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PERSONAL

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EDUCATION

- 2007 **Doctor of Philosophy**
Epidemiology, formal minor in biostatistics
University of North Carolina at Chapel Hill
Dissertation: Genotype-by-smoking interaction and the risk of
atherosclerosis and its clinical sequelae
Advisor: Dr. Kari North
- 2003 **Masters of Public Health**
Epidemiology, specialization in public health genetics
University of Michigan, Ann Arbor
- 1999 **Bachelor of Science**
Human Biology
Michigan State University

PROFESSIONAL EXPERIENCE

- 2023 – present Professor of Epidemiology
University of North Carolina at Chapel Hill
- 2017 – 2023 Associate Professor of Epidemiology
University of North Carolina at Chapel Hill
- 2016 – present Carolina Population Center Fellow
University of North Carolina at Chapel Hill
- 2011 - 2017 Assistant Professor of Epidemiology
University of North Carolina at Chapel Hill
- 2007 – 2011 Postdoctoral Fellow
Department of Epidemiology
University of North Carolina at Chapel Hill

2003 – 2007 Research Assistant
 Department of Epidemiology
 University of North Carolina at Chapel Hill

2001 – 2003 Research Assistant
 Department of Epidemiology
 University of Michigan, Ann Arbor

2000 – 2001 Research Assistant
 Department of Pharmacology
 University of Michigan, Ann Arbor

HONORS

2024 AHA Genomic/Precision Medicine & Epidemiology/Prevention Mid-Career Award
 2021 UNC Gillings School of Public Health Teaching Excellence and Innovation Award
 2016 Invited attendee, AHA/ASA Research Leaders Academy
 2015 Sandra A. Daugherty AHA Epidemiology/Lifestyle junior faculty award
 2013 CHARGE visiting fellow (Mentor: Dr. Bruce Psaty, University of Washington)
 2011 -2017 NIH / NHLBI Clinical Research Loan Repayment Program Award
 2010 University of North Carolina Postdoctoral Award for Research Excellence
 2008 Delta Omega, Honorary Society for Graduate Studies in Public Health
 2007 - 2009 University of North Carolina Cardiovascular Disease postdoctoral training grant
 2006 Rebecca James Baker Memorial Scholarship
 2005 North Carolina State University Institute in Statistical Genetics scholarship
 2003 - 2006 University of North Carolina Cardiovascular Disease predoctoral training grant
 2001 - 2003 University of Michigan Dean’s Scholarship
 1996,97,98-99 Dean’s List of Scholars, Michigan State University

BIBLIOGRAPHY

†First author is a pre- or post-doctoral fellow for whom I am the primary mentor.

**Co-first/senior authorship*

UNDER REVIEW

†Lee HH, **Avery CL**, Graff M, Kim D, Van Horn L, Kooperberg C, North KE. Genetic risk of body mass index and race modify effect of a low-fat dietary trial on 7-year weight change.

†Ballou AF, April-Sanders A, Meyer M, Rodrigues C, Stores-Alvarez D, Vanterpool YE, Vladitiu CJ, Welch B, Graff M, **Avery CL**. Associations between Lipoprotein(a) and Pregnancy Outcomes: The Hispanic Community Health Study/Study of Latinos

†Downie CG, Highland HM, Alotaibi M, Welch BM, Howard AG, Lilly A, Cheng S, Long T, Jain M, Thyagarajan B, Yu B, North KE, **Avery CL**. Genome-wide association study reveals shared and distinct genetic architecture underlying fatty acid and bioactive oxylipin metabolites

†Staley BS, Mhatre S, Sha W, Hullings AG, Krishnan M, Aiello AE, Meyer K, North KE, Kershaw KN, Sumner SCJ, Lewis CE, Gordon-Larsen P, **Avery CL**, Howard AG. Socioeconomic disadvantage is associated with metabolic pathways linked to increased chronic disease risk: the Coronary Artery Risk Development in Young Adults (CARDIA) Study

†Lee MP, Raffield L, Ballou AF, Downie CG, Kassahun H, Kent ST, Lopez AG, Monda KL, North KE, Rhodes SL, Yarosh R, Ballantyne CM, Graff M, **Avery CL**. Phenome-wide association study

Renson A, Keil AP, Dowd JB, **Avery CL**, Robinson WR, Zivich PN, Safford MM, Carson AP, Aiello AE. Effects of a hypothetical U.S. federal minimum wage increase on racial inequities in cardiovascular mortality: a novel difference-in-difference approach.

Shrestha P, Graff M, Gu Y, Wang Y, Avery CL, Ginnis J, Simancas-Pallares MA, Ferreira Zandoná AG, Ahn HS, Nguyen KN, Lin DY, Preisser JS, Slade GD, Marazita ML, North KE, Divaris K. Multi-ancestry Genome-Wide Association Study of Early Childhood Caries. medRxiv [Preprint]. 2024 Mar 18:2024.

PUBLISHED OR IN-PRESS (n= 7,432 citations, h-index = 39)

131. Li Y, Wong KY, Howard AG, Gordon-Larsen P, Highland HM, Graff M, North KE, Downie CG, **Avery CL**, Yu B, Young KL, Buchanan V, Kaplan R, Hou L, Thomas Joyce B, Qi Q, Sofer T, Moon J-Y, Lin D-Y. A method for Mendelian Randomization analyses with incomplete measurements on the exposure with application to metabolomics in HCHS/SOL. *HGG Adv*. 2024 Jan 11;5(1):100245. doi: 10.1016/j.xhgg.2023.100245. Epub 2023 Oct 28.

130. Lundin JI, Peters U, Hu Y, Ammous F, **Avery CL**, Benjamin EJ, Bis JC, Brody JA, Carlson C, Cushman M, Gignoux C, Guo X, Haessler J, Haiman C, Joehanes R, Kasela S, Kenny E, Lapalainien T, Levy D, Liu C, Liu Y, Loos RJF, Lu A, Matise T, North KE, Park SL, Ratliff SM, Reiner A, Rich SS, Rotter JI, Smith JA, Sotoodehnia N, Tracy R, Van den Berg D, Xu H, Ye T, Zhao W, Raffield LM, Kooperberg C; PAGE Study. Methylation patterns associated with C-reactive protein in racially and ethnically diverse populations. *Epigenetics*. 2024 Dec;19(1):2333668. doi: 10.1080/15592294.2024.2333668. Epub 2024 Apr 3.

129. Frank SM, Jaacks LM, Meyer K, Rose D, Adair LS, **Avery CL**, Taillie LS. Dietary quality and dietary greenhouse gas emissions in the USA: a comparison of the planetary health diet index, healthy eating index-2015, and dietary approaches to stop hypertension. *Int J Behav Nutr Phys Act*. 2024 Apr 2;21(1):36. doi: 10.1186/s12966-024-01581-y.

128. Martin SS, Aday AW, Almarzooq ZI, Anderson CAM, Arora P, Avery CL, Baker-Smith CM, Barone Gibbs B, Beaton AZ, Boehme AK, Commodore-Mensah Y, Currie ME, Elkind MSV, Evenson KR, Generoso G, Heard DG, Hiremath S, Johansen MC, Kalani R, Kazi DS, Ko D, Liu J, Magnani JW, Michos ED, Mussolino ME, Navaneethan SD, Parikh NI, Perman SM, Poudel R, Rezk-Hanna M, Roth GA, Shah NS, St-Onge MP, Thacker EL, Tsao CW, Urbut SM, Van Spall HGC, Voeks JH, Wang NY, Wong ND, Wong SS, Yaffe K, Palaniappan LP; American Heart Association Council on Epidemiology and Prevention Statistics Committee and Stroke Statistics Subcommittee. 2024 Heart Disease and Stroke Statistics: A Report of US and Global Data From the American Heart Association. *Circulation*. 2024 Feb 20;149(8):e347-e913.

127. Frank SM, Jaacks LM, **Avery CL**, Adair LS, Meyer K, Rose D, Taillie LS. Dietary quality and cardiometabolic indicators in the USA: A comparison of the Planetary Health Diet Index, Healthy Eating Index-2015, and Dietary Approaches to Stop Hypertension. *PLoS One*. 2024 Jan 10;19(1):e0296069. doi: 10.1371/journal.pone.0296069.

