## Linda M Polfus

## List of Publications by Year

 in descending orderSource: https:||exaly.com/author-pdf/7718615/publications.pdf
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5 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

Allelic Heterogeneity at the CRP Locus Identified by Whole-Genome Sequencing in Multi-ancestry
2.6

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## 6 Cohorts. American Journal of Human Genetics, 2020, 106, 112-120.

7 Whole genome sequence association with E-selectin levels reveals loss-of-function variant in African
7 Americans. Human Molecular Genetics, 2019, 28, 515-523.
1.4

15

8 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.
$0.4 \quad 13$
A common TCN1 loss-of-function variant is associated with lower vitamin B12 concentration in
9 African Americans. Blood, 2018, 131, 2859-2863.

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

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> 11 Association Between Absolute Neutrophil Count and Variation at <i>TCIRG1</i〉: The NHLBI Exome
> Sequencing Project. Genetic Epidemiology, 2016, 40, 470-474.
$0.6 \quad 11$

12 Exome Genotyping Identifies Pleiotropic Variants Associated with Red Blood Cell Traits. American Journal of Human Genetics, 2016, 99, 8-21.
2.6

60
Whole-Exome Sequencing Identifies Loci Associated with Blood Cell Traits and Reveals a Role for
13 Alternative GFIB Splice Variants in Human Hematopoiesis. American Journal of Human Genetics, 2016,
$99,481-488$.

14 Association of Exome Sequences With Cardiovascular Traits Among Blacks in the Jackson Heart Study. Circulation: Cardiovascular Genetics, 2016, 9, 368-374.
$5.1 \quad 8$
15 Platelet-Related Variants Identified by Exomechip Meta-analysis in 157,293 Individuals. American Journal
2.6 of Human Genetics, 2016, 99, 40-55.

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.

