Marie-France Hivert

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	New genetic loci link adipose and insulin biology to body fat distribution. Nature, 2015, 518, 187-196.	27.8	1,328
2	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
3	DNA Methylation in Newborns and Maternal Smoking in Pregnancy: Genome-wide Consortium Meta-analysis. American Journal of Human Genetics, 2016, 98, 680-696.	6.2	717
4	Genetic variation in GIPR influences the glucose and insulin responses to an oral glucose challenge. Nature Genetics, 2010, 42, 142-148.	21.4	591
5	Novel Loci for Adiponectin Levels and Their Influence on Type 2 Diabetes and Metabolic Traits: A Multi-Ethnic Meta-Analysis of 45,891 Individuals. PLoS Genetics, 2012, 8, e1002607.	3.5	419
6	Maternal and fetal genetic effects on birth weight and their relevance to cardio-metabolic risk factors. Nature Genetics, 2019, 51, 804-814.	21.4	402
7	Refining the accuracy of validated target identification through coding variant fine-mapping in type 2 diabetes. Nature Genetics, 2018, 50, 559-571.	21.4	356
8	Impact of common genetic determinants of Hemoglobin A1c on type 2 diabetes risk and diagnosis in ancestrally diverse populations: A transethnic genome-wide meta-analysis. PLoS Medicine, 2017, 14, e1002383.	8.4	341
9	Impact of Type 2 Diabetes Susceptibility Variants on Quantitative Glycemic Traits Reveals Mechanistic Heterogeneity. Diabetes, 2014, 63, 2158-2171.	0.6	297
10	Detailed Physiologic Characterization Reveals Diverse Mechanisms for Novel Genetic Loci Regulating Glucose and Insulin Metabolism in Humans. Diabetes, 2010, 59, 1266-1275.	0.6	237
11	Associations of Adiponectin, Resistin, and Tumor Necrosis Factor-α with Insulin Resistance. Journal of Clinical Endocrinology and Metabolism, 2008, 93, 3165-3172.	3.6	229
12	Genetic Evidence for Causal Relationships Between Maternal Obesity-Related Traits and Birth Weight. JAMA - Journal of the American Medical Association, 2016, 315, 1129.	7.4	220
13	Gestational diabetes mellitus epigenetically affects genes predominantly involved in metabolic diseases. Epigenetics, 2013, 8, 935-943.	2.7	217
14	Maternal BMI at the start of pregnancy and offspring epigenome-wide DNA methylation: findings from the pregnancy and childhood epigenetics (PACE) consortium. Human Molecular Genetics, 2017, 26, 4067-4085.	2.9	211
15	Precision Medicine in Diabetes: A Consensus Report From the American Diabetes Association (ADA) and the European Association for the Study of Diabetes (EASD). Diabetes Care, 2020, 43, 1617-1635.	8.6	204
16	Genetic Evidence for a Normal-Weight "Metabolically Obese―Phenotype Linking Insulin Resistance, Hypertension, Coronary Artery Disease, and Type 2 Diabetes. Diabetes, 2014, 63, 4369-4377.	0.6	185
17	Low-frequency and rare exome chip variants associate with fasting glucose and type 2 diabetes susceptibility. Nature Communications, 2015, 6, 5897.	12.8	173
18	Updated Genetic Score Based on 34 Confirmed Type 2 Diabetes Loci Is Associated With Diabetes Incidence and Regression to Normoglycemia in the Diabetes Prevention Program. Diabetes, 2011, 60, 1340-1348.	0.6	172

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19	Genome-wide association study of offspring birth weight in 86 577 women identifies five novel loci and highlights maternal genetic effects that are independent of fetal genetics. Human Molecular Genetics, 2018, 27, 742-756.	2.9	156
20	Common Variants in the Adiponectin Gene (<i>ADIPOQ</i>) Associated With Plasma Adiponectin Levels, Type 2 Diabetes, and Diabetes-Related Quantitative Traits. Diabetes, 2008, 57, 3353-3359.	0.6	147
21	Clear detection of ADIPOQ locus as the major gene for plasma adiponectin: Results of genome-wide association analyses including 4659 European individuals. Atherosclerosis, 2010, 208, 412-420.	0.8	146