126. Lu Y, Pike JR, Hoozeveer RC, Walker KA, Raffield LM, Selvin E, **Avery CL**, Engel SM, Mielke MM, Garcia T, Palta P. Liver integrity and the risk of Alzheimer's disease and related dementias. *Alzheimers Dement*. 2024 Mar;20(3):1913-1922. doi: 10.1002/alz.13601.

†125. Conners KM, **Avery CL**, Syed FF. Advancing cardiovascular risk assessment with artificial intelligence. Opportunities and implications in North Carolina. *NC Med J* 2024, in press

124. Li Y, Wong KY, Howard AG, Gordon-Larsen P, Highland HM, Graff M, North KE, Downie CG, **Avery CL**, Yu B, Young KL, Buchanan VL, Kaplan R, Hou L, Joyce BT, Qi Q, Sofer T, Moon JY, Lin DY. Mendelian Randomization with Incomplete Measurements on the Exposure in the Hispanic Community Health Study/Study of Latinos. *HGG Adv*. 2023 Oct 9:100245. doi: 10.1016/j.xhgg.2023.100245.

123. Frank SM, Jaacks LM, Adair LS, **Avery CL**, Meyer KT, Rose D, Taillie LS. Adherence to the Planetary Health Diet Index and Correlation with Nutrients of Public Health Concern: An analysis of NHANES 2003-2018. *Am J Clin Nutr*. 2024. Feb;119(2):384-392. doi: 10.1016/j.ajcnut.2023.10.018.

122. Lee MP, Dimos S, Raffield L, Ballou AF, Arehart C, Correa A, de Vries P, Du Z, Gignoux C, Gordon-Larsen P, Guo X, Haessler J, Howard AG, Hu Y, Kent S, Monda KL, North KE, Peters U, Preuss M, Rich S, Rhodes S, Rotter J, Yao J, Yarosh R, Tsai M, Kooperberg C, Ballantyne CM, **Avery CL, Graff M. Ancestral diversity in lipoprotein(a) studies helps address evidence gaps. *Open Heart* 2023. Aug;10(2):e002382. doi: 10.1136/openhrt-2023-002382.

121. **Avery CL**, Howard AG, Lee HH, Lee MP, Koenigsberg SH, Ballou AF, Preuss M, Raffield L, Yarosh R, North KE, Gordon-Larsen P, Graff M. Branched chain amino acids harbor distinct and often opposing effects on health and disease. *Comm Med* 2023. Nov 28;3(1):172. doi: 10.1038/s43856-023-00382-x

120. Tsao CW, Aday AW, Almarzooq ZI, Anderson CAM, Arora P, **Avery CL**, Baker-Smith CM, Beaton AZ, Boehme AK, Buxton AE, Commodore-Mensah Y, Elkind MSV, Evenson KR, Eze-Nliam C, Fugar S, Generoso G, Heard DG, Hiremath S, Ho JE, Kalani R, Kazi DS, Ko D, Levine DA, Liu J, Ma J, Magnani JW, Michos ED, Mussolino ME, Navaneethan SD, Parikh NI, Poudel R, Rezk-Hanna M, Roth GA, Shah NS, St-Onge MP, Thacker EL, Virani SS, Voeks JH, Wang NY, Wong ND, Wong SS, Yaffe K, Martin SS; American Heart Association Council on Epidemiology and Prevention Statistics Committee and Stroke Statistics Subcommittee. Heart Disease and Stroke Statistics-2023 Update: A Report From the American Heart Association. *Circulation*. 2023 Jan 25.

119. Shemirani R, Belbin GM, **Avery CL**, Kenny EE, Gignoux CR, Ambite J-L. Selecting Clustering Algorithms for IBD Mapping. *Pac Symp Biocomput* 2023. 28:121-132

118. Anwar MY, Baldassari AR, Polikowsky HG, Sitlani CM, Highland HM, Chami N, Chen HH,

Graff M, Howard AG, Jung SY, Petty LE, Wang Z, Zhu W, Buyske S, Cheng I, Kaplan R, Kooperberg C, Loos RJF, Peters U, McCormick JB, Fisher-Hoch SP, **Avery CL**, Taylor KC, Below JE, North KE. Genetic pleiotropy underpinning adiposity and inflammation in self-identified Hispanic/Latino populations. *BMC Med Genomics* 2022. Sep 10;15(1):192. doi: 10.1186/s12920-022-01352-3.

117. Young WJ, Lahrouchi N, Isaacs A, Duong T, Foco L, Ahmed F, Brody JA, Salman R, Noordam R, Benjamins JW, Haessler J, Lyytikäinen LP, Repetto L, Concas MP, van den Berg ME, Weiss S, Baldassari AR, Bartz TM, Cook JP, Evans DS, Freudling R, Hines O, Isaksen JL, Lin H, Mei H, Moscati A, Müller-Nurasyid M, Nursyifa C, Qian Y, Richmond A, Roselli C, Ryan KA, Tarazona-Santos E, Thériault S, van Duijvenboden S, Warren HR, Yao J, Raza D, Aeschbacher S, Ahlberg G, Alonso A, Andreasen L, Bis JC, Boerwinkle E, Campbell A, Catamo E, Cocca M, Cutler MJ, Darbar D, De Grandi A, De Luca A, Ding J, Ellervik C, Ellinor PT, Felix SB, Froguel P, Fuchsberger C, Gögele M, Graff C, Graff M, Guo X, Hansen T, Heckbert SR, Huang PL, Huikuri HV, Hutri-Kähönen N, Ikram MA, Jackson RD, Junttila J, Kavousi M, Kors JA, Leal TP, Lemaitre RN, Lin HJ, Lind L, Linneberg A, Liu S, MacFarlane PW, Mangino M, Meitinger T, Mezzavilla M, Mishra PP, Mitchell RN, Mononen N, Montasser ME, Morrison AC, Nauck M, Nauffal V, Navarro P, Nikus K, Pare G, Patton KK, Pelliccione G, Pittman A, Porteous DJ, Pramstaller PP, Preuss MH, Raitakari OT, Reiner AP, Ribeiro ALP, Rice KM, Risch L, Schlessinger D, Schotten U, Schurmann C, Shen X, Shoemaker MB, Sinagra G, Sinner MF, Soliman EZ, Stoll M, Strauch K, Tarasov K, Taylor KD, Tinker A, Trompet S, Uitterlinden A, Völker U, Völzke H, Waldenberger M, Weng LC, Whitsel EA, Wilson JG, **Avery CL**, Conen D, Correa A, Cucca F, Dörr M, Gharib SA, Girotto G, Grarup N, Hayward C, Jamshidi Y, Järvelin MR, Jukema JW, Kääb S, Kähönen M, Kanters JK, Kooperberg C, Lehtimäki T, Lima-Costa MF, Liu Y, Loos RJF, Lubitz SA, Mook-Kanamori DO, Morris AP, O'Connell JR, Olesen MS, Orini M, Padmanabhan S, Pattaro C, Peters A, Psaty BM, Rotter JJ, Stricker B, van der Harst P, van Duijn CM, Verweij N, Wilson JF, Arking DE, Ramirez J, Lambiase PD, Sotoodehnia N, Mifsud B, Newton-Cheh C, Munroe PB. Genetic analyses of the electrocardiographic QT interval and its components identify additional loci and pathways. *Nat Commun.* 2022 Sep 1;13(1):5144. doi: 10.1038/s41467-022-32821-z. PMID: 36050321.

