## Janine F Felix

List of Publications by Year in descending order

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38742 31849 12,255 157 50 citations h-index papers

g-index 164 164 164 19358 docs citations times ranked citing authors all docs

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#	Article	IF	CITATIONS
1	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
2	New loci associated with kidney function and chronic kidney disease. Nature Genetics, 2010, 42, 376-384.	21.4	710
3	The Generation R Study: design and cohort update 2017. European Journal of Epidemiology, 2016, 31, 1243-1264.	5 <b>.</b> 7	608
4	A catalog of genetic loci associated with kidney function from analyses of a million individuals. Nature Genetics, 2019, 51, 957-972.	21.4	549
5	Genetic associations at 53 loci highlight cell types and biological pathways relevant for kidney function. Nature Communications, 2016, 7, 10023.	12.8	412
6	Genome-wide associations for birth weight and correlations with adult disease. Nature, 2016, 538, 248-252.	27.8	406
7	Genome-wide association study identifies six new loci influencing pulse pressure and mean arterial pressure. Nature Genetics, 2011, 43, 1005-1011.	21.4	403
8	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
9	Association Between Telomere Length and Risk of Cancer and Non-Neoplastic Diseases. JAMA Oncology, 2017, 3, 636.	7.1	376
10	The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. Nature Genetics, 2016, 48, 1171-1184.	21.4	362
11	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. Nature Genetics, 2016, 48, 1462-1472.	21.4	284
12	Genome-wide association analysis identifies three new susceptibility loci for childhood body mass index. Human Molecular Genetics, 2016, 25, 389-403.	2.9	275
13	Trans-ancestry meta-analyses identify rare and common variants associated with blood pressure and hypertension. Nature Genetics, 2016, 48, 1151-1161.	21.4	261
14	Life-Course Genome-wide Association Study Meta-analysis of Total Body BMD and Assessment of Age-Specific Effects. American Journal of Human Genetics, 2018, 102, 88-102.	6.2	252
15	The Giessen Pulmonary Hypertension Registry: Survival in pulmonary hypertension subgroups. Journal of Heart and Lung Transplantation, 2017, 36, 957-967.	0.6	221
16	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
17	Maternal plasma folate impacts differential DNA methylation in an epigenome-wide meta-analysis of newborns. Nature Communications, 2016, 7, 10577.	12.8	219
18	Genomic and phenotypic insights from an atlas of genetic effects on DNA methylation. Nature Genetics, 2021, 53, 1311-1321.	21.4	218

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19	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
20	Genetic Variants Associated With Cardiac Structure and Function. JAMA - Journal of the American Medical Association, 2009, 302, 168.	7.4	202
21	Phenotypic Characterization of GeneticallyÂLowered Human Lipoprotein(a) Levels. Journal of the American College of Cardiology, 2016, 68, 2761-2772.	2.8	186
22	Epigenome-Wide Meta-Analysis of Methylation in Children Related to Prenatal NO <sub>2</sub> Air Pollution Exposure. Environmental Health Perspectives, 2017, 125, 104-110.	6.0	176
23	DNA methylation mediates the effect of maternal smoking during pregnancy on birthweight of the offspring. International Journal of Epidemiology, 2015, 44, 1224-1237.	1.9	172
24	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
25	Epigenome-wide meta-analysis of DNA methylation and childhood asthma. Journal of Allergy and Clinical Immunology, 2019, 143, 2062-2074.	2.9	147
26	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
27	Genome-wide association meta-analyses and fine-mapping elucidate pathways influencing albuminuria. Nature Communications, 2019, 10, 4130.	12.8	133
28	Whole-Genome Sequencing Coupled to Imputation Discovers Genetic Signals for Anthropometric Traits. American Journal of Human Genetics, 2017, 100, 865-884.	6.2	131
29	Cystatin C and Cardiovascular Disease. Journal of the American College of Cardiology, 2016, 68, 934-945.	2.8	109
30	Systematic evaluation and validation of reference and library selection methods for deconvolution of cord blood DNA methylation data. Clinical Epigenetics, 2019, 11, 125.	4.1	107
31	Cohort Profile: Pregnancy And Childhood Epigenetics (PACE) Consortium. International Journal of Epidemiology, 2018, 47, 22-23u.	1.9	105
32	Genetic loci associated with heart rate variability and their effects on cardiac disease risk. Nature Communications, 2017, 8, 15805.	12.8	95
33	Novel loci for childhood body mass index and shared heritability with adult cardiometabolic traits. PLoS Genetics, 2020, 16, e1008718.	3.5	95
34	Etiology of Esophageal Atresia and Tracheoesophageal Fistula: "Mind the Gap― Current Gastroenterology Reports, 2010, 12, 215-222.	2.5	88
35	Effects of choline on health across the life course: a systematic review. Nutrition Reviews, 2015, 73, 500-522.	5.8	87
36	GWAS on longitudinal growth traits reveals different genetic factors influencing infant, child, and adult BMI. Science Advances, 2019, 5, eaaw3095.	10.3	86