22	Maternal Gestational Diabetes Mellitus and Newborn DNA Methylation: Findings From the Pregnancy and Childhood Epigenetics Consortium. Diabetes Care, 2020, 43, 98-105.	8.6	145
23	Heterogeneous Contribution of Insulin Sensitivity and Secretion Defects to Gestational Diabetes Mellitus. Diabetes Care, 2016, 39, 1052-1055.	8.6	142
24	Meta-analysis of epigenome-wide association studies in neonates reveals widespread differential DNA methylation associated with birthweight. Nature Communications, 2019, 10, 1893.	12.8	140
25	The nasal methylome as a biomarker of asthma and airway inflammation in children. Nature Communications, 2019, 10, 3095.	12.8	129
26	Polygenic Type 2 Diabetes Prediction at the Limit of Common Variant Detection. Diabetes, 2014, 63, 2172-2182.	0.6	127
27	Identification of <i>HKDC1</i> and <i>BACE2</i> as Genes Influencing Clycemic Traits During Pregnancy Through Genome-Wide Association Studies. Diabetes, 2013, 62, 3282-3291.	0.6	119
28	Association of Variants in <i>RETN</i> With Plasma Resistin Levels and Diabetes-Related Traits in the Framingham Offspring Study. Diabetes, 2009, 58, 750-756.	0.6	107
29	Cohort Profile: Pregnancy And Childhood Epigenetics (PACE) Consortium. International Journal of Epidemiology, 2018, 47, 22-23u.	1.9	105
30	Lower Adiponectin Levels at First Trimester of Pregnancy Are Associated With Increased Insulin Resistance and Higher Risk of Developing Gestational Diabetes Mellitus. Diabetes Care, 2013, 36, 1577-1583.	8.6	102
31	Precision medicine in diabetes: a Consensus Report from the American Diabetes Association (ADA) and the European Association for the Study of Diabetes (EASD). Diabetologia, 2020, 63, 1671-1693.	6.3	102
32	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
33	Persistent DNA methylation changes associated with prenatal mercury exposure and cognitive performance during childhood. Scientific Reports, 2017, 7, 288.	3.3	95
34	Plasma Concentrations of Per- and Polyfluoroalkyl Substances at Baseline and Associations with Glycemic Indicators and Diabetes Incidence among High-Risk Adults in the Diabetes Prevention Program Trial. Environmental Health Perspectives, 2017, 125, 107001.	6.0	88
35	Adaptations of placental and cord blood <i>ABCA1Â</i> DNA methylation profile to maternal metabolic status. Epigenetics, 2013, 8, 1289-1302.	2.7	86
36	Per- and polyfluoroalkyl substances and blood lipid levels in pre-diabetic adults—longitudinal analysis of the diabetes prevention program outcomes study. Environment International, 2019, 129, 343-353.	10.0	80

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37	Realthy Lifestyle Interventions to Combat Noncommunicable Diseaseat A Novel Nonhierarchical Connectivity Model for Key Stakeholders: A Policy Statement From the American Heart Association, European Society of Cardiology, European Association for Cardiovascular Prevention and Rehabilitation, and American College of Preventive Medicine. Mayo Clinic Proceedings, 2015, 90,	3.0	77
38	Epigenome-Wide Association Study of Incident Type 2 Diabetes in a British Population: EPIC-Norfolk Study. Diabetes, 2019, 68, 2315-2326.	0.6	77
39	Hypertensive Disorders of Pregnancy and DNA Methylation in Newborns. Hypertension, 2019, 74, 375-383.	2.7	73
40	Medical Training to Achieve Competency in Lifestyle Counseling: An Essential Foundation for Prevention and Treatment of Cardiovascular Diseases and Other Chronic Medical Conditions: A Scientific Statement From the American Heart Association. Circulation, 2016, 134, e308-e327.	1.6	71
41	Genetics of Glucose regulation in Gestation and Growth (Gen3G): a prospective prebirth cohort of mother–child pairs in Sherbrooke, Canada. BMJ Open, 2016, 6, e010031.	1.9	67
