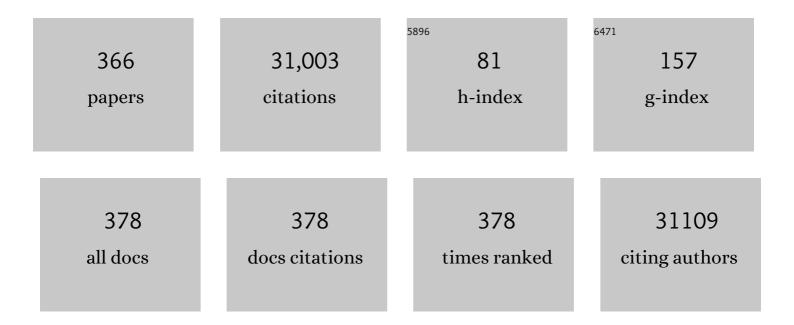
Louis Perusse

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Dietary Mediators of the Genetic Susceptibility to Obesity—Results from the Quebec Family Study. Journal of Nutrition, 2022, 152, 49-58.	2.9	8
2	Effects of sodium intake and cardiorespiratory fitness on body composition and genetic susceptibility to obesity: results from the Quebec Family Study. British Journal of Nutrition, 2022, , 1-10.	2.3	0
3	Understanding Gene-Lifestyle Interaction in Obesity: The Role of Mediation versus Moderation. Lifestyle Genomics, 2022, 15, 67-76.	1.7	5
4	The fit-active profile to better reflect the benefits of a lifelong vigorous physical activity participation: mini-review of literature and population data. Applied Physiology, Nutrition and Metabolism, 2021, 46, 1-8.	1.9	1
5	The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679.	27.8	353
6	Genetic Variation in the Response to Exercise Training. , 2020, , 187-196.		1
7	Integrative Network Analysis of Multi-Omics Data in the Link between Plasma Carotenoid Concentrations and Lipid Profile. Lifestyle Genomics, 2020, 13, 11-19.	1.7	5
8	Genetics of Energy Expenditure in Humans. , 2020, , 135-145.		1
9	Genetics of Obesity: Family Studies. , 2020, , 79-92.		1
10	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.	7.9	44
11	Associations of autozygosity with a broad range of human phenotypes. Nature Communications, 2019, 10, 4957.	12.8	84
12	The Challenge of Stratifying Obesity: Attempts in the Quebec Family Study. Frontiers in Genetics, 2019, 10, 994.	2.3	3
13	Circulating glutamate level as a potential biomarker for abdominal obesity and metabolic risk. Nutrition, Metabolism and Cardiovascular Diseases, 2019, 29, 1353-1360.	2.6	29
14	Associations Between Dietary Protein Sources, Plasma BCAA and Short-Chain Acylcarnitine Levels in Adults. Nutrients, 2019, 11, 173.	4.1	47
15	Network Analysis of the Potential Role of DNA Methylation in the Relationship between Plasma Carotenoids and Lipid Profile. Nutrients, 2019, 11, 1265.	4.1	17
16	Weighted gene co-expression network analysis to explain the relationship between plasma total carotenoids and lipid profile. Genes and Nutrition, 2019, 14, 16.	2.5	9
17	Protein intake and the incidence of pre-diabetes and diabetes in 4 population-based studies: the PREVIEW project. American Journal of Clinical Nutrition, 2019, 109, 1310-1318.	4.7	28
18	Familial resemblances in human plasma metabolites are attributable to both genetic and common environmental effects. Nutrition Research, 2019, 61, 22-30.	2.9	18

