Tsegaselassie Workalemahu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9370267/publications.pdf

Version: 2024-02-01

46 papers

14,783 citations

304743 22 h-index 206112 48 g-index

53 all docs 53 docs citations

53 times ranked

22817 citing authors

| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Genetic studies of body mass index yield new insights for obesity biology. Nature, 2015, 518, 197-206. | 27.8 | 3,823 |
| 2 | Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. Nature Genetics, 2010, 42, 937-948. | 21.4 | 2,634 |
| 3 | 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 |
| 4 | Hundreds of variants clustered in genomic loci and biological pathways affect human height. Nature, 2010, 467, 832-838. | 27.8 | 1,789 |
| 5 | New genetic loci link adipose and insulin biology to body fat distribution. Nature, 2015, 518, 187-196. | 27.8 | 1,328 |
| 6 | Meta-analysis identifies 13 new loci associated with waist-hip ratio and reveals sexual dimorphism in the genetic basis of fat distribution. Nature Genetics, 2010, 42, 949-960. | 21.4 | 836 |
| 7 | Genome-wide meta-analysis identifies 11 new loci for anthropometric traits and provides insights into genetic architecture. Nature Genetics, 2013, 45, 501-512. | 21.4 | 578 |
| 8 | Quality control and conduct of genome-wide association meta-analyses. Nature Protocols, 2014, 9, 1192-1212. | 12.0 | 398 |
| 9 | Sex-stratified Genome-wide Association Studies Including 270,000 Individuals Show Sexual Dimorphism in Genetic Loci for Anthropometric Traits. PLoS Genetics, 2013, 9, e1003500. | 3.5 | 371 |
| 10 | Novel locus including FGF21 is associated with dietary macronutrient intake. Human Molecular Genetics, 2013, 22, 1895-1902. | 2.9 | 167 |
| 11 | Genome-wide physical activity interactions in adiposity $\hat{a} \in A$ meta-analysis of 200,452 adults. PLoS Genetics, 2017, 13, e1006528. | 3.5 | 158 |
| 12 | Genome-wide analysis of BMI in adolescents and young adults reveals additional insight into the effects of genetic loci over the life course. Human Molecular Genetics, 2013, 22, 3597-3607. | 2.9 | 116 |
| 13 | Genomics of body fat percentage may contribute to sex bias in anorexia nervosa. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 428-438. | 1.7 | 87 |
| 14 | Genetic variants, plasma lipoprotein(a) levels, and risk of cardiovascular morbidity and mortality among two prospective cohorts of type 2 diabetes. European Heart Journal, 2012, 33, 325-334. | 2.2 | 81 |
| 15 | Meta-analysis of Gene-Level Associations for Rare Variants Based on Single-Variant Statistics. American Journal of Human Genetics, 2013, 93, 236-248. | 6.2 | 60 |
| 16 | Genetic correlations of psychiatric traits with body composition and glycemic traits are sex- and age-dependent. Nature Communications, 2019, 10, 5765. | 12.8 | 59 |
| 17 | Sex differences in the associations of placental epigenetic aging with fetal growth. Aging, 2019, 11, 5412-5432. | 3.1 | 44 |
| 18 | Genetic and Environmental Influences on Fetal Growth Vary during Sensitive Periods in Pregnancy. Scientific Reports, 2018, 8, 7274. | 3.3 | 38 |