42	PPARGC1α gene DNA methylation variations in human placenta mediate the link between maternal hyperglycemia and leptin levels in newborns. Clinical Epigenetics, 2016, 8, 72.	4.1	66
43	Diabetes Risk Perception and Intention to Adopt Healthy Lifest yles Among Primary Care Patients. Diabetes Care, 2009, 32, 1820-1822.	8.6	64
44	Identifying primary care patients at risk for future diabetes and cardiovascular disease using electronic health records. BMC Health Services Research, 2009, 9, 170.	2.2	63
45	Greater early and midâ€pregnancy gestational weight gains are associated with excess adiposity in midâ€childhood. Obesity, 2016, 24, 1546-1553.	3.0	62
46	An integrative cross-omics analysis of DNA methylation sites of glucose and insulin homeostasis. Nature Communications, 2019, 10, 2581.	12.8	62
47	Birth weight-for-gestational age is associated with DNA methylation at birth and in childhood. Clinical Epigenetics, 2016, 8, 118.	4.1	61
48	Maternal alcohol consumption and offspring DNA methylation: findings from six general population-based birth cohorts. Epigenomics, 2018, 10, 27-42.	2.1	58
49	Prenatal Exposure to Mercury: Associations with Global DNA Methylation and Hydroxymethylation in Cord Blood and in Childhood. Environmental Health Perspectives, 2017, 125, 087022.	6.0	57
50	Trans-ethnic Meta-analysis and Functional Annotation Illuminates theÂGenetic Architecture of Fasting Glucose and Insulin. American Journal of Human Genetics, 2016, 99, 56-75.	6.2	55
51	Susceptibility to type 2 diabetes mellitus—from genes to prevention. Nature Reviews Endocrinology, 2014, 10, 198-205.	9.6	54
52	Association of Perfluoroalkyl and Polyfluoroalkyl Substances With Adiposity. JAMA Network Open, 2018, 1, e181493.	5.9	54
53	ADA/EASD Precision Medicine in Diabetes Initiative: An International Perspective and Future Vision for Precision Medicine in Diabetes. Diabetes Care, 2022, 45, 261-266.	8.6	53
54	Associations of Perfluoroalkyl and Polyfluoroalkyl Substances With Incident Diabetes and Microvascular Disease. Diabetes Care, 2019, 42, 1824-1832.	8.6	49

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55	Mediation by Placental DNA Methylation of the Association of Prenatal Maternal Smoking and Birth Weight. American Journal of Epidemiology, 2019, 188, 1878-1886.	3.4	48
56	Developmental programming: Stateâ€ofâ€ŧheâ€science and future directions–Summary from a Pennington Biomedical symposium. Obesity, 2016, 24, 1018-1026.	3.0	47
57	Multi-ancestry genome-wide association study of gestational diabetes mellitus highlights genetic links with type 2 diabetes. Human Molecular Genetics, 2022, 31, 3377-3391.	2.9	47
58	Patterns of body mass index milestones in early life and cardiometabolic risk in early adolescence. International Journal of Epidemiology, 2019, 48, 157-167.	1.9	45
59	Epigenetic age acceleration is associated with allergy and asthma in children in Project Viva. Journal of Allergy and Clinical Immunology, 2019, 143, 2263-2270.e14.	2.9	43
60	Neighborhood Child Opportunity Index and Adolescent Cardiometabolic Risk. Pediatrics, 2021, 147, .	2.1	43
61	Sustainable food systems and nutrition in the 21st century: a report from the 22nd annual Harvard Nutrition Obesity Symposium. American Journal of Clinical Nutrition, 2022, 115, 18-33.	4.7	43
62	Validation of a DNA methylation reference panel for the estimation of nucleated cells types in cord blood. Epigenetics, 2016, 11, 773-779.	2.7	42
63	Placental DNA Methylation Adaptation to Maternal Glycemic Response in Pregnancy. Diabetes, 2018, 67, 1673-1683.	0.6	42