116. Lee M, Whitsel EA, **Avery CL**, Hughes TM, Griswold ME, Sedaghat S, Gottesman RF, Mosley TH, Heiss G, Lutsey PL. Variation in Population Attributable Fraction of Dementia Associated With Potentially Modifiable Risk Factors by Race and Ethnicity in the US. *JAMA Netw Open.* 2022;5(7):e2219672. doi:10.1001/jamanetworkopen.2022.19672

115. Tcheandjieu C, Zhu X, Hillard AT, Clarke SL, Napolino V, Ma S, Fang H, Sinnott-Armstrong N, Gorman BR, Pyarajan S, Huang J, Song R, Plomondon M, Maddox TM, Waldo SW, Lee K-M, Ho Y-L, Wojcik GL, Bustamente CD, Prichard JK, Buyske S, Kooperberg CL, Haessler J, Loos RFJ, Do R, Berbanck M, Chaudhary K, North KE, **Avery CL**, Graff M, Chen F, Haiman CA, Le Marchand LE, Wilkens LR, Bis JC, Leonard H, Shen B, Lange LA, Giri A, Lu Y, Dikilitas O, Kullo IJ, Stanaway IB, Jarvik GP, Gordon AD, Hebring S, Namjou B, Harley JB, Kaufman K, Koyama S, Ito K, Ishigaki K, Tsao N, Verma SS, Ritchie MD, Kember RL, Natarajan P, Kathiresan S, Bick AG, Levin M, Hauser ER, Miller DR, Vujkovic M, Huffman JE, Raghavan S, Klarin D, Lee JS, Voight BF, Saleheen D, Reaven PD, Lynch JA, Cho K, Gaziano JM, Rader DJ, Chang K-M, Damrauer SM, Wilson PW, Tang H, Sun YV, Tsao PS, O'Donnell CJ, Assimes TL. A large-scale multi-ethnic genome-wide association study of coronary artery disease. In press, *Nat Med* 2022

- †114. Downie CG, Highland HM, Lee MP, Raffield LM, Preuss M, Whitsel EA, Psaty BM, Sitlani CM, Graff M, **Avery CL**. Genome-wide analysis of variant-by-thiazide diuretic interactions on blood lipid concentrations in the UK Biobank identifies a novel LDL-C locus. *Circ Res*. 2022 Jun 14;101161CIRCRESAHA122321120. doi: 10.1161/CIRCRESAHA.122.321120.
113. **Avery CL**, Howard AH, Ballou AF, Buchanan VL, Collins JM, Downie CG, Graff M, Highland HM, Lee MP, Lilly AG, Lu K, Ragner JE, Staley BS, North KE, and Gordon-Larsen P. How genetic data can strengthen causal inference in exposomics. *Environ Health Perspect*. 2022. May;130(5):55001. doi: 10.1289/EHP9098. Epub 2022 May 9.
112. Highland HM, Wojcik GL, Graff M, Nishimura KK, Hodonsky CJ, Baldassari AR, Cheng I, Gignoux CR, Tao R, Li Y, Boerwinkle E, Fornage M, Haessler J, Hindorff LA, Hu Y, Justice AE, Lin BM, Lin D-Y, Stram DO, Haiman CA, Kooperberg C, Le Marchand L, Matisse TC, Kenny EE, Carlson CS, Stahl EA, **Avery CL**, North KE, Ambite JL, Buyske S, Loos RFJ, Peters U, Young KL, Bien SA, Huckins LM. Predicted gene expression in ancestrally diverse populations leads to discovery of novel susceptibility loci for lifestyle and cardiometabolic traits. *Am J Human Genet*. 2022. Apr 7;109(4):669-679
111. Polikowsky HG, Shaw DM, Petty LE, Chen H-H, Pruett D, Linklater JP, Viljoen KZ, Beilby JM, Highland HM, Levitt B, **Avery CL**, Mullan Harris K, Jones R, Below JE, Kraft SJ. Population-based genetic effects for developmental stuttering. *HGG Advances*. Dec 2;3(1):100073. doi: 10.1016/j.xhgg.2021.100073. eCollection 2022 Jan 13.
110. Downie CG, Dimos S, Wang Y, Bien SA, Raffield LM, Armstrong N, Polikowsky HG, Darst B, Polfus LM, Below JE, Irvin MR, Carlson C, Haiman C, Hindorff L, Phillips L, Liu S, Pankow J, Rich SS, Rotter J, Buyske S, Matisse T, Kooperberg C, Loos R, North KE, **Avery CL**, Graff M, Highland HM. Multi-ethnic GWAS and fine-mapping of glycemic traits identify novel loci in the PAGE study. *Diabetologia* 2022. Mar;65(3):477-489.
109. Rebolledo Fuentealba NI, Reyes M, Popkin BM, Adair LS, **Avery CL**, Corval C, Ng S, Tallie Smith L. Changes in Nonnutritive Sweetener Intake in a Cohort of Preschoolers After the Implementation of Chile's Law of Food Labeling and Advertising. *Pediatr Obes*. 2022. Jul;17(7):e12895. doi: 10.1111/ijpo.12895. Epub 2022 Jan 27.
108. Bauer AE, **Avery CL**, Shi M, Weinberg C, Olshan AF, Harmon Q, Luo J, Yang J, Manuck T, Williams N, McGinnis R, Morgan L, Klungsoyr K, Trogstad L, Magnus P, Engel SM. Do genetic variants modify the effect of smoking on risk of preeclampsia in pregnancy? *Am J Perinatol*. 2022. Nov 28. doi: 10.1055/s-0041-1740072.
107. Raffield LM, Howard AG, Graff M, Lin D, Cheng S, Demerath E, Ndumele C, Rebholz C, Seidemann S, Yu B, North KE, Gordon-Larsen P, **Avery CL**. Obesity duration, severity, and distribution trajectories and cardiovascular disease risk in the Atherosclerosis Risk in Communities Study. *J Am Heart Assoc*. 2021. Dec 21;10(24):e019946. doi: 10.1161/JAHA.121.019946.
106. Wen J, Xie M, Rowland B, Rosen JD, Sun Q, Tapia AL, Qian H, Kowalski MH, Shan Y, Young KL, Graff M, Argos M, **Avery CL**, Bien SA, Buyske S, Choquet H, Fornage M, Hodonsky C, Jorgenson E, Kooperberg C, Loos RJF, Liu Y, Moon J-Y, North KE, Rich SS, Rotter JI, Smith J,

Zhao W, Shang L, Wang T, Zhou X, Reiner AP, Raffield LM, Li Y. Transcriptome-wide association study of blood cell traits in African American and Hispanic/Latino Populations. *Genes* (Basel). 2021 Jul 8;12(7):1049. doi: 10.3390/genes12071049.

105. Sun Q, Rowland B, Wen J, Huang L, Lee MP, **Avery CL**, Franceschini N, North KE, Li Y, Raffield L. Analyses of Serum and Urine Biomarker Traits in UK Biobank Participants with African, South Asian, and East Asian Ancestry. *J Hum Genet*. 2021 Aug 11. doi: 10.1038/s10038-021-00968-0.

104. Hu Y, Bien SA, Nishimura KK, Haessler J, Hodonsky CJ, Baldassari AR, Highland HM, Wang Z, Preuss M, Sitlani CM, Wojcik GL, Tao R, Graff M, Huckins LM, Sun Q, Chen M-H, Mousas A, Auer PL, Lettre G, Tang W, Qi L, Thyagarajan B, Buyske S, Fornage M, Hindorf LA, Li Y, Lin D, Reiner AP, North KE, Loos RJF, Raffield LM, Peters U, **Avery CL**, Kooperberg C. Multi-ethnic genome-wide association analyses of white blood cell and platelet traits in the Population Architecture using Genomics and Epidemiology (PAGE) Study. *BMC Genomics*. 2021 Jun 9;22(1):432. doi: 10.1186/s12864-021-07745-5.

103. Sitlani CM, Baldassari AR, Highland HM, Hodonsky CJ, McKnight B, **Avery CL**. Comparison of multiple phenotype association tests using summary statistics in genome-wide association studies. *Hum Mol Genet*. 2021 Jul 9;30(15):1371-1383. doi: 10.1093/hmg/ddab126

102. **Avery CL**, Howard AG, Nichols HB. Comparison of 20-year obesity-related cancer mortality trends with heart disease mortality trends in the US. *JAMA Network Open*. 2021 May 3;4(5):e218356. doi: 10.1001/jamanetworkopen.2021.8356.

101. Wang Y, Wang H, Howard AH, Tsilimigras MCB, **Avery CL**, Meyer KA, Sha W, Sun S, Su C, Wang Z, Fodor AA, Zhang B, Gordon-Larsen P. Gut microbiota and host metabolites associated with blood pressure in Chinese adults. *Hypertension*. 2021. Feb;77(2):706-717

100. **Avery CL**, Howard AG, Nichols HB. Trends in U.S. cancer and heart disease mortality, 1999-2018. *Circulation*. 2021 Jan 19;143(3):287-288

99. Shemirani R, Belbin GM, **Avery CL**, Kenny EE, Gignoux CR, Ambite J-L. Rapid detection of identity-by-descent tracts for mega-scale datasets. *Nat Commun*. 2021 Jun 10;12(1):3546. doi: 10.1038/s41467-021-22910-w.