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|----|--|------|-----------|
| 19 | Placental DNA methylation changes associated with maternal prepregnancy BMI and gestational weight gain. International Journal of Obesity, 2020, 44, 1406-1416. | 3.4 | 31 |
| 20 | Maternal dyslipidemia during early pregnancy and epigenetic ageing of the placenta. Epigenetics, 2019, 14, 1030-1039. | 2.7 | 30 |
| 21 | DNA methylation loci in placenta associated with birthweight and expression of genes relevant for early development and adult diseases. Clinical Epigenetics, 2020, 12, 78. | 4.1 | 28 |
| 22 | Birth Weight, Genetic Susceptibility, and Adulthood Risk of Type 2 Diabetes. Diabetes Care, 2012, 35, 2479-2484. | 8.6 | 24 |
| 23 | Physical Activity and Metabolic Syndrome among Ethiopian Adults. American Journal of Hypertension, 2013, 26, 535-540. | 2.0 | 21 |
| 24 | Differential DNA Methylation in Placenta Associated With Maternal Blood Pressure During Pregnancy. Hypertension, 2020, 75, 1117-1124. | 2.7 | 20 |
| 25 | Genetic Determinants for Body Iron Store and Type 2 Diabetes Risk in US Men and Women. PLoS ONE, 2012, 7, e40919. | 2.5 | 19 |
| 26 | Placental genetic variations in vitamin D metabolism and birthweight. Placenta, 2017, 50, 78-83. | 1.5 | 17 |
| 27 | Shared genetic underpinnings of childhood obesity and adult cardiometabolic diseases. Human Genomics, 2019, 13, 17. | 2.9 | 17 |
| 28 | Genetic variations related to maternal whole blood mitochondrial DNA copy number: a genome-wide and candidate gene study. Journal of Maternal-Fetal and Neonatal Medicine, 2017, 30, 2433-2439. | 1.5 | 15 |
| 29 | Genetic variations and risk of placental abruption: A genome-wide association study and meta-analysis of genome-wide association studies. Placenta, 2018, 66, 8-16. | 1.5 | 15 |
| 30 | Abruptio placentae risk and genetic variations in mitochondrial biogenesis and oxidative phosphorylation: replication of a candidate gene association study. American Journal of Obstetrics and Gynecology, 2018, 219, 617.e1-617.e17. | 1.3 | 15 |
| 31 | Trans-ethnic meta-analysis of genome-wide association studies identifies maternal ITPR1 as a novel locus influencing fetal growth during sensitive periods in pregnancy. PLoS Genetics, 2020, 16, e1008747. | 3.5 | 13 |
| 32 | Maternal cardiometabolic factors and genetic ancestry influence epigenetic aging of the placenta. Journal of Developmental Origins of Health and Disease, 2021, 12, 34-41. | 1.4 | 13 |
| 33 | Placental multi-omics integration identifies candidate functional genes for birthweight. Nature Communications, 2022, 13, 2384. | 12.8 | 13 |
| 34 | High burden of birthweight-lowering genetic variants in Africans and Asians. BMC Medicine, 2018, 16, 70. | 5.5 | 12 |
| 35 | Early pregnancy dyslipidemia is associated with placental DNA methylation at loci relevant for cardiometabolic diseases. Epigenomics, 2020, 12, 921-934. | 2.1 | 12 |
| 36 | Genome-wide and candidate gene association studies of placental abruption. International Journal of Molecular Epidemiology and Genetics, 2013, 4, 128-39. | 0.4 | 11 |

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|----|--|-----|-----------|
| 37 | Race–ethnic differences in the associations of maternal lipid trait genetic risk scores with longitudinal fetal growth. Journal of Clinical Lipidology, 2019, 13, 821-831. | 1.5 | 8 |
| 38 | Placental telomere length and risk of placental abruption. Journal of Maternal-Fetal and Neonatal Medicine, 2016, 29, 2767-2772. | 1.5 | 6 |
| 39 | Influence of Fetal and Maternal Genetic Susceptibility to Obesity on Birthweight in African Ancestry Populations. Frontiers in Genetics, 2018, 9, 511. | 2.3 | 6 |
| 40 | Maternal and Offspring Genetic Risk of Type 2 Diabetes and Offspring Birthweight Among African Ancestry Populations. Journal of Clinical Endocrinology and Metabolism, 2019, 104, 5032-5042. | 3.6 | 5 |
| 41 | Maternal BMIâ€Increasing Genetic Risk Score and Fetal Weights among Diverse US Ethnic Groups. Obesity, 2019, 27, 1150-1160. | 3.0 | 5 |
| 42 | Genetic overlap between birthweight and adult cardiometabolic diseases has implications for genomic medicine. Scientific Reports, 2019, 9, 4076. | 3.3 | 5 |
| 43 | Admixture mapping identifies African and Amerindigenous local ancestry loci associated with fetal growth. Human Genetics, 2021, 140, 985-997. | 3.8 | 5 |
| 44 | Associations of perinatal exposure to PM2.5 with gestational weight gain and offspring birth weight. Environmental Research, 2022, 204, 112087. | 7.5 | 4 |
| 45 | Maternal-fetal genetic interactions, imprinting, and risk of placental abruption. Journal of Maternal-Fetal and Neonatal Medicine, 2022, 35, 3473-3482. | 1.5 | 3 |
| 46 | Associations of maternal blood pressure-raising polygenic risk scores with fetal weight. Journal of Human Hypertension, 2021, , . | 2.2 | 3 |