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19	The relationship between yogurt consumption, body weight, and metabolic profiles in youth with a familial predisposition to obesity. European Journal of Clinical Nutrition, 2019, 73, 541-548.	2.9	11
20	Familial resemblances in human whole blood transcriptome. BMC Genomics, 2018, 19, 300.	2.8	5
21	Acute cardiorespiratory responses in participants with heart disease during cycling at different immersion levels. Clinical Physiology and Functional Imaging, 2018, 38, 100-107.	1.2	3
22	Acute effects of water immersion on heart rate variability in participants with heart disease. Clinical Physiology and Functional Imaging, 2018, 38, 233-239.	1.2	2
23	Yogurt consumption, body composition, and metabolic health in the Québec Family Study. European Journal of Nutrition, 2018, 57, 1591-1603.	3.9	21
24	Genetic and Common Environmental Contributions to Familial Resemblances in Plasma Carotenoid Concentrations in Healthy Families. Nutrients, 2018, 10, 1002.	4.1	7
25	The role of eating behavior traits in mediating genetic susceptibility to obesity. American Journal of Clinical Nutrition, 2018, 108, 445-452.	4.7	39
26	Polygenic risk score for predicting weight loss after bariatric surgery. JCI Insight, 2018, 3, .	5.0	30
27	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.	5.0	20
28	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. Nature Communications, 2017, 8, 14977.	12.8	169
29	Obesity, genes, and sleep habits. American Journal of Clinical Nutrition, 2017, 105, 779-780.	4.7	1
30	Guide for Current Nutrigenetic, Nutrigenomic, and Nutriepigenetic Approaches for Precision Nutrition Involving the Prevention and Management of Chronic Diseases Associated with Obesity. Journal of Nutrigenetics and Nutrigenomics, 2017, 10, 43-62.	1.3	118
31	Acute breathing patterns in healthy and heart disease participants during cycling at different levels of immersion. Respiratory Physiology and Neurobiology, 2017, 235, 1-7.	1.6	4
32	Genome-wide physical activity interactions in adiposity ― A meta-analysis of 200,452 adults. PLoS Genetics, 2017, 13, e1006528.	3.5	158
33	Association between Metabolite Profiles, Metabolic Syndrome and Obesity Status. Nutrients, 2016, 8, 324.	4.1	33
34	No Evidence of a Common DNA Variant Profile Specific to World Class Endurance Athletes. PLoS ONE, 2016, 11, e0147330.	2.5	96
35	The economic consequences of obesity and overweight among adults in Quebec. Canadian Journal of Public Health, 2016, 107, e507-e513.	2.3	7
36	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.	1.9	3

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37	Advances in Exercise, Fitness, and Performance Genomics in 2015. Medicine and Science in Sports and Exercise, 2016, 48, 1906-1916.	0.4	52
38	A CpG-SNP Located within the <i>ARPC3</i> Gene Promoter Is Associated with Hypertriglyceridemia in Severely Obese Patients. Annals of Nutrition and Metabolism, 2016, 68, 203-212.	1.9	12
39	A principal component meta-analysis on multiple anthropometric traits identifies novel loci for body shape. Nature Communications, 2016, 7, 13357.	12.8	74
40	Familial resemblances in blood leukocyte DNA methylation levels. Epigenetics, 2016, 11, 831-838.	2.7	10
41	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.	2.7	15
42	Genome-wide association studies suggest sex-specific loci associated with abdominal and visceral fat. International Journal of Obesity, 2016, 40, 662-674.	3.4	74
43	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. Nature Communications, 2016, 7, 10495.	12.8	245
44	Genome-wide meta-analysis uncovers novel loci influencing circulating leptin levels. Nature Communications, 2016, 7, 10494.	12.8	153
45	Association between yogurt consumption, dietary patterns, and cardio-metabolic risk factors. European Journal of Nutrition, 2016, 55, 577-587.	3.9	51
46	The rare allele of DGKZ SNP rs10838599 is associated with variability in HDL-cholesterol levels among severely obese patients. Integrative Obesity and Diabetes, 2016, 2, 219-224.	0.2	1
47	Impact of NMT1 Gene Polymorphisms on Features of the Metabolic Syndrome among Severely Obese Patients. Obesity Research - Open Journal, 2016, 2, 101-110.	0.4	Ο
48	Estimating genetic effect sizes under joint disease-endophenotype models in presence of gene-environment interactions. Frontiers in Genetics, 2015, 6, 248.	2.3	4
49	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.	3.5	331
50	Natural Rumenâ€Derived <i>trans</i> Fatty Acids Are Associated with Metabolic Markers of Cardiac Health. Lipids, 2015, 50, 873-882.	1.7	36
51	New genetic loci link adipose and insulin biology to body fat distribution. Nature, 2015, 518, 187-196.	27.8	1,328
52	Genetic studies of body mass index yield new insights for obesity biology. Nature, 2015, 518, 197-206.	27.8	3,823
53	C3 Polymorphism Influences Circulating Levels of C3, ASP and Lipids in Schizophrenic Patients. Neurochemical Research, 2015, 40, 906-914.	3.3	10
54	Directional dominance on stature and cognition inÂdiverse human populations. Nature, 2015, 523, 459-462.	27.8	173

