# Audrey E Hendricks

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/5818437/publications.pdf

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|---|---|---|---|
| 47 papers | 3,245 citations | 331538 21 h-index | 233338 45 g-index |
| 50 all docs | 50 docs citations | 50 times ranked | 9249 citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | Genome-wide analysis of copy number variants and normal facial variation in a large cohort of Bantu Africans. Human Genetics and Genomics Advances, 2022, 3, 100082. | 1.0 | 1 |
| 2 | RAREsim: A simulation method for very rare genetic variants. American Journal of Human Genetics, 2022, 109, 680-691. | 2.6 | 1 |
| 3 | Opportunities and challenges for the use of common controls in sequencing studies. Nature Reviews Genetics, 2022, 23, 665-679. | 7.7 | 13 |
| 4 | Lipidomics-Based Comparison of Molecular Compositions of Green, Yellow, and Red Bell Peppers. Metabolites, 2021, 11, 241. | 1.3 | 13 |
| 5 | Improved first trimester maternal iodine status with preconception supplementation: The Women First Trial. Maternal and Child Nutrition, 2021, 17, e13204. | 1.4 | 7 |
| 6 | Genome-wide copy number variations in a large cohort of bantu African children. BMC Medical Genomics, 2021, 14, 129. | 0.7 | 6 |
| 7 | Summix: A method for detecting and adjusting for population structure in genetic summary data. American Journal of Human Genetics, 2021, 108, 1270-1282. | 2.6 | 5 |
| 8 | Different Blood Metabolomics Profiles in Infants Consuming a Meat- or Dairy-Based Complementary Diet. Nutrients, 2021, 13, 388. | 1.7 | 3 |
| 9 | Effects of Complementary Feeding With Different Protein-Rich Foods on Infant Growth and Gut Health: Study Protocol. Frontiers in Pediatrics, 2021, 9, 793215. | 0.9 | 4 |
| 10 | High-throughput analysis suggests differences in journal false discovery rate by subject area and impact factor but not open access status. BMC Bioinformatics, 2020, 21, 564. | 1.2 | 0 |
| 11 | Exome Sequencing Identifies Genes and Gene Sets Contributing to Severe Childhood Obesity, Linking PHIP Variants to Repressed POMC Transcription. Cell Metabolism, 2020, 31, 1107-1119.e12. | 7.2 | 38 |
| 12 | Longitudinal Changes of One-Carbon Metabolites and Amino Acid Concentrations during Pregnancy in the Women First Maternal Nutrition Trial. Current Developments in Nutrition, 2020, 4, nzz132. | 0.1 | 6 |
| 13 | Nutrimetabolomics reveals food-specific compounds in urine of adults consuming a DASH-style diet. Scientific Reports, 2020, 10, 1157. | 1.6 | 18 |
| 14 | Different Gut Microbial Profiles in Sub-Saharan African and South Asian Women of Childbearing Age Are Primarily Associated With Dietary Intakes. Frontiers in Microbiology, 2019, 10, 1848. | 1.5 | 16 |
| 15 | Genetic architecture of human thinness compared to severe obesity. PLoS Genetics, 2019, 15, e1007603. | 1.5 | 98 |
| 16 | Steroid receptor coactivator-1 modulates the function of Pomc neurons and energy homeostasis. Nature Communications, 2019, 10, 1718. | 5.8 | 45 |
| 17 | Insight into genetic regulation of miRNA in mouse brain. BMC Genomics, 2019, 20, 849. | 1.2 | 4 |
| 18 | Different Growth Patterns Persist at 24 Months of Age in Formula-Fed Infants Randomized to Consume a Meat- or Dairy-Based Complementary Diet from 5 to 12 Months of Age. Journal of Pediatrics, 2019, 206, 78-82. | 0.9 | 11 |

