-690.	1.1	15
188	Associations between USF1 gene variants and cardiovascular risk factors in the Quebec Family Study. Clinical Genetics, 2007, 71, 245-253.	1.0	15
189	Methylation quantitative trait loci within the TOMM20 gene are associated with metabolic syndrome-related lipid alterations in severely obese subjects. Diabetology and Metabolic Syndrome, 2016, 8, 55.	1.2	15
190	Development and Validation of the Food Liking Questionnaire in a French-Canadian Population. Nutrients, 2017, 9, 1337.	1.7	15
191	Plasma Triglyceride Levels May Be Modulated by Gene Expression of IQCJ, NXPH1, PHF17 and MYB in Humans. International Journal of Molecular Sciences, 2017, 18, 257.	1.8	15
192	Mendelian Randomization Analysis Identifies Blood Tyrosine Levels as a Biomarker of Non-Alcoholic Fatty Liver Disease. Metabolites, 2022, 12, 440.	1.3	15
193	Detection of a major gene effect for LDL peak particle diameter and association with apolipoprotein H gene haplotype. Atherosclerosis, 2005, 182, 231-239.	0.4	14
194	Differences in Transcriptional Activation by the Two Allelic (L162V Polymorphic) Variants of PPAR <mml:math xmlns:mml="http://www.w3.org/1998/Math/MathML"><mml:mi>α</mml:mi></mml:math> after Omega-3 Fatty Acids Treatment. PPAR Research, 2009, 2009, 1-5.	1.1	14
195	Comparison of the dipeptidyl peptidase-4 gene methylation levels between severely obese subjects with and without the metabolic syndrome. Diabetology and Metabolic Syndrome, 2013, 5, 4.	1.2	14
196	PPAR <b><i>α</i></b> : A Master Regulator of Bilirubin Homeostasis. PPAR Research, 2014, 2014, 1-11.	1.1	14
197	Haplotypes in the phospholipid transfer protein gene are associated with obesity-related phenotypes: the QuA@bec Family Study. International Journal of Obesity, 2005, 29, 1338-1345.	1.6	13
198	Gene Expression Variability in Subcutaneous and Omental Adipose Tissue of Obese Men. Gene Expression, 2007, 14, 35-46.	0.5	13

#	Article	IF	CITATIONS
199	<i>CYR61 </i> polymorphisms are associated with plasma HDLâ€cholesterol levels in obese individuals. Clinical Genetics, 2007, 72, 224-229.	1.0	13
200	Omega-3 fatty acids regulate gene expression levels differently in subjects carrying the PPAR $\hat{l}\pm$ L162V polymorphism. Genes and Nutrition, 2009, 4, 199-205.	1.2	13
201	A polymorphism of the interferon-gamma-inducible protein 30 gene is associated with hyperglycemia in severely obese individuals. Human Genetics, 2012, 131, 57-66.	1.8	13
202	Nutrigenetic Testing for Personalized Nutrition: An Evaluation of Public Perceptions, Attitudes, and Concerns in a Population of French Canadians. Lifestyle Genomics, 2018, 11, 155-162.	0.6	13
203	Hyperinsulinemia and abdominal obesity affect the expression of hypertriglyceridemia in heterozygous familial lipoprotein lipase deficiency. Diabetes, 1997, 46, 2063-2068.	0.3	13
204	Influences of the phosphatidylcholine transfer protein gene variants on the LDL peak particle size. Atherosclerosis, 2007, 195, 297-302.	0.4	12
205	A CpG-SNP Located within the <b><i>ARPC3</i></b> Gene Promoter Is Associated with Hypertriglyceridemia in Severely Obese Patients. Annals of Nutrition and Metabolism, 2016, 68, 203-212.	1.0	12
206	A common variant in ARHGEF10 alters delta-6 desaturase activity and influence susceptibility to hypertriglyceridemia. Journal of Clinical Lipidology, 2018, 12, 311-320.e3.	0.6	12
207	Gut Microbial Signatures of Distinct Trimethylamine N-Oxide Response to Raspberry Consumption. Nutrients, 2022, 14, 1656.	1.7	12
208	Genetic epistasis in the VLDL catabolic pathway is associated with deleterious variations on triglyceridemia in obese subjects. International Journal of Obesity, 2007, 31, 1325-1333.	1.6	11