98. Wang Y, Wang H, Howard AG, Tsilimigras MCB, **Avery CL**, Meyer KA, Sha W, Sumner SJ, Zhang J, Su C, Wang Z, Zhang B, Fodor AA, Gordon-Larsen P. Sodium and Potassium Consumption was Associated with Gut Microbiota and Plasma Metabolome Involved in Cardiometabolic Health: A Population-based Study of Chinese Adults. *Am J Clinical Nutrition*. 2020 Dec 10;112(6):1599-1612.

97. Wang Y, Wang H, Howard AG, Meyer KA, Tsilimigras M, **Avery CL**, Sha W, Sun S, Zhang J, Su C, Wang Z, Zhang B, Fodor A, Gordon-Larsen P. Circulating Short-chain Fatty Acids are Positively Associated with Adiposity Measures in Chinese Adults. *Nutrients*. 2020 Jul 17;12(7):2127

†96. Engeda JC, Lhachimi SK, Rosamond WD, Lund JL, Keyserling TC, Safford MM, Colantonio LD, Muntner P, **Avery CL**. Projections of atherosclerotic cardiovascular disease and type 2 diabetes across statin treatment guidelines. *PLoS Med.* 2020. Aug 26;17(8):e1003280

95. Kats D, Evenson KR, Zeng D, **Avery CL**, Palta P, Kritchevsky SB, Heiss G. Physical activity from mid- to late-life: duration and intensity. The ARIC Study. *Aging (Albany NY)*. 2020 Mar 13;12(5):4592-4602.

94. Martinez-Miller EE, Prather A, Robinson WR, **Avery CL**, Yang YC, Haan MN, Aiello AE. Acculturation, Cognitive Performance and Decline, and Incident Dementia/CIND: The Sacramento Area Latino Study on Aging. *Am J Epidemiol* 2020 Nov 2;189(11):1292-1305

†93. Hodonsky CJ, Bien S, Baldassari AR, Raffield L, Highland HM, Sitlani CM, Wojcik GL, Tao R, Graff M, Tang W, Thyagarajan B, Buyske S, Fornage M, Hindorff LA, Li Y, Lan D, Reiner AP, North KE, Roos RJF, Kooperberg C, **Avery CL**. Ancestry-specific associations identified in genome-wide combined-phenotype study of red blood cell traits emphasize benefits of diversity in genomics. *BMC Genomics*, 2020. Mar 14;21(1):228. doi: 10.1186/s12864-020-6626-9.

92. Ntalla I, Weng, L-C, Cartwright JH, Hall AW, Svinbjornsson G, Tucker NR, Choi SH, Chaffin MD, Roselli C, Barnes MR, Mifsud B, Warren HR, Hayward C, Marten J, Cranley JJ, Concas MP, Gasparini P, Boutin T, Kolcic I, Polasek O, Ruden I, Araujo NM, Lima-Costa MF, Ribeiro ALP, Souza RP, Tarazona-Santos E, Giedraitis V, Ingelsson E, Mahajan A, Morris AP, Del Greco F, Foco L, Gogele M, Hicks AA, Cook JP, Lind L, Lindgren CM, Sundstrom J, Nelson CP, Riaz MB, Samani NJ, Sinagra G, Ulivi S, Kahonen M, Mishra PP, Mononen N, Nikus K, Caulfield MJ, Dominiczak A, Padmanabhan S, Montasser ME, O'Connell JR, Ryan K, Shuldiner AR, Aeschbacher S, Conen D, Risch L, Theriault S, Hutri-Kahonen N, Lehtimaki T, Lyytikainen L-P, Raitakari OT, Barnes CLK, Campbell H, Joshi PK, Wilson JF, Isaacs A, Kors JA, van Juijn CM, Huang PL, Gudnason V, Harris TB, Launer LJ, Smith AV, Bottinger EP, Loos RJF, Nadkarni GN, Preuss MH, Correa A, Mei H, Wilson J, Meitinger T, Nuller-Nurasid M, Peters A, Waldenberger M, Mangino M, Spector TD, Rienstra M, van de Vegte YJ, van der Harst P, Verweij N, Kaab S, Schramm K, Sinner MF, Strauch K, Cutler JM, Fatkin D, London B, Olesen M, Roden DM, Shoemaker B, Smith JG, Biggs ML, Bis JC, Brody JA, Psaty BM, Rice K, Sotoodehnia N, De Grandi A, Fuchsberger C, Pattaro C, Pramstaller PP, Ford I, Jukema JW, Macfarlane PW, Trompet S, Dorr M, Felix SB, Wolker U, Weiss S, Havulinna AS, Jula A, Saaksjarvi K, Salomaa V, Guo X, Heckbert SR, Lin HJ, Rotter JI, Taylor KD, Yao J, de Mutsert R, Maan AC, Mook-Kanamori DO, Noordam R, Cucca F, Ding J, Lakatta EG, Qian Y, Tarasov KV, Levy D, Lin H, Newton-Cheh CH, Lunetta KL, Murray AD, Porteous DJ, Smith BH, Stricker BH, Uitterlinden A, van den Berg ME, Haessler J, Jackson RD, Kooperberg C, Peters U, Reiner AP, Whitsel EA, Alonso A, Arking DE, Boerwinkle E, Ehret GB, Soliman EZ, **Avery CL**, Gogarten SM, Kerr KF, Laurie CC, Seyerle AA, Stilp A, Assa S, Said A, van der Ende MY, Lambiase PD, Orini M, Ramirez J, Van Duijvenboden S, Arnar DO, Gudbjartsson DF, Holm H, Sulem P, Thorleifsson G, Thorolfsson RB, Thorsteinsdottir U, Benjamin EJ, Tinker A, Stefansson K, Ellinor PT, Jamshidi Y, Lubitz SA, Munroe PB. Multi-ancestry GWAS of the electrocardiographic PR interval identifies 210 loci underlying cardiac conduction. 2020 *Nature Communications*, May 21;11(1):2542. doi: 10.1038/s41467-020-15706-x.

†91. Baldassari AR, Sitlani CM, Highland HM, Arking DE, Buyske S, Darbar D, Gondalia R, Graff

M, Guo X, Heckbert SR, Hindorff LA, Hodonsky CJ, Chen Y-DI, Kaplan RC, Peters U, Post W, Reiner AP, Rotter JI, Shohet RV, Sotoodehnia N, Tao R, Taylor KD, Yao J, Kenny EE, Lin HJ, Soliman EZ, Whitsel EA, North KE, Kooperberg C, **Avery CL**. Multi-ethnic genome-wide association study of decomposed cardiac conduction phenotypes illustrates strategies to identify and characterize evidence of shared genetic effects for complex traits. *Circ Precision Med*. 2020 Aug;13(4):e002680. doi: 10.1161/CIRCGEN.119.002680.

90. de Las Fuentes L, Sung YJ, Sitlani CM, **Avery CL**, Bartz TM, Keyser C, Evans DS, Li X, Musani SK, Rutter R, Smith AV, Sun F, Trompet S, Xu H, Arnett DK, Bis JC, Broeckel U, Busch EL, Chen YI, Correa A, Cummings SR, Floyd JS, Ford I, Guo X, Harris TB, Ikram MA, Lange L, Launer LJ, Reiner AP, Schwander K, Smith NL, Sotoodehnia N, Stewart JD, Stott DJ, Stürmer T, Taylor KD, Uitterlinden A, Vasani RS, Wiggins KL, Cupples LA, Gudnason V, Heckbert SR, Jukema JW, Liu Y, Psaty BM, Rao DC, Rotter JI, Stricker B, Wilson JG, Whitsel EA. Genome-wide meta-analysis of variant-by-diuretic interactions as modulators of lipid traits in persons of European and African ancestry. *Pharmacogenomics J*, 2020. Jun;20(3):482-493.

89. Gellert KS, Keil AP, Zeng D, Lesko CR, Aubert RE, **Avery CL**, Lutsey PL, Siega-Riz AM, Windham BG, Heiss G. Reducing the population burden of coronary heart disease by modifying adiposity: Estimates from the Atherosclerosis Risk In Communities (ARIC) Study. *J Am Heart Assoc* 2020. 9(4): e012214

88. Irvin MR, Sitlani CS, Bis JA, Whitsel EA, Steward J, Raffield L, Sun F, Liu C-T, Xu H, Cupples AL, Rossing P, Smith A, Noguiera NRZ, Noordam R, Rotter J, Yao J, Li X, Guo X, Limdi N, Lange L, Ahluwalia TS, Arnett DK, **Avery CL**. Genome wide association study of apparent treatment resistant hypertension in the CHARGE consortium: The CHARGE Pharmacogenetics Working Group. *Pharmacogenomics*. 2020. Nov 15;32(12):1146-1153.

