## Bertha A Hidalgo

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3619535/publications.pdf

Version: 2024-02-01

58 2,873 papers citations

201385 27 h-index 50 g-index

61 all docs 61 does citations

61 times ranked

6758 citing authors

#	Article	IF	Citations
1	Inherited causes of clonal haematopoiesis in 97,691 whole genomes. Nature, 2020, 586, 763-768.	13.7	376
2	Multivariate or Multivariable Regression?. American Journal of Public Health, 2013, 103, 39-40.	1.5	222
3	Use of >100,000 NHLBI Trans-Omics for Precision Medicine (TOPMed) Consortium whole genome sequences improves imputation quality and detection of rare variant associations in admixed African and Hispanic/Latino populations. PLoS Genetics, 2019, 15, e1008500.	1.5	203
4	Low-frequency and rare exome chip variants associate with fasting glucose and type 2 diabetes susceptibility. Nature Communications, 2015, 6, 5897.	5 <b>.</b> 8	173
5	Epigenome-wide study identifies novel methylation loci associated with body mass index and waist circumference. Obesity, 2015, 23, 1493-1501.	1.5	152
6	Epigenome-Wide Association Study of Fasting Measures of Glucose, Insulin, and HOMA-IR in the Genetics of Lipid Lowering Drugs and Diet Network Study. Diabetes, 2014, 63, 801-807.	0.3	149
7	Inhibition of Wnt/ $\hat{l}^2$ -catenin pathway by niclosamide: A therapeutic target for ovarian cancer. Gynecologic Oncology, 2014, 134, 112-120.	0.6	142
8	Effect of Niclosamide on Basal-like Breast Cancers. Molecular Cancer Therapeutics, 2014, 13, 800-811.	1.9	78
9	Longâ€Cycle Structured Intermittent versus Continuous Highly Active Antiretroviral Therapy for the Treatment of Chronic Infection with Human Immunodeficiency Virus: Effects on Drug Toxicity and on Immunologic and Virologic Parameters. Journal of Infectious Diseases, 2003, 188, 388-396.	1.9	77
10	Turning Chutes into Ladders for Women Faculty: A Review and Roadmap for Equity in Academia. Journal of Women's Health, 2020, 29, 721-733.	1.5	76
11	Development of a Theory-Based (PEN-3 and Health Belief Model), Culturally Relevant Intervention on Cervical Cancer Prevention Among Latina Immigrants Using Intervention Mapping. Health Promotion Practice, 2012, 13, 29-40.	0.9	75
12	Evaluation of Initial CD4+T Cell Counts in Individuals with Newly Diagnosed Human Immunodeficiency Virus Infection, by Sex and Race, in Urban Settings. Journal of Infectious Diseases, 2002, 185, 1818-1821.	1.9	70
13	Metabolic and inflammatory biomarkers are associated with epigenetic aging acceleration estimates in the GOLDN study. Clinical Epigenetics, 2018, 10, 56.	1.8	68
14	Niclosamide and its analogs are potent inhibitors of Wnt/ $\hat{l}^2$ -catenin, mTOR and STAT3 signaling in ovarian cancer. Oncotarget, 2016, 7, 86803-86815.	0.8	64
15	Dietary Interventions and Quality of Life: A Systematic Review of the Literature. Journal of Nutrition Education and Behavior, 2014, 46, 90-101.	0.3	63
16	An integrative cross-omics analysis of DNA methylation sites of glucose and insulin homeostasis. Nature Communications, 2019, 10, 2581.	5 <b>.</b> 8	62
17	Genetics of Type 2 Diabetes in U.S. Hispanic/Latino Individuals: Results From the Hispanic Community Health Study/Study of Latinos (HCHS/SOL). Diabetes, 2017, 66, 1419-1425.	0.3	60
18	Epigenomics and metabolomics reveal the mechanism of the APOA2-saturated fat intake interaction affecting obesity. American Journal of Clinical Nutrition, 2018, 108, 188-200.	2.2	54