209	Combining genetic markers and clinical risk factors improves the risk assessment of impaired glucose metabolism. Annals of Medicine, 2010, 42, 196-206.	1.5	11
210	SREBF1 gene variations modulate insulin sensitivity in response to a fish oil supplementation. Lipids in Health and Disease, 2014, 13, 152.	1.2	11
211	Gene-diet interactions with polymorphisms of the MGLL gene on plasma low-density lipoprotein cholesterol and size following an omega-3 polyunsaturated fatty acid supplementation: a clinical trial. Lipids in Health and Disease, 2014, 13, 86.	1.2	11
212	Polymorphisms in FFAR4 (GPR120) Gene Modulate Insulin Levels and Sensitivity after Fish Oil Supplementation. Journal of Personalized Medicine, 2017, 7, 15.	1.1	11
213	Temporal Changes in Gene Expression Profile during Mature Adipocyte Dedifferentiation. International Journal of Genomics, 2017, 2017, 1-11.	0.8	11
214	Social Support, but Not Perceived Food Environment, Is Associated with Diet Quality in French-Speaking Canadians from the PREDISE Study. Nutrients, 2019, 11, 3030.	1.7	11
215	Animal and Cellular Studies Demonstrate Some of the Beneficial Impacts of Herring Milt Hydrolysates on Obesity-Induced Glucose Intolerance and Inflammation. Nutrients, 2020, 12, 3235.	1.7	11
216	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

#	Article	IF	CITATIONS
217	Influence of the angiotensin-converting enzyme gene insertion/deletion polymorphism on lipoprotein/lipid response to gemfibrozil. Clinical Genetics, 2002, 62, 45-52.	1.0	10
218	Combined effects of PPARÎ <sup>3</sup> 2 P12A and PPARα L162V polymorphisms on glucose and insulin homeostasis: the Québec Family Study. Journal of Human Genetics, 2003, 48, 614-621.	1.1	10
219	Interaction between HNF4A polymorphisms and physical activity in relation to type 2 diabetes-related traits: Results from the Quebec Family Study. Diabetes Research and Clinical Practice, 2009, 84, 211-218.	1.1	10
220	Interactions between Dietary Fat Intake and FASN Genetic Variation Influence LDL Peak Particle Diameter. Journal of Nutrigenetics and Nutrigenomics, 2011, 4, 137-145.	1.8	10
221	<i>DUSP1</i> Gene Polymorphisms Are Associated with Obesity-Related Metabolic Complications among Severely Obese Patients and Impact on Gene Methylation and Expression. International Journal of Genomics, 2013, 2013, 1-10.	0.8	10
222	Familial resemblances in blood leukocyte DNA methylation levels. Epigenetics, 2016, 11, 831-838.	1.3	10
223	N-3 Polyunsaturated Fatty Acids Stimulate Bile Acid Detoxification in Human Cell Models. Canadian Journal of Gastroenterology and Hepatology, 2018, 2018, 1-12.	0.8	10
224	Individuals with self-determined motivation for eating have better overall diet quality: Results from the PREDISE study. Appetite, 2021, 165, 105426.	1.8	10
225	Nutrigenetics, omega-3 and plasma lipids/lipoproteins/apolipoproteins with evidence evaluation using the GRADE approach: a systematic review. BMJ Open, 2022, 12, e054417.	0.8	10
226	HDL Cholesterol and TaqIB Cholesteryl Ester Transfer Protein Gene Polymorphism in Renal Transplant Recipients. Nephron, 2000, 84, 333-341.	0.9	9
227	A Sequence Variation in the Mitochondrial Glycerol-3-Phosphate Dehydrogenase Gene Is Associated with Increased Plasma Glycerol and Free Fatty Acid Concentrations among French Canadians. Molecular Genetics and Metabolism, 2001, 72, 209-217.	0.5	9
228	Contribution of Hierarchical Clustering Techniques to the Modeling of the Geographic Distribution of Genetic Polymorphisms Associated with Chronic Inflammatory Diseases in the Québec Population. Public Health Genomics, 2007, 10, 218-226.	0.6	9