87. Wang Y, Howard AG, Adair LS, **Avery CL**, Wang H, Gordon-Larsen P. Waist circumference change is associated with blood pressure change independent of BMI change. *Obesity* (Silver Spring). 2020 Jan;28(1):146-153.

86. Pendergrass SA, Buyske S, Jeff JM, Frase A, Dudek S, Bradford Y, Ambite JL, **Avery CL**, Buzkova P, Deelman E, Fesinmeyer MD, Haiman C, Heiss G, Hindorff LA, Hsu CN, Jackson RD, Lin Y, Le Marchand L, Matisse TC, Monroe KR, Moreland L, North KE, Park SL, Reiner A, Wallace R, Wilkens LR, Kooperberg C, Ritchie MD, Crawford DC. A phenome-wide association study (PheWAS) in the Population Architecture using Genomics and Epidemiology (PAGE) study reveals potential pleiotropy in African Americans. *PLoS One*, 2019. 14(12): e0226771

85. Kowalski MH, Qian H, Hou Z, Rosen JD, Tapia AL, Shan Y, Jain D, Argos M, Arnett DK, **Avery CL**, Barnes KC, Becker LC, Bien SA, Bis JC, Blangero J, Boerwinkle E, Bowden DW, Buyske S, Cai J, Cho MH, Choi SH, Choquet H, Cupples LA, Cushman M, Daya M, de Vries P, Ellinor PT, Faraday N, Fornage M, Gabriel S, Ganesh S, Graff M, Gupta N, He J, Heckbert SR, Hidalgo B, Hodonsky CJ, Irvin MR, Johnson AD, Jorgenson E, Kaplan R, Kardina SLR, Kelly TN, Kooperberg C, Lasky-Su JA, Loos RJJ, Lubitz SA, Mathias RA, McHugh CP, Montgomery C, Moon J-Y, Morrison AC, Palmer ND, Pankratz N, Papanicolaou GJ, Peralta JM, Peyser PA, Rich SS, Rotter JI, Silverman EK, Smith JA, Smith NL, Taylor KD, Thornton TA, Tiwari HK, Tracy RP,

Wang T, Weiss ST, Weng L-C, Wiggins KL, Wilson JG, Yanek LR, Zöllner S, North KE, Auer PL, Raffield LM, Reiner AP, Lu Y. 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. 2019, *PloS Genetics*. 2019 Dec 23;15(12):e1008500.

84. Aburto TC, Gordon-Larsen P, Poti JM, Howard AG, Adair LS, **Avery CL**, Popkin BM. Is a hypertension diagnosis associated with improved dietary outcomes within 2 to 4 years? A fixed-effects analysis from the China Health and Nutrition Survey. 2019, *J Am Heart Assoc*. e012703.

83. Swenson BR, Louie T, Lin HH, Mendez-Giraldez R, Below JE, Laurie CC, Kerr KF, Highland HM, Ryckman K, Kooperberg C, Soliman EZ, Seyerle AA, Guo X, Taylor KD, Yao J, Heckbert SR, Petty LE, Cheng S, Below N, Whitsel EA, Hanis CL, Rotter JI, Sofer T, **Avery CL**, Sotoodenhia N. GWAS of QRS Duration Identifies New Loci Specific to Hispanic/Latino Populations. *Plos One*. 2019. Jun 28;14(6):e0217796

†82. Engeda JC, Stackhouse A, White M, Rosamond WD, Lhachimi SK, Lund JL, Keyserling TC, **Avery CL**. Narrowing the focus to primary prevention populations by broadening the view: Examining the association between statins and diabetes among randomized controlled trials and observational studies. *Diabetes Res Clin Pract*. 2019. May;151:96-105

81. Wojcik GL, Graff M, Nishimura K, Tao R, Haessler J, Gignoux CR, Highland HM, Patel YM, Sorokin EP, **Avery CL**, Belbin GM, Bien SB, Cheng I, Hodonsky CJ, Huckins LM, Jeffs J, Justice AE, Kocarnik JM, Lin BM, Lu Y, Nelson SC, Park S-S L, Preuss M, Richard MA, Schurmann C, Setiawan VW, Vishnu A, Verbanck M, Walker R, Young KL, Zubair N, Ambite JL, Boerwinkle E, Bottinger E, Bustamante CD, Caberto C, Conomos MP, Deelman E, Do R, Doheny K, Fernandez-Rhodes L, Fornage M, Heiss G, Hindorff L, Jackson RD, James R, Laurie CA, Laurie CC, Li Y, Lin DY, Nadkarni G, Pankow J, Pooler LC, Reiner AP, Romm J, Sabati C, Sheng X, Stahl E, Stram D, Thornton TA, Wassel CL, Wilkens LR, Yoneyama S, Buyske S, Haiman C, Kooperberg C, Le Marchand L, Loos RJF, Matise TC, North KE, Peters U, Kenny EE, Carlson CS. Genetic diversity turns a new PAGE in our understanding of complex traits. *Nature*, 2019. Jun;570(7762):514-518

80. Martinez-Miller EE, Prather A, Robinson WR, **Avery CL**, Yang YC, Haan MN, Aiello AE. US acculturation and poor sleep among an intergenerational cohort of adult Latinos in Sacramento, California. *Sleep*. 2019. Mar 1;42(3)

79. Kocarnik JM, Richard M, Graff M, Haessler J, Carlson C, Carty CL, Reiner AP, **Avery CL**, Ballantyne CM, LaCroix AZ, Assimes TL, Barbalić M, Pankratz N, Tang W, Chen D, Pereira R, Bien S, Nishimura K, Best LG, Ambite JL, Cheng I, Crawford D, Hindorff LA, Heiss G, North KE, Haiman CA, Peters U, Le Marchand , Kooperberg C. Discovery, fine-mapping, and conditional analyses of genetic variants associated with C-reactive protein in multiethnic populations using the Metabochip in the Population Architecture using Genomics and Epidemiology (PAGE) study. *Hum Mol Genet*. 2018. Aug 15;27(16):2940-2953

†78. Hodonsky CJ, Schurrman C, Schick UM, Kocarnik J, Tao R, van Rooij FJA, Wassel C, Buyske S, Fornage M, Hindorff LA, Psaty BM, Ganesh SK, Lin DY, North KE, Kooperberg CL, Reiner AP, Loos R, **Avery CL**. Generalization and fine mapping of RBC trait genetic associations to multi-ethnic

populations: The PAGE Study. *Am J Hematol*. 2018 Jun 15;10.1002/ajh.25161

77. van Setten J, Brody J, Campbell H, Del Greco F, Evans DS, Givson Q, Gudbjartsson DF, Kerr FK, Krijthe BP, Muller C, Muller-Nyrasyid M, Nolte IM, Padmanabhan S, Robino A, Smith AV, Sterj M, Swenson B, Tanaka T, Teumer A, Trompet S, Ulivi S, Verweji N, Yin X, Arnar DO, Asselbergs FW, **Avery CL**, Bis J, Butler AM, Blankenberg S, Boerwinkle E, Buckley BM, den Hoed M, Dominczak AF, Ehret GB, Eijgelsheim M, Felix SB, Franco OH, Gharib S, Harris TB, Heckbert S, Holm H, Ilaria G, Lorio A, Kolcic I, Kors JA, Lakatta EG, Launer LJ, Lin HJ, Lin H, Liu Y, Loos R, MacFarlane P, Magnani JW, Mateo I, Meitinger T, Mitchell B, Munzel T, Papanicolaou GJ, Peters A, Pfeufer A, Pramstaller PM, Rotter JI, Rudan I, Samani NJ, Schlessinger D, Aldana CTS, Sinner M, Snieder H, Soliman E, Spector TD, Stott DJ, Strauch K, Tarasov K, Uitterlinden AG, Volker U, Volzke H, Waldenberger M, Wild PS, Zeller T, Alonso A, Bandinelli S, Benjamin EJ, Cucca F, Cummings SR, Dorr M, Ferrucci L, Gasparini P, Gudnason V, Hayward C, Heckbert SR, Hicks AA, Jamshidi Y, Jukema JW, Kaab S, Munroe PB, Parsa A, Polasek O, Psaty BM, Schnabel RB, Sinagra G, Stefansson K, Stricker BH, van der Harst P, van Duijn C, Wilson JF, de Bakker PIW, Isaacs A, Arking DA, Sotoodehnia N. Genome-wide association meta-analysis of PR interval identifies 47 novel loci associated with atrial and atrioventricular electrical activity. *Nat Commun*, 2018.

