

Kristin L Young

List of Publications by Year in descending order

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Version: 2024-02-01

55
papers

3,809
citations

361413

20
h-index

155660

55
g-index

65
all docs

65
docs citations

65
times ranked

8684
citing authors

#	ARTICLE	IF	CITATIONS
1	Do adverse childhood experiences and genetic obesity risk interact in relation to body mass index in young adulthood? Findings from the National Longitudinal Study of Adolescent to Adult Health. <i>Pediatric Obesity</i> , 2022, 17, e12885.	2.8	4
2	Ancestral diversity improves discovery and fine-mapping of genetic loci for anthropometric traits. The Hispanic/Latino Anthropometry Consortium. <i>Human Genetics and Genomics Advances</i> , 2022, 3, 100099.	1.7	3
3	Predicted gene expression in ancestrally diverse populations leads to discovery of susceptibility loci for lifestyle and cardiometabolic traits. <i>American Journal of Human Genetics</i> , 2022, 109, 669-679.	6.2	5
4	Genetic variants in anti-Müllerian hormone-related genes and breast cancer risk: results from the AMBER consortium. <i>Breast Cancer Research and Treatment</i> , 2021, 185, 469-478.	2.5	1
5	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. <i>American Journal of Human Genetics</i> , 2021, 108, 564-582.	6.2	18
6	Genome-wide association study of body fat distribution traits in Hispanics/Latinos from the HCHS/SOL. <i>Human Molecular Genetics</i> , 2021, 30, 2190-2204.	2.9	8
7	Transcriptome-Wide Association Study of Blood Cell Traits in African Ancestry and Hispanic/Latino Populations. <i>Genes</i> , 2021, 12, 1049.	2.4	11
8	Sugar-Sweetened Beverage Consumption May Modify Associations Between Genetic Variants in the CHREBP (Carbohydrate Responsive Element Binding Protein) Locus and HDL-C (High-Density Lipoprotein) Tj ETQq0,0,0 rgBT /Overlock 1 e003288.	3.6	8
9	The power of genetic diversity in genome-wide association studies of lipids. <i>Nature</i> , 2021, 600, 675-679.	27.8	353
10	Genetic Studies of Leptin Concentrations Implicate Leptin in the Regulation of Early Adiposity. <i>Diabetes</i> , 2020, 69, 2806-2818.	0.6	26
11	Serum metabolites reflecting gut microbiome alpha diversity predict type 2 diabetes. <i>Gut Microbes</i> , 2020, 11, 1632-1642.	9.8	65
12	Importance of Genetic Studies of Cardiometabolic Disease in Diverse Populations. <i>Circulation Research</i> , 2020, 126, 1816-1840.	4.5	19
13	Open Chromatin Profiling in Adipose Tissue Marks Genomic Regions with Functional Roles in Cardiometabolic Traits. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 2521-2533.	1.8	19
14	Genetic analyses of diverse populations improves discovery for complex traits. <i>Nature</i> , 2019, 570, 514-518.	27.8	679
15	Exome-Derived Adiponectin-Associated Variants Implicate Obesity and Lipid Biology. <i>American Journal of Human Genetics</i> , 2019, 105, 15-28.	6.2	21
16	Multi-ancestry genome-wide gene-smoking interaction study of 387,272 individuals identifies new loci associated with serum lipids. <i>Nature Genetics</i> , 2019, 51, 636-648.	21.4	112
17	Protein-coding variants implicate novel genes related to lipid homeostasis contributing to body-fat distribution. <i>Nature Genetics</i> , 2019, 51, 452-469.	21.4	89
18	Associations of Mitochondrial and Nuclear Mitochondrial Variants and Genes with Seven Metabolic Traits. <i>American Journal of Human Genetics</i> , 2019, 104, 112-138.	6.2	106

