Lindsay Fernandez-Rhodes

List of Publications by Year
in descending order

44 papers	2,638 citations	393982 19 h-index	301761 39 g-index
47 all docs	47 docs citations	47 times ranked	6249 citing authors

#	Article	IF	Citations
1	Genetic analyses of diverse populations improves discovery for complex traits. Nature, 2019, 570, 514-518.	13.7	679
2	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. Nature Genetics, 2017, 49, 834-841.	9.4	426
3	Genetic Diversity and Association Studies in US Hispanic/Latino Populations: Applications in the Hispanic Community Health Study/Study of Latinos. American Journal of Human Genetics, 2016, 98, 165-184.	2.6	266
4	Clinical features of spinal and bulbar muscular atrophy. Brain, 2009, 132, 3242-3251.	3.7	223
5	Efficacy and safety of dutasteride in patients with spinal and bulbar muscular atrophy: a randomised placebo-controlled trial. Lancet Neurology, The, 2011, 10, 140-147.	4.9	138
6	Fine Mapping and Identification of BMI Loci in African Americans. American Journal of Human Genetics, 2013, 93, 661-671.	2.6	77
7	A trans-ancestral meta-analysis of genome-wide association studies reveals loci associated with childhood obesity. Human Molecular Genetics, 2019, 28, 3327-3338.	1.4	76
8	Replication of loci influencing ages at menarche and menopause in Hispanic women: the Women's Health Initiative SHARe Study. Human Molecular Genetics, 2012, 21, 1419-1432.	1.4	71
9	Genetic Association Analysis under Complex Survey Sampling: The Hispanic Community Health Study/Study of Latinos. American Journal of Human Genetics, 2014, 95, 675-688.	2.6	70
10	Genetic risk factors for BMI and obesity in an ethnically diverse population: Results from the population architecture using genomics and epidemiology (PAGE) study. Obesity, 2013, 21, 835-846.	1.5	68
11	Replication of genetic loci for ages at menarche and menopause in the multi-ethnic Population Architecture using Genomics and Epidemiology (PAGE) study. Human Reproduction, 2013, 28, 1695-1706.	0.4	64
12	Age at menarche and risk of type 2 diabetes among African-American and white women in the Atherosclerosis Risk in Communities (ARIC) study. Diabetologia, 2012, 55, 2371-2380.	2.9	61
13	Association of Adiposity Genetic Variants With Menarche Timing in 92,105 Women of European Descent. American Journal of Epidemiology, 2013, 178, 451-460.	1.6	51
14	Genetic Variation and Reproductive Timing: African American Women from the Population Architecture Using Genomics and Epidemiology (PAGE) Study. PLoS ONE, 2013, 8, e55258.	1.1	39
15	Trans-ethnic fine-mapping of genetic loci for body mass index in the diverse ancestral populations of the Population Architecture using Genomics and Epidemiology (PAGE) Study reveals evidence for multiple signals at established loci. Human Genetics, 2017, 136, 771-800.	1.8	31
16	Assessing Function and Endurance in Adults with Spinal and Bulbar Muscular Atrophy: Validity of the Adult Myopathy Assessment Tool. Rehabilitation Research and Practice, 2014, 2014, 1-16.	0.5	28
17	Genetics of Obesity in Diverse Populations. Current Diabetes Reports, 2018, 18, 145.	1.7	27
18	The genetic underpinnings of variation in ages at menarche and natural menopause among women from the multi-ethnic Population Architecture using Genomics and Epidemiology (PAGE) Study: A trans-ethnic meta-analysis. PLoS ONE, 2018, 13, e0200486.	1.1	25