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37	Associations of autozygosity with a broad range of human phenotypes. Nature Communications, 2019, 10, 4957.	12.8	84
38	The LifeCycle Project-EU Child Cohort Network: a federated analysis infrastructure and harmonized data of more than 250,000 children and parents. European Journal of Epidemiology, 2020, 35, 709-724.	5.7	81
39	Epigenome-wide meta-analysis of blood DNA methylation in newborns and children identifies numerous loci related to gestational age. Genome Medicine, 2020, 12, 25.	8.2	81
40	An epigenome-wide association meta-analysis of prenatal maternal stress in neonates: A model approach for replication. Epigenetics, 2016, 11, 140-149.	2.7	80
41	A trans-ancestral meta-analysis of genome-wide association studies reveals loci associated with childhood obesity. Human Molecular Genetics, 2019, 28, 3327-3338.	2.9	76
42	Nonâ€VACTERLâ€type anomalies are frequent in patients with esophageal atresia/tracheoâ€esophageal fistula and full or partial VACTERL association. Birth Defects Research Part A: Clinical and Molecular Teratology, 2008, 82, 92-97.	1.6	73
43	Hypertensive Disorders of Pregnancy and DNA Methylation in Newborns. Hypertension, 2019, 74, 375-383.	2.7	73
44	Chromosomal anomalies in the aetiology of oesophageal atresia and tracheo-oesophageal fistula. European Journal of Medical Genetics, 2007, 50, 163-175.	1.3	71
45	Using Genetic Variation to Explore the Causal Effect of Maternal Pregnancy Adiposity on Future Offspring Adiposity: A Mendelian Randomisation Study. PLoS Medicine, 2017, 14, e1002221.	8.4	71
46	Cell type specific DNA methylation in cord blood: A 450K-reference data set and cell count-based validation of estimated cell type composition. Epigenetics, 2016, 11, 690-698.	2.7	69
47	Genetic and environmental factors in the etiology of esophageal atresia and/or tracheoesophageal fistula: An overview of the current concepts. Birth Defects Research Part A: Clinical and Molecular Teratology, 2009, 85, 747-754.	1.6	68
48	Epigenome-wide change and variation in DNA methylation in childhood: trajectories from birth to late adolescence. Human Molecular Genetics, 2021, 30, 119-134.	2.9	65
49	Comparison of smoking-related DNA methylation between newborns from prenatal exposure and adults from personal smoking. Epigenomics, 2019, 11, 1487-1500.	2.1	64
50	An integrative cross-omics analysis of DNA methylation sites of glucose and insulin homeostasis. Nature Communications, 2019, 10, 2581.	12.8	62
51	Associations of maternal quitting, reducing, and continuing smoking during pregnancy with longitudinal fetal growth: Findings from Mendelian randomization and parental negative control studies. PLoS Medicine, 2019, 16, e1002972.	8.4	62
52	Maternal alcohol consumption and offspring DNA methylation: findings from six general population-based birth cohorts. Epigenomics, 2018, 10, 27-42.	2.1	58
53	Prevalence of Pulmonary Hypertension in the General Population: The Rotterdam Study. PLoS ONE, 2015, 10, e0130072.	2.5	57
54	Association between DNA methylation and ADHD symptoms from birth to school age: a prospective meta-analysis. Translational Psychiatry, 2020, 10, 398.	4.8	54