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19	Association of DNA Methylation at CPT1A Locus with Metabolic Syndrome in the Genetics of Lipid Lowering Drugs and Diet Network (GOLDN) Study. PLoS ONE, 2016, 11, e0145789.	1.1	54
20	Epigenome-wide association study of metabolic syndrome in African-American adults. Clinical Epigenetics, 2018, 10, 49.	1.8	49
21	Carbohydrate and fat intake associated with risk of metabolic diseases through epigenetics of CPT1A. American Journal of Clinical Nutrition, 2020, 112, 1200-1211.	2.2	48
22	Impact of Rare and Common Genetic Variants on Diabetes Diagnosis by Hemoglobin A1c in Multi-Ancestry Cohorts: The Trans-Omics for Precision Medicine Program. American Journal of Human Genetics, 2019, 105, 706-718.	2.6	44
23	Genetic Characterization of Rebounding Human Immunodeficiency Virus Type 1 in Plasma during Multiple Interruptions of Highly Active Antiretroviral Therapy. Journal of Virology, 2003, 77, 3229-3237.	1.5	42
24	Epigenome-wide association study of triglyceride postprandial responses to a high-fat dietary challenge. Journal of Lipid Research, 2016, 57, 2200-2207.	2.0	40
25	Ideal cardiovascular health, glycaemic status and incident type 2 diabetes mellitus: the REasons for Geographic and Racial Differences in Stroke (REGARDS) study. Diabetologia, 2019, 62, 426-437.	2.9	39
26	Mendelian randomization supports bidirectional causality between telomere length and clonal hematopoiesis of indeterminate potential. Science Advances, 2022, 8, eabl6579.	4.7	36
27	Mapping hot spots of breast cancer mortality in the United States: place matters for Blacks and Hispanics. Cancer Causes and Control, 2018, 29, 737-750.	0.8	34
28	Genetic determinants of telomere length from 109,122 ancestrally diverse whole-genome sequences in TOPMed. Cell Genomics, 2022, 2, 100084.	3.0	29
29	The role of epidemiology in evidence-based policy making: a case study of tobacco use in youth. Annals of Epidemiology, 2015, 25, 360-365.	0.9	27
30	Lipid changes due to fenofibrate treatment are not associated with changes in DNA methylation patterns in the GOLDN study. Frontiers in Genetics, 2015, 6, 304.	1.1	20
31	Sex Differences in Blood HDLâ $\in$ c, the Total Cholesterol/HDLâ $\in$ c Ratio, and Palmitoleic Acid are Not Associated with Variants in Common Candidate Genes. Lipids, 2017, 52, 969-980.	0.7	19
32	Heritable DNA Methylation in CD4+ Cells among Complex Families Displays Genetic and Non-Genetic Effects. PLoS ONE, 2016, 11, e0165488.	1.1	19
33	Associations of the MCM6-rs3754686 proxy for milk intake in Mediterranean and American populations with cardiovascular biomarkers, disease and mortality: Mendelian randomization. Scientific Reports, 2016, 6, 33188.	1.6	18
34	An epigenome-wide association study of inflammatory response to fenofibrate in the Genetics of Lipid Lowering Drugs and Diet Network. Pharmacogenomics, 2017, 18, 1333-1341.	0.6	16
35	Addressing refugee health through evidence-based policies: aÂcaseÂstudy. Annals of Epidemiology, 2018, 28, 411-419.	0.9	16
36	Immunogenicity of Seven New Recombinant Yellow Fever Viruses 17D Expressing Fragments of SIVmac239 Gag, Nef, and Vif in Indian Rhesus Macaques. PLoS ONE, 2013, 8, e54434.	1.1	15