229	Apolipoprotein E and lipoprotein lipase gene polymorphisms interaction on the atherogenic combined expression of hypertriglyceridemia and hyperapobetalipoproteinemia phenotypes. Journal of Endocrinological Investigation, 2007, 30, 551-557.	1.8	9
230	LIPE C-60G influences the effects of physical activity on body fat and plasma lipid concentrations: the Quebec Family Study. Human Genomics, 2009, 3, 157.	1.4	9
231	Dairy Product Consumption Interacts with Glucokinase (GCK) Gene Polymorphisms Associated with Insulin Resistance. Journal of Personalized Medicine, 2017, 7, 8.	1.1	9
232	Consumption of low nutritive value foods and cardiometabolic risk factors among French-speaking adults from Quebec, Canada: the PREDISE study. Nutrition Journal, 2019, 18, 49.	1.5	9
233	Weighted gene co-expression network analysis to explain the relationship between plasma total carotenoids and lipid profile. Genes and Nutrition, 2019, 14, 16.	1.2	9
234	Raspberry consumption: identification of distinct immune-metabolic response profiles by whole blood transcriptome profiling. Journal of Nutritional Biochemistry, 2022, 101, 108946.	1.9	9

#	Article	IF	Citations
235	Associations between Polymorphisms in Genes Involved in Fatty Acid Metabolism and Dietary Fat Intakes. Journal of Nutrigenetics and Nutrigenomics, 2012, 5, 1-12.	1.8	8
236	Polymorphisms in Genes Involved in the Triglyceride Synthesis Pathway and Marine Omega-3 Polyunsaturated Fatty Acid Supplementation Modulate Plasma Triglyceride Levels. Journal of Nutrigenetics and Nutrigenomics, 2013, 6, 268-280.	1.8	8
237	Current knowledge and interest of French Canadians regarding nutrigenetics. Genes and Nutrition, 2019, 14, 5.	1.2	8
238	An 8-week freeze-dried blueberry supplement impacts immune-related pathways: a randomized, double-blind placebo-controlled trial. Genes and Nutrition, 2021, 16, 7.	1.2	8
239	Salmon peptides limit obesityâ€associated metabolic disorders by modulating a gutâ€liver axis in vitamin Dâ€deficient mice. Obesity, 2021, 29, 1635-1649.	1.5	8
240	Associations between Dietary Patterns and LDL Peak Particle Diameter: A Cross-Sectional Study. Journal of the American College of Nutrition, 2010, 29, 630-637.	1.1	7
241	Fine Mapping of the Insulin-Induced Gene 2 Identifies a Variant Associated With LDL Cholesterol and Total Apolipoprotein B Levels. Circulation: Cardiovascular Genetics, 2010, 3, 454-461.	5.1	7
242	Transcriptomic profiles of skeletal muscle tissue following an euglycemic-hyperinsulinemic clamp in insulin-resistant obese subjects. Genes and Nutrition, 2013, 8, 91-98.	1.2	7
243	Modulation of C-reactive protein and plasma omega-6 fatty acid levels by phospholipase A2 gene polymorphisms following a 6-week supplementation with fish oil. Prostaglandins Leukotrienes and Essential Fatty Acids, 2015, 102-103, 37-45.	1.0	7
244	Ethical considerations in the implementation of nutrigenetics/nutrigenomics. Personalized Medicine, 2017, 14, 75-83.	0.8	7
245	Genetic and Common Environmental Contributions to Familial Resemblances in Plasma Carotenoid Concentrations in Healthy Families. Nutrients, 2018, 10, 1002.	1.7	7
246	Associations between self-reported vegetable and fruit intake assessed with a new web-based 24-h dietary recall and serum carotenoids in free-living adults: a relative validation study. Journal of Nutritional Science, 2019, 8, e26.	0.7	7
247	Intakes of Total, Free, and Naturally Occurring Sugars in the French-Speaking Adult Population of the Province of Québec, Canada: The PREDISE Study. Nutrients, 2019, 11, 2317.	1.7	7
248	Prevention of Potential Adverse Metabolic Effects of a Supplementation with Omega-3 Fatty Acids Using a Genetic Score Approach. Lifestyle Genomics, 2020, 13, 32-42.	0.6	7