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55	Early- and late-onset preeclampsia and the tissue-specific epigenome of the placenta and newborn. Placenta, 2017, 58, 122-132.	1.5	52
56	Variants in the fetal genome near pro-inflammatory cytokine genes on 2q13 associate with gestational duration. Nature Communications, 2019, 10, 3927.	12.8	49
57	Body mass index, gestational weight gain and fatty acid concentrations during pregnancy: the Generation R Study. European Journal of Epidemiology, 2015, 30, 1175-1185.	5.7	48
58	Newborn DNA-methylation, childhood lung function, and the risks of asthma and COPD across the life course. European Respiratory Journal, 2019, 53, 1801795.	6.7	48
59	Prenatal maternal antidepressants, anxiety, and depression and offspring DNA methylation: epigenome-wide associations at birth and persistence into early childhood. Clinical Epigenetics, 2019, 11, 56.	4.1	46
60	Genome-wide Trans-ethnic Meta-analysis Identifies Seven Genetic Loci Influencing Erythrocyte Traits and a Role for RBPMS in Erythropoiesis. American Journal of Human Genetics, 2017, 100, 51-63.	6.2	45
61	Epigenome-wide association study reveals methylation pathways associated with childhood allergic sensitization. Epigenetics, 2019, 14, 445-466.	2.7	43
62	Association of Birth Weight With Type 2 Diabetes and Glycemic Traits. JAMA Network Open, 2019, 2, e1910915.	5.9	41
63	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
64	Environmental factors in the etiology of esophageal atresia and congenital diaphragmatic hernia: Results of a caseâ€control study. Birth Defects Research Part A: Clinical and Molecular Teratology, 2008, 82, 98-105.	1.6	40
65	Genome-wide association study identifies nine novel loci for 2D:4D finger ratio, a putative retrospective biomarker of testosterone exposure in utero. Human Molecular Genetics, 2018, 27, 2025-2038.	2.9	36
66	Discovery of Genetic Variation on Chromosome 5q22 Associated with Mortality in Heart Failure. PLoS Genetics, 2016, 12, e1006034.	3.5	34
67	Retinal Microvasculature and Cardiovascular Health in Childhood. Pediatrics, 2015, 135, 678-685.	2.1	31
68	Validated inference of smoking habits from blood with a finite DNA methylation marker set. European Journal of Epidemiology, 2019, 34, 1055-1074.	5.7	31
69	Associations of Fetal and Infant Weight Change With General, Visceral, and Organ Adiposity at School Age. JAMA Network Open, 2019, 2, e192843.	5.9	31
70	Vitamin D and risk of pregnancy related hypertensive disorders: mendelian randomisation study. BMJ: British Medical Journal, 2018, 361, k2167.	2.3	31
71	Ethnic disparities in maternal obesity and weight gain during pregnancy. The Generation R Study. European Journal of Obstetrics, Gynecology and Reproductive Biology, 2015, 193, 51-60.	1.1	30
72	Influence of Maternal Angiogenic Factors During Pregnancy on Microvascular Structure in School-Age Children. Hypertension, 2015, 65, 722-728.	2.7	30

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73	Low-frequency variation in TP53 has large effects on head circumference and intracranial volume. Nature Communications, 2019, 10, 357.	12.8	30
74	The effects of lutein on respiratory health across the life course: AÂsystematic review. Clinical Nutrition ESPEN, 2016, 13, e1-e7.	1.2	28
75	The Early Growth Genetics (EGG) and EArly Genetics and Lifecourse Epidemiology (EAGLE) consortia: design, results and future prospects. European Journal of Epidemiology, 2019, 34, 279-300.	5.7	26
76	Evaluation of commonly used analysis strategies for epigenome- and transcriptome-wide association studies through replication of large-scale population studies. Genome Biology, 2019, 20, 235.	8.8	26
77	Maternal body mass index, gestational weight gain, and childhood abdominal, pericardial, and liver fat assessed by magnetic resonance imaging. International Journal of Obesity, 2019, 43, 581-593.	3.4	26
78	Aptamer-Based Proteomic Platform Identifies Novel Protein Predictors of Incident Heart Failure and Echocardiographic Traits. Circulation: Heart Failure, 2020, 13, e006749.	3.9	26
79	Liver Fat and Cardiometabolic Risk Factors Among Schoolâ€Age Children. Hepatology, 2020, 72, 119-129.	7.3	25
80	Agenesis of the trachea: Phenotypic expression of a rare cause of fatal neonatal respiratory insufficiency in six patients. International Journal of Pediatric Otorhinolaryngology, 2006, 70, 365-370.	1.0	24
81	Folate, vitamin B12, and homocysteine in smokingâ€exposed pregnant women: A systematic review. Maternal and Child Nutrition, 2019, 15, e12675.	3.0	24
82	Maternal anxiety during pregnancy and newborn epigenome-wide DNA methylation. Molecular Psychiatry, 2021, 26, 1832-1845.	7.9	24
83	The EU Child Cohort Network's core data: establishing a set of findable, accessible, interoperable and re-usable (FAIR) variables. European Journal of Epidemiology, 2021, 36, 565-580.	5.7	24
84	Meta-analysis of epigenome-wide association studies in newborns and children show widespread sex differences in blood DNA methylation. Mutation Research - Reviews in Mutation Research, 2022, 789, 108415.	5.5	24
85	Esophageal atresia and tracheoesophageal fistula in children of women exposed to diethylstilbestrol in utero. American Journal of Obstetrics and Gynecology, 2007, 197, 38.e1-38.e5.	1.3	21
86	Associations of genetic risk scores based on adult adiposity pathways with childhood growth and adiposity measures. BMC Genetics, 2016, 17, 120.	2.7	21
87	DNA methylation signatures of aggression and closely related constructs: A meta-analysis of epigenome-wide studies across the lifespan. Molecular Psychiatry, 2021, 26, 2148-2162.	7.9	21
88	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
89	Exploring the role of genetic confounding in the association between maternal and offspring body mass index: evidence from three birth cohorts. International Journal of Epidemiology, 2020, 49, 233-243.	1.9	18
90	Differential and shared genetic effects on kidney function between diabetic and non-diabetic individuals. Communications Biology, 2022, 5, .	4.4	17