64	Primary Prevention of ASCVD and T2DM in Patients at Metabolic Risk: An Endocrine Society* Clinical Practice Guideline. Journal of Clinical Endocrinology and Metabolism, 2019, 104, 3939-3985.	3.6	42
65	DNA methylation and body mass index from birth to adolescence: meta-analyses of epigenome-wide association studies. Genome Medicine, 2020, 12, 105.	8.2	41
66	Placental DNA methylation signatures of maternal smoking during pregnancy and potential impacts on fetal growth. Nature Communications, 2021, 12, 5095.	12.8	41
67	Higher maternal leptin levels at second trimester are associated with subsequent greater gestational weight gain in late pregnancy. BMC Pregnancy and Childbirth, 2016, 16, 62.	2.4	40
68	Associations of Prenatal and Postnatal Maternal Depressive Symptoms with Offspring Cognition and Behavior in Mid-Childhood: A Prospective Cohort Study. International Journal of Environmental Research and Public Health, 2019, 16, 1007.	2.6	40
69	Who will deliver comprehensive healthy lifestyle interventions to combat non-communicable disease? Introducing the healthy lifestyle practitioner discipline. Expert Review of Cardiovascular Therapy, 2016, 14, 15-22.	1.5	39
70	First and second trimester gestational weight gains are most strongly associated with cord blood levels of hormones at delivery important for glycemic control and somatic growth. Metabolism: Clinical and Experimental, 2017, 69, 112-119.	3.4	38
71	Placental lipoprotein lipase DNA methylation alterations are associated with gestational diabetes and body composition at 5Âyears of age. Epigenetics, 2017, 12, 616-625.	2.7	38
72	Timing of Complementary Feeding Introduction and Adiposity Throughout Childhood. Pediatrics, 2019, 144	2.1	38

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73	Epigenetic dysregulation of the IGF system in placenta of newborns exposed to maternal impaired glucose tolerance. Epigenomics, 2014, 6, 193-207.	2.1	37
74	DNA methylation of blood cells is associated with prevalent type 2 diabetes in a meta-analysis of four European cohorts. Clinical Epigenetics, 2021, 13, 40.	4.1	37
75	Pre-, Perinatal, and Parental Predictors of Body Mass Index Trajectory Milestones. Journal of Pediatrics, 2018, 201, 69-77.e8.	1.8	36
76	Association of Weight for Length vs Body Mass Index During the First 2 Years of Life With Cardiometabolic Risk in Early Adolescence. JAMA Network Open, 2018, 1, e182460.	5.9	35
77	Maternal lipid profile differs by gestational diabetes physiologic subtype. Metabolism: Clinical and Experimental, 2019, 91, 39-42.	3.4	35
78	TNFα Dynamics During the Oral Glucose Tolerance Test Vary According to the Level of Insulin Resistance in Pregnant Women. Journal of Clinical Endocrinology and Metabolism, 2014, 99, 1862-1869.	3.6	34
79	Lifestyle and Metformin Ameliorate Insulin Sensitivity Independently of the Genetic Burden of Established Insulin Resistance Variants in Diabetes Prevention Program Participants. Diabetes, 2016, 65, 520-526.	0.6	34
80	Interplay of Placental DNA Methylation and Maternal Insulin Sensitivity in Pregnancy. Diabetes, 2020, 69, 484-492.	0.6	34
81	Mendelian Randomization Analysis of Hemoglobin A1c as a Risk Factor for Coronary Artery Disease. Diabetes Care, 2019, 42, 1202-1208.	8.6	33
82	Branched Chain Amino Acids, Androgen Hormones, and Metabolic Risk Across Early Adolescence: A Prospective Study in Project Viva. Obesity, 2018, 26, 916-926.	3.0	31
83	The association of tumor necrosis factor $\hat{I}\pm$ receptor 2 and tumor necrosis factor $\hat{I}\pm$ with insulin resistance and the influence of adipose tissue biomarkers in humans. Metabolism: Clinical and Experimental, 2010, 59, 540-546.	3.4	30
84	Cross-tissue comparisons of leptin and adiponectin. Adipocyte, 2014, 3, 132-140.	2.8	30