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55	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.	1.3	24
56	Advances in Exercise, Fitness, and Performance Genomics in 2014. Medicine and Science in Sports and Exercise, 2015, 47, 1105-1112.	0.4	38
57	Novel loci associated with usual sleep duration: the CHARGE Consortium Genome-Wide Association Study. Molecular Psychiatry, 2015, 20, 1232-1239.	7.9	112
58	Replication of 6 Obesity Genes in a Meta-Analysis of Genome-Wide Association Studies from Diverse Ancestries. PLoS ONE, 2014, 9, e96149.	2.5	56
59	Differential methylation in visceral adipose tissue of obese men discordant for metabolic disturbances. Physiological Genomics, 2014, 46, 216-222.	2.3	43
60	FTO genetic variants, dietary intake and body mass index: insights from 177 330 individuals. Human Molecular Genetics, 2014, 23, 6961-6972.	2.9	143
61	Crossâ€sectional associations of acylation stimulating protein (<scp>ASP</scp>) and adipose tissue gene expression with estradiol and progesterone in preâ€and postmenopausal women. Clinical Endocrinology, 2014, 81, 736-745.	2.4	11
62	SREBF1 gene variations modulate insulin sensitivity in response to a fish oil supplementation. Lipids in Health and Disease, 2014, 13, 152.	3.0	11
63	Advances in Exercise, Fitness, and Performance Genomics in 2013. Medicine and Science in Sports and Exercise, 2014, 46, 851-859.	0.4	25
64	Findings from the Quebec Family Study on the Etiology of Obesity: Genetics and Environmental Highlights. Current Obesity Reports, 2014, 3, 54-66.	8.4	71
65	Defining the role of common variation in the genomic and biological architecture of adult human height. Nature Genetics, 2014, 46, 1173-1186.	21.4	1,818
66	Yogurt intake is associated with a healthier dietary pattern and is a lower contributor of energy intake in obese individuals (1018.6). FASEB Journal, 2014, 28, 1018.6.	0.5	0
67	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.	2.7	14
68	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.	0.8	6
69	Association between plasma omega-3 fatty acids and cardiovascular disease risk factors. Applied Physiology, Nutrition and Metabolism, 2013, 38, 243-248.	1.9	5
70	<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.	1.6	10
71	A variant in the <i>LRRFIP1</i> gene is associated with adiposity and inflammation. Obesity, 2013, 21, 185-192.	3.0	29
72	Parental eating behavior traits are related to offspring BMI in the Québec Family Study. International Journal of Obesity, 2013, 37, 1422-1426.	3.4	11

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73	Advances in Exercise, Fitness, and Performance Genomics in 2012. Medicine and Science in Sports and Exercise, 2013, 45, 824-831.	0.4	50
74	Advances in Exercise, Fitness, and Performance Genomics in 2011. Medicine and Science in Sports and Exercise, 2012, 44, 809-817.	0.4	55
75	Past dieting is related to rigid control and disinhibition in adolescents from the Québec Family Study. British Journal of Nutrition, 2012, 108, 1976-1979.	2.3	6
76	Thymic stromal lymphopoietin: an immune cytokine gene associated with the metabolic syndrome and blood pressure in severe obesity. Clinical Science, 2012, 123, 99-109.	4.3	16
77	Associations between Polymorphisms in Genes Involved in Fatty Acid Metabolism and Dietary Fat Intakes. Journal of Nutrigenetics and Nutrigenomics, 2012, 5, 1-12.	1.3	8
78	Omega-3 fatty acids status in human subjects estimated using a food frequency questionnaire and plasma phospholipids levels. Nutrition Journal, 2012, 11, 46.	3.4	32
79	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.	4.1	62
80	Individualized Weight Management: What Can Be Learned from Nutrigenomics and Nutrigenetics?. Progress in Molecular Biology and Translational Science, 2012, 108, 347-382.	1.7	25
81	Association of <i>LIPA</i> Gene Polymorphisms With Obesityâ€Related Metabolic Complications Among Severely Obese Patients. Obesity, 2012, 20, 2075-2082.	3.0	23
82	Prevalence and Familial Patterns of Night Eating in the Québec Adipose and Lifestyle InvesTigation in Youth (QUALITY) Study. Obesity, 2012, 20, 1598-1603.	3.0	14
83	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.	21.4	762
84	Association between olfactory receptor genes, eating behavior traits and adiposity: Results from the Quebec Family Study. Physiology and Behavior, 2012, 105, 772-776.	2.1	41
85	A polymorphism of the interferon-gamma-inducible protein 30 gene is associated with hyperglycemia in severely obese individuals. Human Genetics, 2012, 131, 57-66.	3.8	13
86	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.	4.6	51
87	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.	1.3	24
88	Effects of neuromedin-Î ² on caloric compensation, eating behaviours and habitual food intake. Appetite, 2011, 57, 21-27.	3.7	5
89	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.	1.1	5
90	Interactions between Dietary Fat Intake and FASN Genetic Variation Influence LDL Peak Particle Diameter. Journal of Nutrigenetics and Nutrigenomics, 2011, 4, 137-145.	1.3	10