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91	Associations of maternal and fetal vitamin D status with childhood body composition and cardiovascular risk factors. Maternal and Child Nutrition, 2019, 15, e12672.	3.0	16
92	Epigenomics of being bullied: changes in DNA methylation following bullying exposure. Epigenetics, 2020, 15, 750-764.	2.7	16
93	Health in children: A conceptual framework for use in healthy ageing research. Maturitas, 2014, 77, 47-51.	2.4	15
94	Sildenafil versus Nitric Oxide for Acute Vasodilator Testing in Pulmonary Arterial Hypertension. Pulmonary Circulation, 2015, 5, 305-312.	1.7	15
95	Influence of genetic variants associated with body mass index on eating behavior in childhood. Obesity, 2017, 25, 765-772.	3.0	15
96	Residential Proximity to Major Roadways at Birth, DNA Methylation at Birth and Midchildhood, and Childhood Cognitive Test Scores: Project Viva(Massachusetts, USA). Environmental Health Perspectives, 2018, 126, 97006.	6.0	15
97	Genomeâ€wide DNA methylation patterns associated with sleep and mental health in children: a populationâ€based study. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2020, 61, 1061-1069.	<b>5.</b> 2	15
98	Effects of protein intake on blood pressure, insulin sensitivity and blood lipids in children: a systematic review. British Journal of Nutrition, 2015, 113, 383-402.	2.3	14
99	Body Fat Distribution, Overweight, and Cardiac Structures in Schoolâ€Age Children: A Populationâ€Based Cardiac Magnetic Resonance Imaging Study. Journal of the American Heart Association, 2020, 9, e014933.	3.7	14
100	Associations of Maternal Psychological Distress during Pregnancy with Childhood General and Organ Fat Measures. Childhood Obesity, 2019, 15, 313-322.	1.5	13
101	Associations of maternal early-pregnancy blood glucose and insulin concentrations with DNA methylation in newborns. Clinical Epigenetics, 2020, 12, 134.	4.1	13
102	A population-based resource for intergenerational metabolomics analyses in pregnant women and their children: the Generation R Study. Metabolomics, 2020, $16$ , $43$ .	3.0	13
103	The tissue-specific aspect of genome-wide DNA methylation in newborn and placental tissues: implications for epigenetic epidemiologic studies. Journal of Developmental Origins of Health and Disease, 2021, 12, 113-123.	1.4	13
104	Associations Between Intake of Sugarâ€Containing Beverages in Infancy With Liver Fat Accumulation at School Age. Hepatology, 2021, 73, 560-570.	7.3	13
105	Meta-analysis of epigenome-wide associations between DNA methylation at birth and childhood cognitive skills. Molecular Psychiatry, 2022, 27, 2126-2135.	7.9	13
106	Cardioprotective Effects of <i>MTSS1</i> Enhancer Variants. Circulation, 2019, 139, 2073-2076.	1.6	12
107	Epigenome-wide association study of seizures in childhood and adolescence. Clinical Epigenetics, 2020, 12, 8.	4.1	12
108	Newborn and childhood differential DNA methylation and liver fat in school-age children. Clinical Epigenetics, 2020, 12, 3.	4.1	12