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19	Sugar-sweetened beverage intake associations with fasting glucose and insulin concentrations are not modified by selected genetic variants in a ChREBP-FGF21 pathway: a meta-analysis. <i>Diabetologia</i> , 2018, 61, 317-330.	6.3	32
20	Genome-Wide Interactions with Dairy Intake for Body Mass Index in Adults of European Descent. <i>Molecular Nutrition and Food Research</i> , 2018, 62, 1700347.	3.3	9
21	Genetics of Obesity in Diverse Populations. <i>Current Diabetes Reports</i> , 2018, 18, 145.	4.2	27
22	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. <i>BMC Obesity</i> , 2018, 5, 26.	3.1	6
23	Characterization of the contribution of shared environmental and genetic factors to metabolic syndrome methylation heritability and familial correlations. <i>BMC Genetics</i> , 2018, 19, 69.	2.7	3
24	A survey of microRNA single nucleotide polymorphisms identifies novel breast cancer susceptibility loci in a case-control, population-based study of African-American women. <i>Breast Cancer Research</i> , 2018, 20, 45.	5.0	15
25	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. <i>Nature Genetics</i> , 2018, 50, 26-41.	21.4	286
26	Rare and low-frequency coding variants alter human adult height. <i>Nature</i> , 2017, 542, 186-190.	27.8	544
27	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. <i>Nature Communications</i> , 2017, 8, 14977.	12.8	169
28	Fifteen new risk loci for coronary artery disease highlight arterial-wall-specific mechanisms. <i>Nature Genetics</i> , 2017, 49, 1113-1119.	21.4	260
29	BMI loci and longitudinal BMI from adolescence to young adulthood in an ethnically diverse cohort. <i>International Journal of Obesity</i> , 2017, 41, 759-768.	3.4	23
30	Genetic identification of a common collagen disease in Puerto Ricans via identity-by-descent mapping in a health system. <i>ELife</i> , 2017, 6, .	6.0	65
31	Genome-wide physical activity interactions in adiposity â€• A meta-analysis of 200,452 adults. <i>PLoS Genetics</i> , 2017, 13, e1006528.	3.5	158
32	Discovery and fine-mapping of adiposity loci using high density imputation of genome-wide association studies in individuals of African ancestry: African Ancestry Anthropometry Genetics Consortium. <i>PLoS Genetics</i> , 2017, 13, e1006719.	3.5	98
33	The interaction between physical activity and obesity gene variants in association with BMI: Does the obesogenic environment matter?. <i>Health and Place</i> , 2016, 42, 159-165.	3.3	10
34	Evidence for Association between <i>SH2B1</i> Gene Variants and Glycated Hemoglobin in Nondiabetic European American Young Adults: The Add Health Study. <i>Annals of Human Genetics</i> , 2016, 80, 294-305.	0.8	3
35	Genome-wide association of trajectories of systolic blood pressure change. <i>BMC Proceedings</i> , 2016, 10, 321-327.	1.6	8
36	Comparison of 2 models for gene-environment interactions: an example of simulated gene-medication interactions on systolic blood pressure in family-based data. <i>BMC Proceedings</i> , 2016, 10, 371-377.	1.6	3

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37	Genetic Diversity and Association Studies in US Hispanic/Latino Populations: Applications in the Hispanic Community Health Study/Study of Latinos. <i>American Journal of Human Genetics</i> , 2016, 98, 165-184.	6.2	266
38	Influence of <sc>SNP</sc>*<sc>SNP</sc> interaction on <sc>BMI</sc> in <sc>E</sc>uropean <sc>A</sc>merican adolescents: findings from the <sc>N</sc>ational <sc>L</sc>ongitudinal <sc>S</sc>tudy of <sc>A</sc>dolescent <sc>H</sc>ealth. <i>Pediatric Obesity</i> , 2016, 11, 95-101.	2.8	10
39	Interaction of smoking and obesity susceptibility loci on adolescent BMI: The National Longitudinal Study of Adolescent to Adult Health. <i>BMC Genetics</i> , 2015, 16, 131.	2.7	10
40	Decisional stage distribution for colorectal cancer screening among diverse, low-income study participants. <i>Health Education Research</i> , 2015, 30, 400-411.	1.9	14
41	Sequence Variation in <i>TMEM18</i> in Association With Body Mass Index. <i>Circulation: Cardiovascular Genetics</i> , 2014, 7, 344-349.	5.1	8
42	Lipoprotein lipase variants interact with polyunsaturated fatty acids for obesity traits in women: Replication in two populations. <i>Nutrition, Metabolism and Cardiovascular Diseases</i> , 2014, 24, 1323-1329.	2.6	10
43	Abstract 21: Accounting For Smoking Behavior In Genome-wide Analysis Of Obesity Phenotypes: The Giant (genetic Investigation Of Anthropometric Traits) Consortium. <i>Circulation</i> , 2014, 129, .	1.6	0
44	Abstract P157: Does Physical Activity Modify the Association of 15 Well-established Obesity Loci with BMI: The ARIC Study. <i>Circulation</i> , 2013, 127, .	1.6	0
45	The interaction between physical activity and obesity gene variants in association with BMI: Does the obesogenic environment matter?. <i>FASEB Journal</i> , 2013, 27, 236.5.	0.5	0
46	Estimation of genetic effects on BMI during adolescence in an ethnically diverse cohort: The National Longitudinal Study of Adolescent Health. <i>Nutrition and Diabetes</i> , 2012, 2, e47-e47.	3.2	24
47	American Indian/Alaska Native Willingness to Provide Biological Samples for Research Purposes. <i>Journal of Community Health</i> , 2012, 37, 701-705.	3.8	14
48	Paternal Genetic History of the Basque Population of Spain. <i>Human Biology</i> , 2011, 83, 455-475.	0.2	12
49	Autosomal short tandem repeat genetic variation of the Basques in Spain. <i>Croatian Medical Journal</i> , 2011, 52, 372-383.	0.7	21
50	Genetic Architecture of a Small, Recently Aggregated Aleut Population: Bering Island, Russia. <i>Human Biology</i> , 2010, 82, 719-736.	0.2	12
51	Postfamine stature and socioeconomic status in Ireland. <i>American Journal of Human Biology</i> , 2008, 20, 726-731.	1.6	11
52	Genetic structure of Algerian populations. <i>American Journal of Human Biology</i> , 2006, 18, 492-501.	1.6	13
53	HLA Genes in the Chuvashian Population from European Russia: Admixture of Central European and Mediterranean Populations. <i>Human Biology</i> , 2003, 75, 375-392.	0.2	47
54	Genetic Evidence for the Phylogenetic Relationship between Na-Dene and Yeniseian Speakers. <i>Human Biology</i> , 2002, 74, 743-760.	0.2	27

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55	Demic expansion or cultural diffusion: migration and Basque origins. , 0, , 224-249.		0