| # | Article | IF | Citations |
|---|---|---|---|
| 19 | Human Semaphorin 3 Variants Link Melanocortin Circuit Development and Energy Balance. Cell, 2019, 176, 729-742.e18. | 13.5 | 80 |
| 20 | A meat- or dairy-based complementary diet leads to distinct growth patterns in formula-fed infants: a randomized controlled trial. American Journal of Clinical Nutrition, 2018, 107, 734-742. | 2.2 | 33 |
| 21 | ProxECAT: Proxy External Controls Association Test. A new case-control gene region association test using allele frequencies from public controls. PLoS Genetics, 2018, 14, e1007591. | 1.5 | 23 |
| 22 | Body mass index is negatively associated with telomere length: a collaborative cross-sectional meta-analysis of 87 observational studies. American Journal of Clinical Nutrition, 2018, 108, 453-475. | 2.2 | 137 |
| 23 | Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. Nature Genetics, 2018, 50, 26-41. | 9.4 | 286 |
| 24 | Whole-Genome Sequencing Coupled to Imputation Discovers Genetic Signals for Anthropometric Traits. American Journal of Human Genetics, 2017, 100, 865-884. | 2.6 | 131 |
| 25 | Disruption of the homeodomain transcription factor orthopedia homeobox ( Otp ) is associated with obesity and anxiety. Molecular Metabolism, 2017, 6, 1419-1428. | 3.0 | 15 |
| 26 | Rare Variant Analysis of Human and Rodent Obesity Genes in Individuals with Severe Childhood Obesity. Scientific Reports, 2017, 7, 4394. | 1.6 | 50 |
| 27 | Iron in Micronutrient Powder Promotes an Unfavorable Gut Microbiota in Kenyan Infants. Nutrients, 2017, 9, 776. | 1.7 | 65 |
| 28 | Gene-gene Interaction Analyses for Atrial Fibrillation. Scientific Reports, 2016, 6, 35371. | 1.6 | 15 |
| 29 | Whole-exome sequencing in an isolated population from the Dalmatian island of Vis. European Journal of Human Genetics, 2016, 24, 1479-1487. | 1.4 | 11 |
| 30 | The UK10K project identifies rare variants in health and disease. Nature, 2015, 526, 82-90. | 13.7 | 1,014 |
| 31 | Genetic associations with expression for genes implicated in GWAS studies for atherosclerotic cardiovascular disease and blood phenotypes. Human Molecular Genetics, 2014, 23, 782-795. | 1.4 | 49 |
| 32 | Correction for multiple testing in a gene region. European Journal of Human Genetics, 2014, 22, 414-418. | 1.4 | 39 |
| 33 | KSR2 Mutations Are Associated with Obesity, Insulin Resistance, and Impaired Cellular Fuel Oxidation. Cell, 2013, 155, 765-777. | 13.5 | 154 |
| 34 | A Comparison of Gene Region Simulation Methods. PLoS ONE, 2012, 7, e40925. | 1.1 | 5 |
| 35 | Assessment of cortical and striatal involvement in 523 Huntington disease brains. Neurology, 2012, 79, 1708-1715. | 1.5 | 52 |
| 36 | TAA repeat variation in the GRIK2 gene does not influence age at onset in Huntingtonâ€™s disease. Biochemical and Biophysical Research Communications, 2012, 424, 404-408. | 1.0 | 20 |

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|---|---|---|---|
| 37 | Postmortem Interval InfluencesÎ±-Synuclein Expression in Parkinson Disease Brain. Parkinson's Disease, 2012, 2012, 1-8. | 0.6 | 11 |
| 38 | Comparison of statistical approaches to rare variant analysis for quantitative traits. BMC Proceedings, 2011, 5, S113. | 1.8 | 9 |
| 39 | Genomewide linkage study of modifiers of LRRK2â€related Parkinson's disease. Movement Disorders, 2011, 26, 2039-2044. | 2.2 | 8 |
| 40 | Genome-wide association and linkage analysis of quantitative traits: comparison of likelihood-ratio test and conditional score statistic. BMC Proceedings, 2009, 3, S100. | 1.8 | 3 |
| 41 | Somatic expansion of the Huntington's disease CAG repeat in the brain is associated with an earlier age of disease onset. Human Molecular Genetics, 2009, 18, 3039-3047. | 1.4 | 255 |
| 42 | Intergenerational and striatal CAG repeat instability in Huntington's disease knock-in mice involve different DNA repair genes. Neurobiology of Disease, 2009, 33, 37-47. | 2.1 | 189 |
| 43 | Estimating the probability of de novo HD cases from transmissions of expanded penetrant CAG alleles in the Huntington disease gene from male carriers of high normal alleles (27â€“35 CAG). American Journal of Medical Genetics, Part A, 2009, 149A, 1375-1381. | 0.7 | 48 |
| 44 | Replication of association between ELAVL4 and Parkinson disease: the GenePD study. Human Genetics, 2008, 124, 95-99. | 1.8 | 34 |
| 45 | The Gly2019Ser mutation in LRRK2is not fully penetrant in familial Parkinson's disease: the GenePD study. BMC Medicine, 2008, 6, 32. | 2.3 | 102 |
| 46 | Haplotypes and gene expression implicate the MAPT region for Parkinson disease. Neurology, 2008, 71, 28-34. | 1.5 | 103 |
| 47 | Beyond Aggravating and Mitigating Factors: The Analysis of Colorado's Death Penalty Cases (1999-2010):. Justice Evaluation Journal, 0, , 1-35. | 0.7 | 0 |