#	Article	IF	Citations
19	Generalization of adiposity genetic loci to US Hispanic women. Nutrition and Diabetes, 2013, 3, e85-e85.	1.5	24
20	Transethnic insight into the genetics of glycaemic traits: fine-mapping results from the Population Architecture using Genomics and Epidemiology (PAGE) consortium. Diabetologia, 2017, 60, 2384-2398.	2.9	20
21	Importance of Genetic Studies of Cardiometabolic Disease in Diverse Populations. Circulation Research, 2020, 126, 1816-1840.	2.0	19
22	Reproductive aging-associated common genetic variants and the risk of breast cancer. Breast Cancer Research, 2012, 14, R54.	2.2	17
23	Generalization and fine mapping of European ancestry-based central adiposity variants in African ancestry populations. International Journal of Obesity, 2017, 41, 324-331.	1.6	16
24	Adverse childhood experiences and lifetime adverse maternal outcomes (gestational diabetes and) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 of Epidemiology, 2020, 50, 1-6.	0.9	14
25	Accuracy of Self-reported Weight in Hispanic/Latino Adults of the Hispanic Community Health Study/Study of Latinos. Epidemiology, 2017, 28, 847-853.	1.2	13
26	Trans-ethnic analysis of metabochip data identifies two new loci associated with BMI. International Journal of Obesity, 2018, 42, 384-390.	1.6	13
27	Genome-wide association of trajectories of systolic blood pressure change. BMC Proceedings, 2016, 10, 321-327.	1.8	8
28	Demographic and sociocultural risk factors for adulthood weight gain in Hispanic/Latinos: results from the Hispanic Community Health Study / Study of Latinos (HCHS/SOL). BMC Public Health, 2021, 21, 2064.	1.2	8
29	Complex patterns of direct and indirect association between the transcription Factor-7 like 2 gene, body mass index and type 2 diabetes diagnosis in adulthood in the Hispanic Community Health Study/Study of Latinos. BMC Obesity, 2018, 5, 26.	3.1	6
30	Methylome-wide association study of central adiposity implicates genes involved in immune and endocrine systems. Epigenomics, 2020, 12, 1483-1499.	1.0	6
31	Genetic Risk Factors for BMI and Obesity in an Ethnically Diverse Population: Results From the Population Architecture Using Genomics and Epidemiology (PAGE) Study. Obesity, 0, , .	1.5	6
32	Findings from the Hispanic Community Health Study/Study of Latinos on the Importance of Sociocultural Environmental Interactors: Polygenic Risk Score-by-Immigration and Dietary Interactions. Frontiers in Genetics, 2021, 12, 720750.	1.1	6
33	Direct and indirect genetic effects on triglycerides through omics and correlated phenotypes. BMC Proceedings, 2018, 12, 22.	1.8	5
34	Association Between Immigration History and Inflammatory Marker Profiles Among Older Adult Mexican Americans. Biodemography and Social Biology, 2018, 64, 30-42.	0.4	5
35	Comparison of 2 models for gene–environment interactions: an example of simulated gene–medication interactions on systolic blood pressure in family-based data. BMC Proceedings, 2016, 10, 371-377.	1.8	3
36	Characterization of the contribution of shared environmental and genetic factors to metabolic syndrome methylation heritability and familial correlations. BMC Genetics, 2018, 19, 69.	2.7	3

#	Article	IF	Citations
37	Spanish Language Use Across Generations and Depressive Symptoms Among US Latinos. Child Psychiatry and Human Development, 2019, 50, 61-71.	1.1	3
38	Intergenerational Educational Attainment and Cardiometabolic Health in Latino Individuals Living in the United States. Obesity, 2021, 29, 1178-1185.	1.5	3
39	Ancestral diversity improves discovery and fine-mapping of genetic loci for anthropometric traitsâ€"The Hispanic/Latino Anthropometry Consortium. Human Genetics and Genomics Advances, 2022, 3, 100099.	1.0	3
40	Local-Level Immigration Enforcement and Risk of Pediatric Hospitalization for Ambulatory Care Sensitive Conditions. Journal of Immigrant and Minority Health, 2021, , 1.	0.8	1
41	Obesity and climate adaptation. Evolution, Medicine and Public Health, 2019, 2019, 104-105.	1.1	0
42	Intergenerational educational mobility and type 2 diabetes in the Sacramento Area Latino Study of Aging. Annals of Epidemiology, 2021, 65, 93-93.	0.9	0
43	Abstract 002: Adherence to Ideal Lifeâ€™s Simple 7 Metrics is Associated With Epigenetic Biomarkers of Aging in African Americans: The Atherosclerosis Risk in Communities (ARIC) Study. Circulation, 2019, 139, .	1.6	0
44	Abstract P271: Dna Methylation Measures of Aging in Midlife are Associated With Frailty Components in African American and European American Older Adults: The Atherosclerosis Risk in Communities (ARIC) Study. Circulation, 2020, 141, .	1.6	0