76. Jarmul J, Pletcher MJ, Hassmiller Lich K, Wheeler SB, Weinberger M, **Avery CL**, Jonas DE, Earnshaw S, Pignone M. Cardiovascular Genetic Risk Testing for Targeting Statin Therapy in the Primary Prevention of Atherosclerotic Cardiovascular Disease: A Cost-Effectiveness Analysis.. Interpreting a 27-SNP cardiovascular genetic risk score in combination with traditional cardiovascular risk factors for prediction of atherosclerotic cardiovascular disease. *Circ Cardiovasc Qual Outcomes*. 2018; Apr;11(4): e00417

75. Hardy ST, **Avery CL**, Zeng D, Heiss G. Primary Prevention of Chronic Kidney Disease Through Population-Based Strategies for Blood Pressure Control: The ARIC Study. *J Clin Hypertens*. 2018. Jun;20(6):1018-1026

74. Napier MD, Franceschini N, Gondalia R, **Avery CL**, Stewart JD, Mendez-Giraldez R, Sitlani CM, Seyerle AA, Highland HM, Li Y, Wilhelmsen KC, Yan S, Duan Q, Roach J, North KE, Reiner AP, Zhang Z-M, Tinker LF, Liao D, Laurie C, Gogarten SM, Lin HJ, Brody JA, Sotoodehnia N, Whitsel EA. A trans-ethnic meta-analysis and genome-wide association study of ventricular and supraventricular ectopy. *Sci Rep*, 2018 Apr. 8(1): 5675

73. Irvin MR, Sitlani CM, Noordam R, **Avery CL**, Bis JC, Floyd JS, Hofman A, Li J, Limdi NA, Srinivasasainagendra V, Stewart J, Li J, Wilson JG, Lipovich L, Smith A, Bartz TM, Hofman A, Whitsel EA, Uitterlinden AG, Wiggins KL, Wilson JG, Zhi D, Stricker BH, Arnett DK, Psaty BM, Lange LA. Genome-wide meta-analysis of SNP-by-thiazide diuretic and SNP-by-ACEI/ARB effect on serum potassium in cohorts of European and African ancestry in CHARGE. *Pharmacogenomics J*. 2018. Feb;19(1):97-108

†72. Engeda JC, Holliday KM, Hardy ST, Chakladar S, Zeng D, Lin DY, Howard BV, Daviglius ML, Schreiner PJ, **Avery CL**. Why metrics matter: how distinct cholesterol metrics could alter prevention efforts. *Sci Rep*, 2018 Apr. 8(1): 6238

71. Busch EL, Don PK, Chu H, Richardson DB, Keku TO, Eberhard DA, **Avery CL**, Sandler RS. Diagnostic accuracy and prediction increment of markers of epithelial-mesenchymal transition to assess cancer cell detachment from primary tumors. *BMC Cancer*, 2018 Jan. 18(1): 82.

†70. Seyerle AA, Lin HJ, Gogarten SM, Stilp A, Mendez-Giraldez R, Soliman EZ, Baldassari A, Graff M, Heckbert SR, Kerr KF, Kooperberg C, Rodriguez CJ, Guo X, Ryckman KK, Yao J, Sotoodehnia N, Taylor KD, Whitsel EA, Rotter JI, Laurie CC, **Avery CL**. Genome-Wide Association Study of PR Interval in Hispanic/Latinos Identifies Novel Locus at *ID2*. *Heart*, 2018 Jun. 104(11):904-911.

69. Bauer AE, **Avery CL**, Shi M, Weinberg CR, Olshan AF, Harmon QE, Luo J, Yang J, Manuck T, Wu MC, Klungsoyr K, Trogstad L, Magnus P, Engel SM. Investigation of the inverse smoking-preeclampsia relationship through the exploration of gene by smoking interactions. *Paediatr Perinat Epidemiol*, 2018 Jan. 32(1): 1-12.

†68. Seyerle AA, Sitlani CM, Noordam R, Gogarten SM, Li J, Li X, Evans DS, Sun F, Isaacs A, Stewart J, Trompet S, Bis JC, Peloso GM, Brody JA, Broer L, Busch EL, Duan Q, Stilp AM, O'Donnell C, MacFarlane P, Floyd JS, Kors JA, Lin HJ, Li-Gao R, Sofer T, Giráldez RM, Cummings SR, Heckbert SR, Hofman A, Ford I, Li Y, Newton-Cheh C, Napier MD, Kerr KK, Prener AP, Rice KM, Roach J, Buckley BM, Soliman EZ, de Mutsert R, Sotoodehnia N, Uitterlinden AG, North KE, Lee C, Strümer T, Taylor KD, Wiggins KL, Wilson JG, Chen Y-DI, Kaplan RC, Wilhelmsen K, Cupples LA, Salomaa V, Jukema JW, Liu Y, Rosendaal FR, Lange LA, Vasana RS, Smith AV, Stricker BH, Laurie CC, Rotter JI, Whitsel EA, Psaty BM, **Avery CL**. Pharmacogenomics Study of Thiazide Diuretics and QT Interval in Multi-Ethnic Populations: The Cohorts for Heart and Aging Research using Genomic Epidemiology Pharmacogenetics Working Group (CHARGE PWG). *Pharmacogenomics J*, 2018 Apr. 18(2): 215-226.

67. Akinkugbe AA, **Avery CL**, Barritt AS, Cole SR, Lerch M, Mayerle J, Offenbacher S, Petersmann A, Nauck M, Völzke H, Slade GD, Heiss G, Kocher T, Holtfreter B. Do genetic markers of inflammation modify the relationship between periodontitis and nonalcoholic fatty liver disease? Findings from the SHIP study. *J Dent Res*, 2017 Nov. 96(12):1392-1399.

†66. Méndez-Giráldez R, Gogarten SM, Below JE, Yao J, Seyerle AA, Kooperberg C, Soliman EZ, Rotter JI, Kerr KF, Rickman K, Taylor KD, Petty LE, Conomos MP, Sotoodehnia N, Cheng S, Highland HM, Heckbert SR, Sofer T, Guo X, Whitsel EA, Lin HH, Hanis CL, Laurie CC, **Avery CL**. GWAS of QT interval in Hispanics/Latinos generalizes previously identified loci and identifies population-specific signals. *Sci Rep*, 2017 Dec. 7(1): 17075

65. Kent ST, Rosenson RS, **Avery CL**, Correa A, Cummings SR, Cupples LA, Cushman M, Evans DS, Gudnason V, Harris TB, Howard G, Irvin MR, Judd SE, Jukema JW, Lange L, Levitan EB, Li X, Liu Y, Postmus I, Psaty MB, Rotter JI, Safford MM, Sitlani CM, Smith AV, Stewart JD, Trompet S, Sun F, Vasana RS, Woolley JM, Whitsel EA, Wiggins KL, Wilson JG, Munter P. *PCSK9* loss-of-function mutations, low-density lipoprotein cholesterol, and coronary heart disease and stroke risk: data from nine population-based cohorts of African Americans and whites. *Circ Cardiovasc Genet*, 2017 Aug. 10(4): e001632.

64. Kerr KF, **Avery CL**, Lin HJ, Raffield LM, Zhang Q, Gogarten S, Jain D, Sofer T, Laurie CC,

Mendez-Giraldez R, Kooperberg C, Hohensee C, Li Y, Perez M, Yao J, Guo X, Taylor KD, Lange LA, Rotter JI, Heckbert SR, Sotoodehnia N, Wilson JG, Whitsel EA. Genome-wide Association Study of Heart Rate and Its Variability in Hispanic/Latino Cohorts. *Heart Rhythm*, 2017 Nov. 14(11): 1675-1684.