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109	Maternal Body Mass Index, Early-Pregnancy Metabolite Profile, and Birthweight. Journal of Clinical Endocrinology and Metabolism, 2022, 107, e315-e327.	3.6	11
110	Sugar-containing beverage intake at the age of 1Âyear and cardiometabolic health at the age of 6Âyears: the Generation R Study. International Journal of Behavioral Nutrition and Physical Activity, 2015, 12, 114.	4.6	10
111	Early origins of ethnic disparities in cardiovascular risk factors. Preventive Medicine, 2015, 76, 84-91.	3.4	10
112	Influence of genetic variants on childhood lung function – The Generation R Study. Pediatric Allergy and Immunology, 2018, 29, 589-595.	2.6	10
113	Cohort Profile: The DynaHEALTH consortium – a European consortium for a life-course bio-psychosocial model of healthy ageing of glucose homeostasis. International Journal of Epidemiology, 2019, 48, 1051-1051k.	1.9	10
114	Psychological Distress and Weight Gain in Pregnancy: a Population-Based Study. International Journal of Behavioral Medicine, 2020, 27, 30-38.	1.7	10
115	Histological, immunohistochemical and transcriptomic characterization of human tracheoesophageal fistulas. PLoS ONE, 2020, 15, e0242167.	2.5	10
116	Maternal Dietary Glycemic Index and Glycemic Load in Pregnancy and Offspring Cord Blood DNA Methylation. Diabetes Care, 2022, 45, 1822-1832.	8.6	10
117	Associations of genetic variants for adult lipid levels with lipid levels in children. The Generation R Study. Journal of Lipid Research, 2016, 57, 2185-2192.	4.2	9
118	Altered DNA methylation in children born to mothers with rheumatoid arthritis during pregnancy. Annals of the Rheumatic Diseases, 2019, 78, 1198-1204.	0.9	9
119	Cord blood DNA methylation reflects cord blood C-reactive protein levels but not maternal levels: a longitudinal study and meta-analysis. Clinical Epigenetics, 2020, 12, 60.	4.1	9
120	Mendelian randomization analysis does not support causal associations of birth weight with hypertension risk and blood pressure in adulthood. European Journal of Epidemiology, 2020, 35, 685-697.	5.7	9
121	Associations of Hair Cortisol Concentrations with General and Organ Fat Measures in Childhood. Journal of Clinical Endocrinology and Metabolism, 2021, 106, e551-e561.	3.6	9
122	Identifying causative mechanisms linking early-life stress to psycho-cardio-metabolic multi-morbidity: The EarlyCause project. PLoS ONE, 2021, 16, e0245475.	2.5	9
123	Associations of Early Pregnancy and Neonatal Circulating Folate, Vitamin B-12, and Homocysteine Concentrations with Cardiometabolic Risk Factors in Children at 10 y of Age. Journal of Nutrition, 2021, 151, 1628-1636.	2.9	9
124	Associations of circulating folate, vitamin B12 and homocysteine concentrations in early pregnancy and cord blood with epigenetic gestational age: the Generation R Study. Clinical Epigenetics, 2021, 13, 95.	4.1	9
125	Maternal Earlyâ€Pregnancy Glucose Concentrations and Liver Fat Among Schoolâ€Age Children. Hepatology, 2021, 74, 1902-1913.	<b>7.</b> 3	9
126	Genome-wide DNA methylation patterns associated with general psychopathology in children. Journal of Psychiatric Research, 2021, 140, 214-220.	3.1	8