249	Are Machine Learning Algorithms More Accurate in Predicting Vegetable and Fruit Consumption Than Traditional Statistical Models? An Exploratory Analysis. Frontiers in Nutrition, 2022, 9, 740898.	1.6	7
250	The Genetic and Metabolic Determinants of Cardiovascular Complications in Type 2 Diabetes: Recent Insights from Animal Models and Clinical Investigations. Canadian Journal of Diabetes, 2013, 37, 351-358.	0.4	6
251	An interaction effect between glucokinase gene variation and carbohydrate intakes modulates the plasma triglyceride response to a fish oil supplementation. Genes and Nutrition, 2014, 9, 395.	1.2	6
252	Remodeling adipose tissue through in silico modulation of fat storage for the prevention of type 2 diabetes. BMC Systems Biology, 2017, 11, 60.	3.0	6

#	Article	IF	CITATIONS
253	Social support for healthy eating: development and validation of a questionnaire for the French-Canadian population. Public Health Nutrition, 2018, 21, 2360-2366.	1.1	6
254	Genetic Risk Score Predictive of the Plasma Triglyceride Response to an Omega-3 Fatty Acid Supplementation in a Mexican Population. Nutrients, 2019, 11, 737.	1.7	6
255	Genetic risk prediction of the plasma triglyceride response to independent supplementations with eicosapentaenoic and docosahexaenoic acids: the ComparED Study. Genes and Nutrition, 2020, 15, 10.	1.2	6
256	Identification of Phenotypic Lipidomic Signatures in Response to Long Chain nâ€3 Polyunsaturated Fatty Acid Supplementation in Humans. Journal of the American Heart Association, 2021, 10, e018126.	1.6	6
257	Omega-3 Polyunsaturated Fatty Acid: A Pharmaco-Nutraceutical Approach to Improve the Responsiveness to Ursodeoxycholic Acid. Nutrients, 2021, 13, 2617.	1.7	6
258	Influences of Gestational Obesity on Associations between Genotypes and Gene Expression Levels in Offspring following Maternal Gastrointestinal Bypass Surgery for Obesity. PLoS ONE, 2015, 10, e0117011.	1.1	6
259	Adipose methylome integrative-omic analyses reveal genetic and dietary metabolic health drivers and insulin resistance classifiers. Genome Medicine, 2022, 14, .	3.6	6
260	Detection of a MspI restriction fragment length polymorphism for the human sex hormone-binding globulin (SHBG) gene. Human Genetics, 1994, 93, 84.	1.8	5
261	ASSOCIATION BETWEEN µâ€OPIOID RECEPTORâ€1 102T>C POLYMORPHISM AND INTERMEDIATE TYPE 2 DIABETES PHENOTYPES: RESULTS FROM THE QUEBEC FAMILY STUDY (QFS). Clinical and Experimental Pharmacology and Physiology, 2008, 35, 1018-1022.	0.9	5
262	Interaction between diets, polymorphisms and plasma lipid levels. Clinical Lipidology, 2010, 5, 421-438.	0.4	5
263	Effects of neuromedin- $\hat{l}^2$ on caloric compensation, eating behaviours and habitual food intake. Appetite, 2011, 57, 21-27.	1.8	5
264	Investigation of LRP8 gene in 1p31 QTL linked to LDL peak particle diameter in the Quebec family study. Molecular Genetics and Metabolism, 2011, 102, 448-452.	0.5	5
265	Association between plasma omega-3 fatty acids and cardiovascular disease risk factors. Applied Physiology, Nutrition and Metabolism, 2013, 38, 243-248.	0.9	5
266	Familial resemblances in human whole blood transcriptome. BMC Genomics, 2018, 19, 300.	1.2	5
267	Assessment of the American Heart Association's "Life's simple 7―score in French-speaking adults from Québec. Nutrition, Metabolism and Cardiovascular Diseases, 2019, 29, 684-691.	1.1	5
268	Integrative Network Analysis of Multi-Omics Data in the Link between Plasma Carotenoid Concentrations and Lipid Profile. Lifestyle Genomics, 2020, 13, 11-19.	0.6	5