85	Genetic Determinants of Glycemic Traits and the Risk of Gestational Diabetes Mellitus. Diabetes, 2018, 67, 2703-2709.	0.6	30
86	Defining Heterogeneity Among Women With Gestational Diabetes Mellitus. Diabetes, 2020, 69, 2064-2074.	0.6	29
87	Dietary characteristics associated with plasma concentrations of per- and polyfluoroalkyl substances among adults with pre-diabetes: Cross-sectional results from the Diabetes Prevention Program Trial. Environment International, 2020, 137, 105217.	10.0	28
88	Prospective Associations of Early Pregnancy Metal Mixtures with Mitochondria DNA Copy Number and Telomere Length in Maternal and Cord Blood. Environmental Health Perspectives, 2021, 129, 117007.	6.0	28
89	Preeclampsia is associated with an increased pro-inflammatory profile in newborns. Journal of Reproductive Immunology, 2015, 112, 111-114.	1.9	27
90	Early pregnancy exposure to metal mixture and birth outcomes – A prospective study in Project Viva. Environment International, 2021, 156, 106714.	10.0	27

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91	Early-pregnancy plasma per- and polyfluoroalkyl substance (PFAS) concentrations and hypertensive disorders of pregnancy in the Project Viva cohort. Environment International, 2022, 165, 107335.	10.0	27
92	Leptin trajectories from birth to mid-childhood and cardio-metabolic health in early adolescence. Metabolism: Clinical and Experimental, 2019, 91, 30-38.	3.4	26
93	Longitudinal Changes in the Relationship Between Hemoglobin A1c and Clucose Tolerance Across Pregnancy and Postpartum. Journal of Clinical Endocrinology and Metabolism, 2020, 105, e1999-e2007.	3.6	26
94	<i>LRP1B, BRD2</i> and <i>CACNA1D</i> : new candidate genes in fetal metabolic programming of newborns exposed to maternal hyperglycemia. Epigenomics, 2015, 7, 1111-1122.	2.1	24
95	Per- and polyfluoroalkyl substances and blood pressure in pre-diabetic adults—cross-sectional and longitudinal analyses of the diabetes prevention program outcomes study. Environment International, 2020, 137, 105573.	10.0	24
96	Maternal anxiety during pregnancy and newborn epigenome-wide DNA methylation. Molecular Psychiatry, 2021, 26, 1832-1845.	7.9	24
97	Per- and polyfluoroalkyl substances and kidney function: Follow-up results from the Diabetes Prevention Program trial. Environment International, 2021, 148, 106375.	10.0	24
98	Temporal trends of concentrations of per- and polyfluoroalkyl substances among adults with overweight and obesity in the United States: Results from the Diabetes Prevention Program and NHANES. Environment International, 2021, 157, 106789.	10.0	24
99	Lifestyle interventions in pregnancy targeting GDM prevention: looking ahead to precision medicine. Diabetologia, 2022, 65, 1814-1824.	6.3	24
100	Gestational Perfluoroalkyl Substance Exposure and DNA Methylation at Birth and 12 Years of Age: A Longitudinal Epigenome-Wide Association Study. Environmental Health Perspectives, 2022, 130, 37005.	6.0	24
101	Peripheral Blood Transcriptomic Signatures of Fasting Clucose and Insulin Concentrations. Diabetes, 2016, 65, 3794-3804.	0.6	22
102	funtooNorm: an R package for normalization of DNA methylation data when there are multiple cell or tissue types. Bioinformatics, 2016, 32, 593-595.	4.1	22
103	Associations of prenatal or infant exposure to acetaminophen or ibuprofen with midâ€childhood executive function and behaviour. Paediatric and Perinatal Epidemiology, 2020, 34, 287-298.	1.7	22
104	Metabolomic Profiles of Overweight/Obesity Phenotypes During Adolescence: A Crossâ€5ectional Study in Project Viva. Obesity, 2020, 28, 379-387.	3.0	22
105	Hypertensive Disorders of Pregnancy and Offspring Cardiometabolic Health at Midchildhood: Project Viva Findings. Journal of the American Heart Association, 2018, 7, .	3.7	21
