

Linda M Polfus

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7718615/publications.pdf>

Version: 2024-02-01

22
papers

607
citations

686830

13
h-index

676716

22
g-index

23
all docs

23
docs citations

23
times ranked

2278
citing authors

#	ARTICLE	IF	CITATIONS
1	Platelet-Related Variants Identified by Exomechip Meta-analysis in 157,293 Individuals. American Journal of Human Genetics, 2016, 99, 40-55.	2.6	82
2	Exome Genotyping Identifies Pleiotropic Variants Associated with Red Blood Cell Traits. American Journal of Human Genetics, 2016, 99, 8-21.	2.6	60
3	Examining the Incidence of Human Papillomavirus-Associated Head and Neck Cancers by Race and Ethnicity in the U.S., 1995–2005. PLoS ONE, 2012, 7, e32657.	1.1	52
4	Genome-Wide Association Study of Gene by Smoking Interactions in Coronary Artery Calcification. PLoS ONE, 2013, 8, e74642.	1.1	51
5	Large-Scale Exome-wide Association Analysis Identifies Loci for White Blood Cell Traits and Pleiotropy with Immune-Mediated Diseases. American Journal of Human Genetics, 2016, 99, 22-39.	2.6	50
6	Analysis of loss-of-function variants and 20 risk factor phenotypes in 8,554 individuals identifies loci influencing chronic disease. Nature Genetics, 2015, 47, 640-642.	9.4	49
7	Whole-Exome Sequencing Identifies Loci Associated with Blood Cell Traits and Reveals a Role for Alternative GF11B Splice Variants in Human Hematopoiesis. American Journal of Human Genetics, 2016, 99, 481-488.	2.6	45
8	Coronary Heart Disease and Genetic Variants with Low Phospholipase A ₂ Activity. New England Journal of Medicine, 2015, 372, 295-296.	13.9	36
9	Genetic discovery and risk characterization in type 2 diabetes across diverse populations. Human Genetics and Genomics Advances, 2021, 2, 100029.	1.0	23
10	Genetic invalidation of Lp-PLA2 as a therapeutic target: Large-scale study of five functional Lp-PLA2-lowering alleles. European Journal of Preventive Cardiology, 2017, 24, 492-504.	0.8	22
11	Risk of breast cancer and prediagnostic urinary excretion of bisphenol A, triclosan and parabens: The Multiethnic Cohort Study. International Journal of Cancer, 2021, 149, 1426-1434.	2.3	21
12	Lipoprotein associated phospholipase A2 activity, apolipoprotein C3 loss-of-function variants and cardiovascular disease: The Atherosclerosis Risk In Communities Study. Atherosclerosis, 2015, 241, 641-648.	0.4	16
13	Whole genome sequence association with E-selectin levels reveals loss-of-function variant in African Americans. Human Molecular Genetics, 2019, 28, 515-523.	1.4	15
14	Multi-ethnic GWAS and fine-mapping of glycaemic traits identify novel loci in the PAGE Study. Diabetologia, 2022, 65, 477-489.	2.9	15
15	Whole-genome sequencing in diverse subjects identifies genetic correlates of leukocyte traits: The NHLBI TOPMed program. American Journal of Human Genetics, 2021, 108, 1836-1851.	2.6	14
16	Lipoprotein-associated phospholipase A2 and risk of incident peripheral arterial disease: Findings from The Atherosclerosis Risk in Communities study (ARIC). Atherosclerosis, 2018, 268, 12-18.	0.4	13
17	Association Between Absolute Neutrophil Count and Variation at <i>TCIRG1</i> : The NHLBI Exome Sequencing Project. Genetic Epidemiology, 2016, 40, 470-474.	0.6	11
18	Allelic Heterogeneity at the CRP Locus Identified by Whole-Genome Sequencing in Multi-ancestry Cohorts. American Journal of Human Genetics, 2020, 106, 112-120.	2.6	9

#	ARTICLE	IF	CITATIONS
19	Association of Exome Sequences With Cardiovascular Traits Among Blacks in the Jackson Heart Study. <i>Circulation: Cardiovascular Genetics</i> , 2016, 9, 368-374.	5.1	8
20	A common TCN1 loss-of-function variant is associated with lower vitamin B12 concentration in African Americans. <i>Blood</i> , 2018, 131, 2859-2863.	0.6	7
21	Genes Influencing the Development and Severity of Chronic ITP Identified through Whole Exome Sequencing. <i>Blood</i> , 2015, 126, 73-73.	0.6	6
22	Genome-wide association study of pancreatic fat: The Multiethnic Cohort Adiposity Phenotype Study. <i>PLoS ONE</i> , 2021, 16, e0249615.	1.1	2