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37	Racial and socioeconomic disparity associates with differences in cardiac DNA methylation among men with end-stage heart failure. American Journal of Physiology - Heart and Circulatory Physiology, 2021, 320, H2066-H2079.	1.5	15
38	Preventive and Curative Care Utilization Among Mexican Immigrant Women in Birmingham, AL. Journal of Immigrant and Minority Health, 2012, 14, 983-989.	0.8	14
39	Interaction of an S100A9 gene variant with saturated fat and carbohydrates to modulate insulin resistance in 3 populations of different ancestries1–3. American Journal of Clinical Nutrition, 2016, 104, 508-517.	2.2	11
40	Genome- and CD4 + T-cell methylome-wide association study of circulating trimethylamine-N-oxide in the Genetics of Lipid Lowering Drugs and Diet Network (GOLDN). Journal of Nutrition & Intermediary Metabolism, 2017, 8, 1-7.	1.7	11
41	An exome-wide sequencing study of lipid response to high-fat meal and fenofibrate in Caucasians from the GOLDN cohort. Journal of Lipid Research, 2018, 59, 722-729.	2.0	10
42	Associations between SLC16A11 variants and diabetes in the Hispanic Community Health Study/Study of Latinos (HCHS/SOL). Scientific Reports, 2019, 9, 843.	1.6	9
43	Will Podcasting and Social Media Replace Journals and Traditional Science Communication? No, but American Journal of Epidemiology, 2021, 190, 1625-1631.	1.6	9
44	CPT1A: the future of heart disease detection and personalized medicine?. Clinical Lipidology, 2014, 9, 9-12.	0.4	8
45	Salivary AMY1 Copy Number Variation Modifies Age-Related Type 2 Diabetes Risk. Clinical Chemistry, 2020, 66, 718-726.	1.5	7
46	A 6-CpG validated methylation risk score model for metabolic syndrome: The HyperGEN and GOLDN studies. PLoS ONE, 2021, 16, e0259836.	1.1	7
47	Genetic Contributors of Incident Stroke in 10,700 African Americans With Hypertension: A Meta-Analysis From the Genetics of Hypertension Associated Treatments and Reasons for Geographic and Racial Differences in Stroke Studies. Frontiers in Genetics, 2021, 12, 781451.	1.1	7
48	Research Goes Red: Early Experience With a Participant-Centric Registry. Circulation Research, 2022, 130, 343-351.	2.0	6
49	Diagnostic accuracy of self-reported racial composition of residential neighborhood. Annals of Epidemiology, 2015, 25, 597-604.	0.9	5
50	Whole Genome Sequencing Identifies CRISPLD2 as a Lung Function Gene in Children With Asthma. Chest, 2019, 156, 1068-1079.	0.4	5
51	Linkage and association analysis of circulating vitamin D and parathyroid hormone identifies novel loci in Alaska Native Yup'ik people. Genes and Nutrition, 2016, 11, 23.	1.2	4
52	Validation of Self-Reported Measures in Health Disparities Research. Journal of Biometrics $\&$ Biostatistics, 2012, 03, .	4.0	4
53	A family-specific linkage analysis of blood lipid response to fenofibrate in the Genetics of Lipid Lowering Drug and Diet Network. Pharmacogenetics and Genomics, 2015, 25, 511-514.	0.7	3
54	Motherâ€child cardiometabolic health 4–10Âyears after pregnancy complicated by obesity with and without gestational diabetes. Obesity Science and Practice, 2022, 8, 627-640.	1.0	3

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55	Racial/ethnic distribution of graduates from doctorate and masters epidemiology degree programs in the United States, 2008 to 2018. Annals of Epidemiology, 2022, 68, 32-36.	0.9	3
56	An Exome-Wide Sequencing Study of the GOLDN Cohort Reveals Novel Associations of Coding Variants and Fasting Plasma Lipids. Frontiers in Genetics, 2019, 10, 158.	1.1	2
57	Behavior related genes, dietary preferences and anthropometric traits. FASEB Journal, 2017, 31, .	0.2	1
58	<b><i>APOL1</i></b> Risk Variants Associated with Serum Albumin in a Population-Based Cohort Study. American Journal of Nephrology, 2022, 53, 182-190.	1.4	0