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91	Impact of Nutritional Epigenomics on Disease Risk and Prevention: Introduction. Journal of Nutrigenetics and Nutrigenomics, 2011, 4, 245-247.	1.3	16
92	Single nucleotide polymorphisms in the myostatin (<i>MSTN</i>) and muscle creatine kinase (<i>CKM</i>) genes are not associated with elite endurance performance. Scandinavian Journal of Medicine and Science in Sports, 2011, 21, 841-845.	2.9	11
93	<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.	3.0	52
94	Contributions of Cardiorespiratory Fitness and Visceral Adiposity to Six-Year Changes in Cardiometabolic Risk Markers in Apparently Healthy Men and Women. Journal of Clinical Endocrinology and Metabolism, 2011, 96, 1462-1468.	3.6	38
95	Insulin Resistance, Low Cardiorespiratory Fitness, and Increased Exercise Blood Pressure. Hypertension, 2011, 58, 1036-1042.	2.7	30
96	Set points, settling points and some alternative models: theoretical options to understand how genes and environments combine to regulate body adiposity. DMM Disease Models and Mechanisms, 2011, 4, 733-745.	2.4	266
97	Advances in Exercise, Fitness, and Performance Genomics in 2010. Medicine and Science in Sports and Exercise, 2011, 43, 743-752.	0.4	64
98	Physical Activity Attenuates the Influence of FTO Variants on Obesity Risk: A Meta-Analysis of 218,166 Adults and 19,268 Children. PLoS Medicine, 2011, 8, e1001116.	8.4	446
99	Associations between Dietary Patterns and LDL Peak Particle Diameter: A Cross-Sectional Study. Journal of the American College of Nutrition, 2010, 29, 630-637.	1.8	7
100	The Three-Factor Eating Questionnaire and BMI in adolescents: results from the Québec Family Study. British Journal of Nutrition, 2010, 104, 1074-1079.	2.3	60
101	Advances in Exercise, Fitness, and Performance Genomics. Medicine and Science in Sports and Exercise, 2010, 42, 835-846.	0.4	111
102	Improvements in glucose homeostasis in response to regular exercise are influenced by the PPARG Pro12Ala variant: results from the HERITAGE Family Study. Diabetologia, 2010, 53, 679-689.	6.3	61
103	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.	2.1	21
104	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
105	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.	4.7	193
106	Combining genetic markers and clinical risk factors improves the risk assessment of impaired glucose metabolism. Annals of Medicine, 2010, 42, 196-206.	3.8	11
107	Effect of Implementation Intentions to Change Behaviour: Moderation by Intention Stability. Psychological Reports, 2010, 106, 147-159.	1.7	29
108	A common haplotype and the Pro582Ser polymorphism of the hypoxia-inducible factor-1α (<i>HIF1A</i>) gene in elite endurance athletes. Journal of Applied Physiology, 2010, 108, 1497-1500.	2.5	53