269	Exploring Attitudes, Subjective Norms and Perceived Behavioural Control in a Genetic-Based and a Population-Based Weight Management Intervention: A One-Year Randomized Controlled Trial. Nutrients, 2020, 12, 3768.	1.7	5
270	Cholecalciferol Supplementation Does Not Prevent the Development of Metabolic Syndrome or Enhance the Beneficial Effects of Omega-3 Fatty Acids in Obese Mice. Journal of Nutrition, 2021, 151, 1175-1189.	1.3	5

#	Article	IF	CITATIONS
271	The apoB-100 gene EcoRI polymorphism influences the relationship between features of the insulin resistance syndrome and the hyper-apoB and dense LDL phenotype in men. Diabetes, 1996, 45, 1405-1411.	0.3	5
272	Myeloperoxidase gene sequence variations are associated with low-density-lipoprotein characteristics. Journal of Human Genetics, 2008, 53, 439-446.	1.1	4
273	Acylation stimulating protein is higher in Inuit from nunavik compared to a southern Quebec population. International Journal of Circumpolar Health, 2009, 68, 421-432.	0.5	4
274	Eating behaviours of non-obese individuals with and without familial history of obesity. British Journal of Nutrition, 2009, 101, 1103-1109.	1.2	4
275	Estimating genetic effect sizes under joint disease-endophenotype models in presence of gene-environment interactions. Frontiers in Genetics, 2015, 6, 248.	1.1	4
276	Impact of systemic enzyme supplementation on low-grade inflammation in humans. PharmaNutrition, 2015, 3, 83-88.	0.8	4
277	Factors Associated with the Intention of Registered Dietitians to Discuss Nutrigenetics with their Patients/Clients. Canadian Journal of Dietetic Practice and Research, 2016, 77, 163-169.	0.5	4
278	Acute Effects of Single Doses of Bonito Fish Peptides and Vitamin D on Whole Blood Gene Expression Levels: A Randomized Controlled Trial. International Journal of Molecular Sciences, 2019, 20, 1944.	1.8	4
279	Associations of Intake of Free and Naturally Occurring Sugars from Solid Foods and Drinks with Cardiometabolic Risk Factors in a Quebec Adult Population: The PREDISE (PRÉDicteurs Individuels,) Tj ETQq1	l <b>1 017:84</b> 31	.4 ngBT /Over
280	A Variant in the LRRFIP1 Gene Is Associated With Adiposity and Inflammation. Obesity, $0, \dots$	1.5	4
280	A Variant in the LRRFIP1 Gene Is Associated With Adiposity and Inflammation. Obesity, 0, , .  A Systematic Review and Recommendations Around Frameworks for Evaluating Scientific Validity in Nutritional Genomics. Frontiers in Nutrition, 2021, 8, 789215.	1.5	4
	A Systematic Review and Recommendations Around Frameworks for Evaluating Scientific Validity in		4 4 3
281	A Systematic Review and Recommendations Around Frameworks for Evaluating Scientific Validity in Nutritional Genomics. Frontiers in Nutrition, 2021, 8, 789215.  Effect of the factor VII R353Q missense mutation on plasma apolipoproteinÂB levels: impact of visceral	1.6	4
281	A Systematic Review and Recommendations Around Frameworks for Evaluating Scientific Validity in Nutritional Genomics. Frontiers in Nutrition, 2021, 8, 789215.  Effect of the factor VII R353Q missense mutation on plasma apolipoproteinÂB levels: impact of visceral obesity. Journal of Human Genetics, 2003, 48, 367-373.  Interaction between Familial History of Obesity and Fat Intakes on Obesity Phenotypes. Journal of	1.6	3
281 282 283	A Systematic Review and Recommendations Around Frameworks for Evaluating Scientific Validity in Nutritional Genomics. Frontiers in Nutrition, 2021, 8, 789215.  Effect of the factor VII R353Q missense mutation on plasma apolipoproteinÂB levels: impact of visceral obesity. Journal of Human Genetics, 2003, 48, 367-373.  Interaction between Familial History of Obesity and Fat Intakes on Obesity Phenotypes. Journal of Nutrigenetics and Nutrigenomics, 2009, 2, 37-42.  A GWAS followâ€up of obesityâ€related SNPs in SYPL2 reveals sexâ€specific association with hip	1.6 1.1 1.8	3 3