106	Characterization of longitudinal wheeze phenotypes from infancy to adolescence in Project Viva, a prebirth cohort study. Journal of Allergy and Clinical Immunology, 2020, 145, 716-719.e8.	2.9	21
107	Higher Adiponectin Levels Predict Greater Weight Gain in Healthy Women in the Nurses' Health Study. Obesity, 2011, 19, 409-415.	3.0	20
108	Timing of Excessive Weight Gain During Pregnancy Modulates Newborn Anthropometry. Journal of Obstetrics and Gynaecology Canada, 2016, 38, 108-117.	0.7	20

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109	Early-Life Exposures and Risk of Diabetes Mellitus and Obesity. Current Diabetes Reports, 2018, 18, 89.	4.2	20
110	Diet and erythrocyte metal concentrations in early pregnancy—cross-sectional analysis in Project Viva. American Journal of Clinical Nutrition, 2021, 114, 540-549.	4.7	20
111	Prenatal metal exposure, cord blood DNA methylation and persistence in childhood: an epigenome-wide association study of 12 metals. Clinical Epigenetics, 2021, 13, 208.	4.1	20
112	Mediation Analysis Supports a Causal Relationship between Maternal Hyperglycemia and Placental DNA Methylation Variations at the Leptin Gene Locus and Cord Blood Leptin Levels. International Journal of Molecular Sciences, 2020, 21, 329.	4.1	19
113	Maternal Glycemic Dysregulation During Pregnancy and Neonatal Blood DNA Methylation: Meta-analyses of Epigenome-Wide Association Studies. Diabetes Care, 2022, 45, 614-623.	8.6	19
114	Cord blood DNA methylation and adiposity measures in early and mid-childhood. Clinical Epigenetics, 2017, 9, 86.	4.1	18
115	Associations of Gestational Glucose Tolerance With Offspring Body Composition and Estimated Insulin Resistance in Early Adolescence. Diabetes Care, 2018, 41, e164-e166.	8.6	18
116	DNA methylation mediates the association between breastfeeding and early-life growth trajectories. Clinical Epigenetics, 2021, 13, 231.	4.1	18
117	Metabolic trajectories across early adolescence: differences by sex, weight, pubertal status and race/ethnicity. Annals of Human Biology, 2019, 46, 205-214.	1.0	17
118	Polygenic risk score for obesity and the quality, quantity, and timing of workplace food purchases: A secondary analysis from the ChooseWell 365 randomized trial. PLoS Medicine, 2020, 17, e1003219.	8.4	17
119	Human plasma pregnancy-associated miRNAs and their temporal variation within the first trimester of pregnancy. Reproductive Biology and Endocrinology, 2022, 20, 14.	3.3	17
120	Early Infant Nutrition and Metabolic Programming: What Are the Potential Molecular Mechanisms?. Current Nutrition Reports, 2014, 3, 281-288.	4.3	16
121	Parent-of-Origin Effects of the APOB Gene on Adiposity in Young Adults. PLoS Genetics, 2015, 11, e1005573.	3.5	16
122	Pregnancy Per- and Polyfluoroalkyl Substance Concentrations and Postpartum Health in Project Viva: A Prospective Cohort. Journal of Clinical Endocrinology and Metabolism, 2020, 105, e3415-e3426.	3.6	16
123	The Obesity-Fertility Protocol: a randomized controlled trial assessing clinical outcomes and costs of a transferable interdisciplinary lifestyle intervention, before and during pregnancy, in obese infertile women. BMC Obesity, 2015, 2, 47.	3.1	15
124	A qualitative study of gestational weight gain goal setting. BMC Pregnancy and Childbirth, 2016, 16, 317.	2.4	15
125	Associations of maternal prenatal smoking with umbilical cord blood hormones: the Project Viva cohort. Metabolism: Clinical and Experimental, 2017, 72, 18-26.	3.4	15
126	Supporting healthful lifestyles during pregnancy: a health coach intervention pilot study. BMC Pregnancy and Childbirth, 2018, 18, 375.	2.4	15