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109	Prediction of daily fruit and vegetable consumption among overweight and obese individuals. Appetite, 2010, 54, 480-484.	3.7	28
110	ACTN3R577X and other polymorphisms are not associated with elite endurance athlete status in the Genathlete study. Journal of Sports Sciences, 2010, 28, 1355-1359.	2.0	41
111	Correlation between $n\hat{a}\in 3$ fatty acid intakes estimated using a food frequency questionnaire and concentrations measured in plasma phospholipids. FASEB Journal, 2010, 24, 939.2.	0.5	0
112	Positional identification of variants of Adamts16 linked to inherited hypertension. Human Molecular Genetics, 2009, 18, 2825-2838.	2.9	57
113	Sex differences in inflammatory markers: what is the contribution of visceral adiposity?. American Journal of Clinical Nutrition, 2009, 89, 1307-1314.	4.7	172
114	Association Study between Candidate Genes and Obesity-Related Phenotypes Using a Sample of Lumberjacks. Public Health Genomics, 2009, 12, 253-258.	1.0	1
115	Low Cardiorespiratory Fitness Levels and Elevated Blood Pressure. Hypertension, 2009, 54, 91-97.	2.7	51
116	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.	1.2	16
117	Meta-Analysis of the INSIG2 Association with Obesity Including 74,345 Individuals: Does Heterogeneity of Estimates Relate to Study Design?. PLoS Genetics, 2009, 5, e1000694.	3.5	62
118	The Human Gene Map for Performance and Health-Related Fitness Phenotypes. Medicine and Science in Sports and Exercise, 2009, 41, 34-72.	0.4	409
119	Body Composition, Cardiorespiratory Fitness, and Low-Grade Inflammation in Middle-Aged Men and Women. American Journal of Cardiology, 2009, 104, 240-246.	1.6	50
120	Analysis of inherited genetic variations at the <i>UGT1</i> locus in the French-Canadian population. Human Mutation, 2009, 30, 677-687.	2.5	28
121	Association between insulin secretion, insulin sensitivity and type 2 diabetes susceptibility variants identified in genome-wide association studies. Acta Diabetologica, 2009, 46, 217-226.	2.5	91
122	Associations between dietary patterns and obesity phenotypes. International Journal of Obesity, 2009, 33, 1419-1426.	3.4	108
123	Prediction of Leisureâ€ŧime Physical Activity Among Obese Individuals. Obesity, 2009, 17, 706-712.	3.0	23
124	Risk Factors for Adult Overweight and Obesity in the Quebec Family Study: Have We Been Barking Up the Wrong Tree?. Obesity, 2009, 17, 1964-1970.	3.0	125
125	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.	2.8	10
126	Phosphoinositide cycle gene polymorphisms affect the plasma lipid profile in the Quebec Family Study. Molecular Genetics and Metabolism, 2009, 97, 149-154.	1.1	24

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127	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.	2.1	24
128	Age-related differences in inflammatory markers in men: contribution of visceral adiposity. Metabolism: Clinical and Experimental, 2009, 58, 1452-1458.	3.4	72
129	Association of <i>OSBPL11</i> Gene Polymorphisms With Cardiovascular Disease Risk Factors in Obesity. Obesity, 2009, 17, 1466-1472.	3.0	31
130	Interaction between Familial History of Obesity and Fat Intakes on Obesity Phenotypes. Journal of Nutrigenetics and Nutrigenomics, 2009, 2, 37-42.	1.3	3
131	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.	1.3	27
132	Eating behaviours of non-obese individuals with and without familial history of obesity. British Journal of Nutrition, 2009, 101, 1103-1109.	2.3	4
133	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.	2.9	9
134	Endothelial nitric oxide synthase gene polymorphism and elite endurance athlete status: the Genathlete study. Scandinavian Journal of Medicine and Science in Sports, 2008, 18, 485-490.	2.9	24
135	Myeloperoxidase gene sequence variations are associated with low-density-lipoprotein characteristics. Journal of Human Genetics, 2008, 53, 439-446.	2.3	4
136	Genome-wide linkage analysis for circulating levels of adipokines and C-reactive protein in the Quebec family study (QFS). Journal of Human Genetics, 2008, 53, 629-636.	2.3	11
137	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.	1.9	5
138	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.	2.3	23
139	Validity of a self-reported measure of familial history of obesity. Nutrition Journal, 2008, 7, 27.	3.4	26
140	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.	4.6	54
141	Genetic and Nutritional Determinants of the Metabolic Syndrome: Introduction. Journal of Nutrigenetics and Nutrigenomics, 2008, 1, 97-99.	1.3	1
142	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.	0.6	206
143	Evidence of a quantitative trait locus for energy and macronutrient intakes on chromosome 3q27.3: the Québec Family Study. American Journal of Clinical Nutrition, 2008, 88, 1142-1148.	4.7	20
144	<i>Dietary Intakes</i> And Familial History of Obesity. Canadian Journal of Dietetic Practice and Research, 2008, 69, 97-100.	0.6	2