281 282 283 284	A Systematic Review and Recommendations Around Frameworks for Evaluating Scientific Validity in Nutritional Genomics. Frontiers in Nutrition, 2021, 8, 789215.  Effect of the factor VII R353Q missense mutation on plasma apolipoproteinÂB levels: impact of visceral obesity. Journal of Human Genetics, 2003, 48, 367-373.  Interaction between Familial History of Obesity and Fat Intakes on Obesity Phenotypes. Journal of Nutrigenetics and Nutrigenomics, 2009, 2, 37-42.  A GWAS followâ€up of obesityâ€related SNPs in SYPL2 reveals sexâ€specific association with hip circumference. Obesity Science and Practice, 2016, 2, 407-414.  The Challenge of Stratifying Obesity: Attempts in the Quebec Family Study. Frontiers in Genetics, 2019,	1.6 1.1 1.8 1.0	3 3
281 282 283 284	A Systematic Review and Recommendations Around Frameworks for Evaluating Scientific Validity in Nutritional Genomics. Frontiers in Nutrition, 2021, 8, 789215.  Effect of the factor VII R353Q missense mutation on plasma apolipoproteinÂB levels: impact of visceral obesity. Journal of Human Genetics, 2003, 48, 367-373.  Interaction between Familial History of Obesity and Fat Intakes on Obesity Phenotypes. Journal of Nutrigenetics and Nutrigenomics, 2009, 2, 37-42.  A GWAS followâ€up of obesityâ€related SNPs in SYPL2 reveals sexâ€specific association with hip circumference. Obesity Science and Practice, 2016, 2, 407-414.  The Challenge of Stratifying Obesity: Attempts in the Quebec Family Study. Frontiers in Genetics, 2019, 10, 994.  Response to the Consensus Report of the Academy of Nutrition and Dietetics: Incorporating Genetic	1.6 1.1 1.8 1.0	3 3 3

#	Article	IF	CITATIONS
289	Changes in systolic blood pressure, postprandial glucose, and gut microbial composition following mango consumption in individuals with overweight and obesity. Applied Physiology, Nutrition and Metabolism, 2022, 47, 565-574.	0.9	3
290	Eco-Evolutionary Dynamics of the Human-Gut Microbiota Symbiosis in a Changing Nutritional Environment. Evolutionary Biology, 2022, 49, 255-264.	0.5	3
291	Effect of a Six-Week National Cholesterol Education Program Step 1 Diet on Plasma Sex Hormone-Binding Globulin Levels In Overweight Premenopausal Women. Metabolic Syndrome and Related Disorders, 2007, 5, 22-33.	0.5	2
292	<i>Dietary Intakes</i> And Familial History of Obesity. Canadian Journal of Dietetic Practice and Research, 2008, 69, 97-100.	0.5	2
293	Effect of the Mediterranean Diet on the Lipid-Lipoprotein Profile: Is It Influenced by the Family History of Dyslipidemia?. Journal of Nutrigenetics and Nutrigenomics, 2015, 7, 177-187.	1.8	2
294	Electronic Health Record-Based Genome-Wide Meta-Analysis Provides New Insights on the Genetic Architecture of Non-Alcoholic Fatty Liver Disease. SSRN Electronic Journal, 0, , .	0.4	2
295	Authors' Response. Journal of the Academy of Nutrition and Dietetics, 2021, 121, 1216-1217.	0.4	2
296	Molecular remodeling of adipose tissue is associated with metabolic recovery after weight loss surgery. Journal of Translational Medicine, 2022, 20, .	1.8	2
297	1.P.168 Abdominal obesity and hyperinsulinemia as predictors of higher cost of coronary artery bypass grafting among men with or without familial hypercholesterolemia. Atherosclerosis, 1997, 134, 52.	0.4	1
298	Presence of palmar xanthomas in myotonic dystrophy identifies different patterns of linkage disequilibrium between the apolipoprotein E and myotonic dystrophy protein kinase loci. Genetics in Medicine, 2005, 7, 213-215.	1.1	1