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127	Comparison of Illumina 450K and EPIC arrays in placental DNA methylation. Epigenetics, 2019, 14, 1177-1182.	2.7	15
128	Self-Monitoring of Blood Glucose: A Complementary Method Beyond the Oral Glucose Tolerance Test to Identify Hyperglycemia During Pregnancy. Canadian Journal of Diabetes, 2019, 43, 627-635.	0.8	14
129	Early life exposure to green space and insulin resistance: An assessment from infancy to early adolescence. Environment International, 2020, 142, 105849.	10.0	14
130	A Polygenic Lipodystrophy Genetic Risk Score Characterizes Risk Independent of BMI in the Diabetes Prevention Program. Journal of the Endocrine Society, 2019, 3, 1663-1677.	0.2	13
131	Early pregnancy essential and non-essential metal mixtures and gestational glucose concentrations in the 2nd trimester: Results from project viva. Environment International, 2021, 155, 106690.	10.0	13
132	Analysis of Early-Life Growth and Age at Pubertal Onset in US Children. JAMA Network Open, 2022, 5, e2146873.	5.9	13
133	Genetic ancestry markers and difference in A1c between African-American and White in the Diabetes Prevention Program. Journal of Clinical Endocrinology and Metabolism, 2019, 104, 328-336.	3.6	12
134	Impact of Genetic Determinants of HbA1c on Type 2 Diabetes Risk and Diagnosis. Current Diabetes Reports, 2018, 18, 52.	4.2	12
135	Placental surface area mediates the association between FGFR2 methylation in placenta and full-term low birth weight in girls. Clinical Epigenetics, 2018, 10, 39.	4.1	12
136	Locus-specific DNA methylation prediction in cord blood and placenta. Epigenetics, 2019, 14, 405-420.	2.7	12
137	Detecting differentially methylated regions with multiple distinct associations. Epigenomics, 2021, 13, 451-464.	2.1	12
138	Calcifediol Decreases Interleukin-6 Secretion by Cultured Human Trophoblasts From GDM Pregnancies. Journal of the Endocrine Society, 2019, 3, 2165-2178.	0.2	11
139	Per- and polyfluoroalkyl substances and calcifications of the coronary and aortic arteries in adults with prediabetes: Results from the diabetes prevention program outcomes study. Environment International, 2021, 151, 106446.	10.0	11
140	DNA methylation changes associated with prenatal mercury exposure: A meta-analysis of prospective cohort studies from PACE consortium. Environmental Research, 2022, 204, 112093.	7.5	11
141	Gestational Diabetes Mellitus Identification Based on Self-Monitoring of Blood Glucose. Canadian Journal of Diabetes, 2015, 39, 162-168.	0.8	10
142	Training Health Professionals to Deliver Healthy Living Medicine. Progress in Cardiovascular Diseases, 2017, 59, 471-478.	3.1	10
143	Genetic determinants of adiponectin regulation revealed by pregnancy. Obesity, 2017, 25, 935-944.	3.0	10
144	Comparison of novel and existing methods for detecting differentially methylated regions. BMC Genetics, 2018, 19, 84.	2.7	10

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145	Genetic Loci and Physiologic Pathways Involved in Gestational Diabetes Mellitus Implicated Through Clustering. Diabetes, 2021, 70, 268-281.	0.6	10
146	Residential PM2.5 exposure and the nasal methylome in children. Environment International, 2021, 153, 106505.	10.0	10
147	Early life exposure to greenness and executive function and behavior: An application of inverse probability weighting of marginal structural models. Environmental Pollution, 2021, 291, 118208.	7.5	10
148	${\sf HNF1\hat{l}}\pm$ defect influences post-prandial lipid regulation. PLoS ONE, 2017, 12, e0177110.	2.5	10
149	Tissue differences in DNA methylation changes at AHRR in full term low birth weight in maternal blood, placenta and cord blood in Chinese. Placenta, 2017, 52, 49-57.	1.5	9