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127	Epigenetic age acceleration and cardiovascular outcomes in school-age children: The Generation R Study. Clinical Epigenetics, 2021, 13, 205.	4.1	8
128	Maternal Mediterranean diet in pregnancy and newborn DNA methylation: a meta-analysis in the PACE Consortium. Epigenetics, 2022, 17, 1419-1431.	2.7	8
129	Timing- and Dose-Specific Associations of Prenatal Smoke Exposure With Newborn DNA Methylation. Nicotine and Tobacco Research, 2020, 22, 1917-1922.	2.6	7
130	Pro-inflammatory Diet Pictured in Children With Atopic Dermatitis or Food Allergy: Nutritional Data of the LiNA Cohort. Frontiers in Nutrition, 2022, 9, 868872.	3.7	7
131	Pulmonary function and diffusion capacity are associated with pulmonary arterial systolic pressure in the general population: The Rotterdam Study. Respiratory Medicine, 2017, 132, 50-55.	2.9	6
132	Vitamin B12, folate and homocysteine concentrations during pregnancy and early signs of atherosclerosis at school-age. Clinical Nutrition, 2021, 40, 5133-5140.	5.0	6
133	LonglTools: Dynamic longitudinal exposome trajectories in cardiovascular and metabolic noncommunicable diseases. Environmental Epidemiology, 2022, 6, e184.	3.0	6
134	Longitudinal associations of DNA methylation and sleep in children: a meta-analysis. Clinical Epigenetics, 2022, 14, .	4.1	6
135	Maternal fish consumption, fatty acid levels and angiogenic factors: The Generation R Study. Placenta, 2015, 36, 1178-1184.	1.5	5
136	Understanding the cumulative risk of maternal prenatal biopsychosocial factors on birth weight: a DynaHEALTH study on two birth cohorts. Journal of Epidemiology and Community Health, 2020, 74, jech-2019-213154.	3.7	5
137	Associations of Hair Cortisol Concentrations With Cardiometabolic Risk Factors in Childhood. Journal of Clinical Endocrinology and Metabolism, 2021, 106, e3400-e3413.	3.6	5
138	Phenotypic Consequences of the <i>GJD2</i> Risk Genotype in Myopia Development., 2021, 62, 16.		5
139	Genetic and clinical determinants of abdominal aortic diameter: genome-wide association studies, exome array data and Mendelian randomization study. Human Molecular Genetics, 2022, 31, 3566-3579.	2.9	5
140	Maternal iron status in early pregnancy and DNA methylation in offspring: an epigenome-wide meta-analysis. Clinical Epigenetics, 2022, 14, 59.	4.1	5
141	Maternal plasma fatty acid patterns in mid-pregnancy and offspring epigenetic gestational age at birth. Epigenetics, 2022, 17, 1562-1572.	2.7	5
142	The Influence of Known Genetic Variants on Subclinical Cardiovascular Outcomes in Childhood. Circulation: Cardiovascular Genetics, 2015, 8, 596-602.	5.1	4
143	Gene Set Enrichment Analyses: lessons learned from the heart failure phenotype. BioData Mining, 2017, 10, 18.	4.0	4
144	Epigenome-wide associations between observed maternal sensitivity and offspring DNA methylation: a population-based prospective study in children. Psychological Medicine, 2022, 52, 2481-2491.	4.5	4

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145	Infant weight growth patterns, childhood BMI, and arterial health at age 10 years. Obesity, 2022, 30, 770-778.	3.0	4
146	Impact of maternal smoking during pregnancy on microvasculature in childhood. The Generation R Study. Early Human Development, 2015, 91, 607-611.	1.8	3
147	Maternal haemoglobin levels in pregnancy and child DNA methylation: a study in the pregnancy and childhood epigenetics consortium. Epigenetics, 2022, 17, 19-31.	2.7	3
148	Body fat, pericardial fat, liver fat and arterial health at age 10 years. Pediatric Obesity, 2022, 17, e12926.	2.8	3
149	Influence of genetic variants for birth weight on fetal growth and placental haemodynamics. Archives of Disease in Childhood: Fetal and Neonatal Edition, 2020, 105, 393-398.	2.8	2
150	Neonatal DNA methylation and childhood low prosocial behavior: An epigenomeâ€wide association metaâ€analysis. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2021, 186, 228-241.	1.7	2
151	Associations of maternal and infant metabolite profiles with foetal growth and the odds of adverse birth outcomes. Pediatric Obesity, 2021, , e12844.	2.8	2
152	Genetics of early-life head circumference and genetic correlations with neurological, psychiatric and cognitive outcomes. BMC Medical Genomics, 2022, $15$ , .	1.5	2
153	Influence of common genetic variants on childhood kidney outcomes. Pediatric Research, 2016, 80, 60-66.	2.3	1
154	Epigenome-wide contributions to individual differences in childhood phenotypes: a GREML approach. Clinical Epigenetics, 2022, 14, 53.	4.1	1
155	Title is missing!. , 2019, 16, e1002972.		0
156	Title is missing!. , 2019, 16, e1002972.		0
157	Title is missing!. , 2019, 16, e1002972.		0