299	Association Study between Candidate Genes and Obesity-Related Phenotypes Using a Sample of Lumberjacks. Public Health Genomics, 2009, 12, 253-258.	0.6	1
300	Genetic Variation of PPARs. PPAR Research, 2009, 2009, 1-1.	1.1	1
301	Thymic stromal lymphopoietin: an immune cytokine gene associated with the metabolic syndrome and blood pressure in severe obesity. Clinical Science, 2012, 123, 271-271.	1.8	1
302	Liking for foods high in salt and fat is associated with a lower diet quality but liking for foods high in sugar is not $\hat{a} \in \text{``Results from the PREDISE study. Food Quality and Preference, 2021, 88, 104073.}$	2.3	1
303	Electronic Health Record-Based Genome-Wide Meta-Analysis Identifies New Susceptibility Loci for Non-Alcoholic Fatty Liver Disease. Journal of the Endocrine Society, 2021, 5, A501-A501.	0.1	1
304	Fatty acids of plasma phospholipids and erythrocytes are reliable biomarkers of nâ€3 polyunsaturated fatty acid supplementation. FASEB Journal, 2010, 24, 939.6.	0.2	1
305	Clinical Practice Guidelines Using GRADE and AGREE II for the Impact of Genetic Variants on Plasma Lipid/Lipoprotein/Apolipoprotein Responsiveness to Omega-3 Fatty Acids. Frontiers in Nutrition, 2021, 8, 768474.	1.6	1
306	1.P.288 Relative contribution of LDL-receptor and LPL gene mutations to angiographically assessed coronary artery disease among French Canadians. Atherosclerosis, 1997, 134, 77.	0.4	0

#	Article	IF	CITATIONS
307	Erratum to "Interaction between HNF4A polymorphisms and physical activity in relation to type 2 diabetes-related traits: Results from the Quebec Family Study―[Diabetes Res. Clin. Pract. 84 (2009) 211–218]. Diabetes Research and Clinical Practice, 2010, 90, 126.	1.1	O
308	Editorial: Dietary Factors, Epigenetics and Their Implications for Human Obesity. Frontiers in Endocrinology, 2020, 11, 601.	1.5	0
309	Genome-Wide Meta-Analysis and Mendelian Randomization Identify Early Biomarkers of Non-Alcoholic Fatty Liver Disease. Journal of the Endocrine Society, 2021, 5, A315-A315.	0.1	0
310	Visceral adipose tissue DNA methylation at dipeptidyl peptidaseâ€4 gene locus is associated with gene expression and plasma lipid levels in severe obesity. FASEB Journal, 2010, 24, .	0.2	0
311	Regulation of the PBMCs gene expression profile with the Western dietary pattern in healthy men and women. FASEB Journal, 2012, 26, 647.5.	0.2	0
312	Association between polymorphisms in the FADS gene cluster and the plasma triacylglycerol response to an ωâ€3 PUFA supplementation. FASEB Journal, 2012, 26, 647.14.	0.2	0
313	Interaction effects between nâ€3 polyunsaturated fatty acids and genetic variations in genes involved in de novo lipogenesis on plasma triglyceride levels. FASEB Journal, 2013, 27, 222.1.	0.2	0
314	DUSP1 gene polymorphisms are associated with obesityrelated metabolic complications and gene methylation levels in severely obese patients. FASEB Journal, 2013, 27, 226.1.	0.2	0
315	Cardiometabolic risk factors are influenced by Stearoylâ€CoA Desaturaseâ€1 (SCD1) polymorphisms and nâ€3 polyunsaturated fatty acids supplementation. FASEB Journal, 2013, 27, 640.10.	0.2	O
316	Polymorphisms in the MGLL gene are associated with plasma LDL  response to a marine nâ€3 PUFA supplementation (1038.1). FASEB Journal, 2014, 28, 1038.1.	0.2	0
317	Bariatric Surgery Induces Hypomethylation of Genes Related to Type 2 Diabetes and Insulin Resistance. International Journal of Molecular Biology Open Access, 2017, 2, .	0.2	0
318	Predicting Adherence to Canada's Food Guide Recommendations on Healthy Food Choices Using Machine Learning Algorithms. Current Developments in Nutrition, 2022, 6, 99.	0.1	0