150	Associations of prenatal exposure to impaired glucose tolerance with eating in the absence of hunger in early adolescence. International Journal of Obesity, 2019, 43, 1903-1913.	3.4	9
151	Placental Epigenome-Wide Association Study Identified Loci Associated with Childhood Adiposity at 3 Years of Age. International Journal of Molecular Sciences, 2020, 21, 7201.	4.1	9
152	Separating Algorithms From Questions and Causal Inference With Unmeasured Exposures: An Application to Birth Cohort Studies of Early Body Mass Index Rebound. American Journal of Epidemiology, 2021, 190, 1414-1423.	3.4	9
153	Dietary fat intake during early pregnancy is associated with cord blood DNA methylation at <i>IGF2</i> and <i>H19</i> genes in newborns. Environmental and Molecular Mutagenesis, 2021, 62, 388-398.	2.2	9
154	Impact of the Creation of a Specialized Clinic for Prenatal Blood Sampling and Follow-up Care in Pregnant Women. Journal of Obstetrics and Gynaecology Canada, 2012, 34, 236-242.	0.7	8
155	Parental Obesity and Offspring Pubertal Development: Project Viva. Journal of Pediatrics, 2019, 215, 123-131.e2.	1.8	8
156	DNA methylation at <i>LRP1</i> gene locus mediates the association between maternal total cholesterol changes in pregnancy and cord blood leptin levels. Journal of Developmental Origins of Health and Disease, 2020, 11, 369-378.	1.4	8
157	Childhood patterns of overweight and wheeze and subsequent risk of current asthma and obesity in adolescence. Paediatric and Perinatal Epidemiology, 2021, 35, 569-577.	1.7	8
158	Maternal Mediterranean diet in pregnancy and newborn DNA methylation: a meta-analysis in the PACE Consortium. Epigenetics, 2022, 17, 1419-1431.	2.7	8
159	Mode of delivery, type of labor, and measures of adiposity from childhood to teenage: Project Viva. International Journal of Obesity, 2021, 45, 36-44.	3.4	7
160	Maternal glucose tolerance in pregnancy and child cognitive and behavioural problems in early and midâ€childhood. Paediatric and Perinatal Epidemiology, 2021, 35, 109-119.	1.7	7
161	Per- and polyfluoroalkyl substance plasma concentrations and metabolomic markers of type 2 diabetes in the Diabetes Prevention Program trial. International Journal of Hygiene and Environmental Health, 2021, 232, 113680.	4.3	7
162	Physiological subtypes of gestational glucose intolerance and risk of adverse pregnancy outcomes. American Journal of Obstetrics and Gynecology, 2022, 226, 241.e1-241.e14.	1.3	7

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163	Mid-Pregnancy Fructosamine Measurement—Predictive Value for Gestational Diabetes and Association with Postpartum Glycemic Indices. Nutrients, 2018, 10, 2003.	4.1	6
164	Maternal corticotropin-releasing hormone is associated with LEP DNA methylation at birth and in childhood: an epigenome-wide study in Project Viva. International Journal of Obesity, 2019, 43, 1244-1255.	3.4	6
165	Associations of sleep duration, sedentary behaviours and energy expenditure with maternal glycemia in pregnancy. Sleep Medicine, 2020, 65, 54-61.	1.6	6
166	Associations of Early Parental Concerns and Feeding Behaviors with Child's Diet Quality through Mid-Childhood. Nutrients, 2020, 12, 3231.	4.1	6
167	First trimester plasma microRNAs levels predict Matsuda Index-estimated insulin sensitivity between 24th and 29th week of pregnancy. BMJ Open Diabetes Research and Care, 2022, 10, e002703.	2.8	6
168	Association of cow's milk intake in early childhood with adiposity and cardiometabolic risk in early adolescence. American Journal of Clinical Nutrition, 2022, 116, 561-571.	4.7	6
169	DNA Methylation and Type 2 Diabetes: the Use of Mendelian Randomization to Assess Causality. Current Genetic Medicine Reports, 2019, 7, 191-207.	1.9	5
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