63. Christophersen IE, Magnani JW, Yin X, Arking D, Niemeijer MN, Lubitz SA, **Avery CL**, Duan Q, Felix SB, Bis J, Kerr K, Isaacs A, Muller-Nurysid M, Muller C, North KE, Reiner AP, Tinker LF, Kors JA, Teumer A, Petersmann A, Sinner MF, Waldenberger M, Peters A, Meitinger T, Limacher MC, Wilhelmsen KC, Hofman A, Uitterlinden A, Krijthe BP, Zhang Z-M, Schnabel RB, Kaab S, van Duijn C, Rotter JI, Sotoodehnia N, Volker W, Dorr M, Li Y, Soliman EZ, Alonso A, Whitsel EA, Stricker BH, Benjamin EJ, Heckbert SR, Ellinor PT. Sixteen genetic loci associated with the electrocardiographic P wave. *Circ Cardiovasc Genet*, 2017 Aug. 10(4): e001667.

62. Nolte IM, Munoz ML, Traganta V, Amare A, Jansen R, Vaes A, der Heyde B, **Avery CL**, Bis JC, Dierckx B, Gogarten SM, Goyette P, Hernesniemi J, Huikari V, Hwang S-J, Jaju D, Kerr KF, Kluttig A, Krijthe BP, Kumar J, Lyytikainen L-P, Maihofer AX, Minassian A, Muller-Nurasyd M, Nivard M, Salvi E, Stewart JD, Thayer JF, van der Laan SW, van der Most PJ, Verweij N, Wong A, Zabaneh D, Zafarmand MH, Abdellaoui A, Albert C, Alonso A, Ashar F, Auvinen J, Axelsson T, Baker DG, Barcella M, Bayoumi R, Beringa RJ, Boomsma D, Boucher G, Britton AR, Christophersen I, de Bakker PI, Dietrich A, Ehret GB, Ellinor PT, Eskola M, Felix JF, Floras J, Franco OH, Friberg P, Gademan MGJ, Geyer MA, Giedraitis V, Hartman CA, Hemerich D, Hofman A, Hottenga J-J, Huikuri H, Hutri-Kahonen N, Jouven X, Junttila J, Juonala M, Kiviniemi AM, Kors JA, Kumari M, Kuznetsova T, Laurie CC, Lefrandt JD, Li Y, Li Y, Liao D, Limacher MC, Lin HJ, Lindgren CM, Lubitz SA, Mahajan A, McKnight B, Schwabedissen HM, Milanese Y, Mononen N, Morris AP, Nalls MA, Navis G, Neijts M, Nikus K, North KE, O'Connor DT, Ormel J, Perz S, Peters A, Psaty BM, Raitakari OT, Risbrough VB, Sinner M, Siscovick D, Smit JH, Smith NL, Soliman EZ, Sotoodehnia N, Staessen JA, Stein PK, Stilp AM, Stolarz-Skrzpek K, Strauch K, Albarwani S, Sundstrom J, Sweene CA, Syvanen A-C, Tardif J-C, Taylor KD, Teumer A, Thornton TA, Tinker LE, Uitterlinden AG, van Setten J, Voss A, Waldenberger M, Wilhelmsen KC, Willemsen G, Wong Q, Zhang Z-M, Zonderman AB, Cusi D, Evans MK, Greiser HK, Hassan M, Ingelsson E, Jarvelin M-R, Kaab S, Kahonen M, Kivimaki M, Kooperberg CL, Kuh D, Lehtimaki T, Lind L, Nievergelt CM, O'Donnell CJ, Oldehinkel AJ, Penninx B, Reiner AP, Reise H, Rioux JD, Rotter JI, Sofer T, Stricker BH, Tiemeier H, van der Harst P, van Rooon AM, Vrikkotte TGM, Asselbergs FW, Brundel BJJM, Heckbert SR, Whitsel EA, den Hoed M, Sneider H, de Gues E. Identification of genetic loci associated with heart rate variability and their effects on heart rate and cardiac disease risk. *Nat Commun*, 2017. Jun 14;8:15805

†61. Hardy ST, Holliday KM, Chakladar S, Engeda JC, Allen NB, Jones DL, Schreiner PJ, Shay CM, Heiss G, Lin DY, Zeng D, **Avery CL**. Heterogeneity in blood pressure transitions over the life course: age-specific emergence of gender and race/ethnic disparities in the US. *JAMA Cardiol*. 2017 Jun 1;2(6):653-661.

†60. Holliday KM, Lin DY, Chakladar S, Cai J, Castaneda SF, Daviglius ML, Evenson KR, Marquez DX, Qi Q, Shay CM, Sotres-Alvarez D, Vidot DC, Zeng D, **Avery CL**. Targeting Physical Activity Interventions for Adults: When do we need to intervene? *Prev Med*. 2017 Apr;97:13-18

59. **Avery CL**, Wassel CL, Richard MA, Highland HM, Bien S, Zubair N, Soliman E, Z., Fornage M, Bielinski SJ, Tao R, Seyerle AA, Shah SJ, Lloyd-Jones DM, Buyske S, Rotter JI, Post WS, Rich SS, Hindorff LA, Jeff JM, Shohet RV, Sotoodehnia N, Lin DY, Whitsel EA, Peters U, Haiman CA, Crawford DC, Kooperberg C, North KE. Fine-Mapping of QT Regions in Global Populations Refines Previously Identified QT Loci and Identifies Signals Unique to African and Hispanic Descent Populations. *Heart Rhythm*, 2017 Apr;14(4):572-580

*58. Noordam R, Sitlani CM, **Avery CL**, Stewart DJ, Gogarten JD, Wiggins KL, Trompet S, Warren HR, Sun F, Evans DS, Li X, Li J, Smith AV, Bis JC, Brody JA, Busch EL, Caulfield MJ, Chen Y-D I, Cummings SR, Cupples LA, Duan Q, Franco OH, Giraldez RM, Harris TB, Heckbert SR, van Heemst D, Hofman A, Floyd JF, Kors JA, Launer LJ, Li Y, Li-Gao R, Lange LA, Lin HJ, Mook-Kanamori DO, de Mutsert R, Napier MD, Newton-Cheh C, Poulter N, Reiner AP, Rice KM, Roach J, Sattar N, Seer P, Seyerle AA, Slagboom PE, Soliman EZ, Sotoodehnia N, Stott DJ, Stürmer T, Taylor KD, Uitterlinden AG, Wihlemlsen KC, Wilson JG, Gudnason V, Jukema JW, Laurie CC, Liu Y, Munroe PB, Rosendaal FR, Rotter JI, Vasani R, Psaty BM, Stricker B, Whitsel EA. A genome-wide interaction analysis of tri/tetracyclic antidepressants and heart rate and QT interval: a pharmacogenomics study from the CHARGE consortium. *J Med Genet*, 2017 May;54(5):313-323

*57. Floyd JS, Sitlani CM, **Avery CL**, Noordam R, Li X, Smith AV, Gogarten SM, Lin J, Broer L, Evans DS, Trompet S, Brody JA, Stewart JD, Seyerle AA, Roach J, Lange LA, Lin HJ, Kors JA, Harris TB, Li-Gao R, Sattar N, Cummings SR, Aviles-Santa L, Wiggins KL, Napier MD, Stürmer T, Bis JC, Kerr KK, Uitterlinden AG, Taylor KD, Stott DJ, de Mutsert R, Launer LJ, Busch EL, Giraldez RM, Sotoodehnia N, Soliman EZ, Li Y, Duan Q, Hofman A, Mook-Kanamori DO, Slagboom PE, Wihlemlsen KC, Reiner AP, Chen Y-DI, Heckbert SR, Kaplan RC, Rice KM, Jukema JW, Liu Y, Rosendaal FR, Gudnason V, Wilson JG, Rotter JI, Laurie CC, Psaty BM, Whitsel EA, Cupples LA, Stricker BH. Large-scale pharmacogenomic study of sulfonylureas and the QT, JT, and QRS intervals: CHARGE Pharmacogenomics Working Group. *Pharmacogenomics J*, 2018 Jan. 18(1):127-135.