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145	A Simple Method to Assess Fruit and Vegetable Intake among Obese and Non-obese Individuals. Canadian Journal of Public Health, 2008, 99, 494-498.	2.3	22
146	Abstract 5083: Body Composition, Cardiorespiratory Fitness and Low-Grade Inflammation in Middle-Aged Men and Women. Circulation, 2008, 118, .	1.6	0
147	Visceral Adipose Tissue Accumulation, Cardiorespiratory Fitness, and Features of the Metabolic Syndrome. Archives of Internal Medicine, 2007, 167, 1518.	3.8	118
148	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.	1.0	9
149	A pharmacogenomics study of the human estrogen glucuronosyltransferase UGT1A3. Pharmacogenetics and Genomics, 2007, 17, 481-495.	1.5	33
150	The lipoprotein/lipid profile is modulated by a gene–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.	2.3	28
151	Influences of the phosphatidylcholine transfer protein gene variants on the LDL peak particle size. Atherosclerosis, 2007, 195, 297-302.	0.8	12
152	Association between a β2-adrenergic receptor polymorphism and elite endurance performance. Metabolism: Clinical and Experimental, 2007, 56, 1649-1651.	3.4	59
153	Adiponectin and adiponectin receptor gene variants in relation to resting metabolic rate, respiratory quotient, and adiposity-related phenotypes in the Québec Family Study. American Journal of Clinical Nutrition, 2007, 85, 26-34.	4.7	53
154	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.	3.4	50
155	Contribution of several candidate gene polymorphisms in the determination of adiposity changes: results from the Québec Family Study. International Journal of Obesity, 2007, 31, 891-899.	3.4	20
156	Evidence of Linkage and Association with Body Fatness and Abdominal Fat on Chromosome 15q26. Obesity, 2007, 15, 2061-2070.	3.0	13
157	Genes, Fat Intake, and Cardiovascular Disease Risk Factors in the Quebec Family Study. Obesity, 2007, 15, 2336-2347.	3.0	21
158	Association of <i>Lipin 1</i> Gene Polymorphisms with Measures of Energy and Glucose Metabolism. Obesity, 2007, 15, 2723-2732.	3.0	44
159	Quantitative Trait Locus on 15q for a Metabolic Syndrome Variable Derived from Factor Analysis*. Obesity, 2007, 15, 544-550.	3.0	29
160	Associations between USF1 gene variants and cardiovascular risk factors in the Quebec Family Study. Clinical Genetics, 2007, 71, 245-253.	2.0	15
161	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.	3.9	33
162	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.	4.6	20

Louis Perusse

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163	Resting metabolic rate and respiratory quotient: results from a genome-wide scan in the Quebec Family Study. American Journal of Clinical Nutrition, 2006, 84, 1527-1533.	4.7	25
164	The Human Gene Map for Performance and Health-Related Fitness Phenotypes. Medicine and Science in Sports and Exercise, 2006, 38, 1863-1888.	0.4	323
165	An overview of obesityâ€specific quality of life questionnaires. Obesity Reviews, 2006, 7, 347-360.	6.5	66
166	A Quantitative Trait Locus for Body Fat on Chromosome 1q43 in French Canadians: Linkage and Association Studies. Obesity, 2006, 14, 1605-1615.	3.0	24
167	The Human Obesity Gene Map: The 2005 Update. Obesity, 2006, 14, 529-644.	3.0	962
168	Polymorphisms in the leptin and leptin receptor genes in relation to resting metabolic rate and respiratory quotient in the Québec Family Study. International Journal of Obesity, 2006, 30, 183-190.	3.4	36
169	Health-Related Quality of Life in Morbid Obesity. Obesity Surgery, 2006, 16, 574-579.	2.1	57
170	Influence of Nonsynonymous Polymorphisms of UGT1A8 and UGT2B7 Metabolizing Enzymes on the Formation of Phenolic and Acyl Glucuronides of Mycophenolic Acid. Drug Metabolism and Disposition, 2006, 34, 1539-1545.	3.3	91
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