56. Sofer T, Heller R, Bogomolov M, **Avery CL**, Graff M, North KE, Reiner AP, Thornton T, Rice K, Benjamin Y, Laurie CC, Kerr K. Leveraging generalization-based design to increase discovery power in GWAS, with application to the Hispanic Community Health Study/Study of Latinos. *Genet Epidemiol*, 2017 Apr;41(3):251-258

55. Gondalia RB, **Avery CL**, Napier MD, Mendez RG, Stewart JD, Sitlani CM, Li Y, Wihlemlsen KC, Duan Q, Roach J, North KE, Reiner AP, Zhang Z-M, Tinker LF, Liao D, Whitsel EA. Genome-wide susceptibility to particulate matter-associated QT prolongation. *Environ Health Perspect*. 2017 Jun 8;125(6):067002

54. Postmus I, Warren HR, Trompet S, Arsenault BJ, **Avery CL**, Bis J, Chasman DI, de Keyser CE, Deshmukh HA, Evans DS, Feng Qi, Li X, Smit R, Smith AV, Sun F, Taylor KD, Arnold AM, Barnes MR, Barratt BJ, Betteridge J, Boekholdt SM, Boerwinkle E, Buckley BM, Chen Y-D, de Craen AJM, Cummings SR, Denny JC, Dube MP, Durrington PN, Eiriksdottir G, Ford I, Guo X, Harris TB, Heckbert SR, Hofman A, Hovingh GK, Kastelein JJP, Launer LJ, Liu C-T, Liu Y, Lumley T, McKeigue PM, Munroe PB, Neil A, Nickerson DA, Nyberg F, O'Brien E, O'Donnell

CJ, Post W, Poulter N, Vasan RS, Rice K, Rich SS, Rivadeneira F, Sattar N, Sever P, Shaw-Hawkins S, Shields DC, Slagboom PE, Smith JL, Smith JD, Sotoodehnia N, Stanton A, Stott DJ, Strickler BH, Sturmer T, Uitterlinden AG, Wie W-Q, Westendorp RGJ, Whitsel EA, Wiggins KL, Wilke RA, Ballantyne CM, Caulfield MJ, Colhoun HM, Cupples LA, Franco OH, Gudnason V, Hitman G, Palmer CAN, Psaty BM, Ridker PM, Stafford JM, Stein CM, Tardif J-C, Jukema JW, Rotter JI, Kraus RM. Meta-analysis of genome-wide association studies of HDL cholesterol response to statins. *J Med Genet.* 2016 Dec;53(12):835-845

*53. Evans DS, **Avery CL**, Nalls MA, Li G, Barnard J, Smith EN, Tanaka T, Butler AM, Keller MF, Buxbaum SG, Akylbekova E, Alonso A, Arking DE, Berenson GS, Bis JC, Buyske S, Carty CL, Chen W, Chung MK, Cummings SR, Deo R, Eaton CB, Fox ER, Heckbert SR, Heiss G, Hindorff LA, Hsueh W-C, Isaacs A, Jamshidi Y, Kerr KF, Liu F, Liu Y, Lohman KK, Maher JF, Mehra R, Meng YA, Musani SK, Newton-Cheh C, North KE, Psaty BM, Redline S, Rotter JI, Schork NJ, Shohet RV, Singleton AB, Smith JD, Soliman EZ, Srinivasan SR, Taylor HA, Van Wagener DR, Wilson JG, Yin X, Young T, Zhang Z-M, Zonderman AB, Evans MK, Ferrucci L, Murray SS, Tranah GJ, Whitsel EA, Reiner AP, Sotoodehnia N. Fine-mapping, Novel Loci Identification, and SNP Association Transferability in a Genome-Wide Association Study of QRS Duration in African Americans. *Hum Mol Genet* 2016 Oct 1;25(19):4350-4368

52. Fernández-Rhodes L, Graff M, Hodonsky CJ, Love SM, Howard AG, Seyerle AA, **Avery CL**, Chittoor G, Franceschini N, Voruganti VS, Young K, O'Connell J, North KE, Justice A. Comparison of Gene-Environment Methods using Simulated GAW19 Family-Based Data. *BMC Proc* 2016; 10(suppl 7); 371-77.

51. Justice AE, Howard A-G, Chittoor G, Fernandez-Rhodes L, Graff M, Voruganti VS, Diao G, Love S-A M, Franceschini N, O'Connell J, **Avery CL**, Young KL, North KE. Genome-Wide Association of SEM-Informed Latent Class Growth Modeling of Systolic Blood Pressure Change. *BMC Proc* 2016; 10(suppl 7): 321-27

50. **Avery CL**, Holliday KM, Chakladar S, Engeda JC, Hardy ST, Reis JP, Schreiner PJ, Shay CM, Daviglus ML, Lin DY, Heiss G, Zeng D. Disparities in early transitions to obesity in contemporary multi-ethnic U.S. population. *PLoS One* 2016; 11(6): e0158025

49. Noordam R. **Avery CL**, Visser LE, Stricker BH. Identifying genetic loci affecting drug response in depression using drug-gene interaction models: the challenges ahead of us. *Pharmacogenomics* 2016; 17(9): 1029-40

48. Petruski-Ivleva N, Viera A, Shimbo D, Munter P, **Avery CL**, Schneider ALC, Couper D, Kucharska-Newton A. Longitudinal patterns of change in systolic blood pressure and incidence of cardiovascular disease. The Atherosclerosis Risk in Communities (ARIC) study. *Hypertension* 2016; 67(6): 1150-6.

47. Busch EL, Keku TO, Richardson DB, Cohen SM, Eberhard DA, **Avery CL**, Sandler RS. Translational measures of markers of epithelial-mesenchymal transition and mortality in a population-based prospective cohort of colorectal cancer patients. *Clinical and Experimental Metastasis* 2016; 33:53-62

46. He Q, Zhang HH, **Avery CL**, Lin DY. Sparse meta-analysis with application to high-dimensional data. *Biostatistics* 2016; 17:205-20
45. Conomos MP, Laurie CA, Stilp AM, Gogarten SM, McHugh CP, Nelson SC, Sofer T, Fernández-Rhodes L, Justice AE, Graff M, Seyerle AA, **Avery CL**, Taylor KD, Rotter JI, Talavera GA, Daviglus ML, Wassertheil-Smoller S, Schneiderman N, Heiss G, Kaplan RC, Kerr KF, Browning SR, Browning BL, Weir BS, Avilés-Santa ML, Papanicolaou GJ, Lumley T, Szpiro AA, North KE, Rice K, Thornton TA, Laurie CC. Genetic Diversity and Association Studies in U.S. Hispanic/Latino Populations: Applications in the Hispanic Community Health Study/Study of Latinos. *Am J Hum Genet* 2016;98:165-84
44. Jones SB, Loehr LR, **Avery CL**, Gottesman R, Wruck L, Henderson L, Shahar E, Rosamond WD. Midlife Alcohol Consumption and the Risk of Stroke in the Atherosclerosis Risk in Communities Study. *Stroke* 2015; 46:3124-30
43. Bis J, Sitlani C, Irvin R, **Avery CL**, Smith AV, Sun F, Evans DS, Musani SK, Li X, Trompet S, Krijthe BP, Harris TB, Quibrera PM, Brody JA, Demissie S, Davis BR, Wiggins K, Tranah GJ, Lange LA, Sotoodehnia N, Stott DJ, Franco OH, Launuer LJ, Stürmer T, Taylor KD, Cupples LA, Eckfeldt J, Smith NL, Liu Y, Wilson JG, Heckbert S, Buckley BM, Ikram MA, Boerwinkle E, Chen YDI, de Craen AJM, Uitterlinden AG, Rotter JI, Ford I, Hofman A, Sattar N, Slagboom PE, Westendorp RGJ, Gudnason V, Vasani RS, Lumley T, Cummings SR, Taylor HA, Post W, Jukema JW, Strickler BH, Whitsel EA, Psaty BM, Arnett D. Drug-gene interactions of anti-hypertensive medications and risk of incident cardiovascular disease: a longitudinal pharmacogenomics study from the CHARGE consortium. *Plos One* 2015; 10(10):e0140496
- †42. Hardy ST, Loehr LR, Butler KR, Chang PP, Folsom AR, Chakladar S, Heiss G, MacLehose RF, Matsushita K, **Avery CL**. Reducing the blood pressure-related burden of cardiovascular disease: Impact of achievable improvements in blood pressure prevention and control. *J Am Heart Assoc* 2015;4(10):e002276
41. Folsom AR, Shah AM, Lutsey PL, Roetker NS, Alonso A, **Avery CL**, Miedema MD, Konety S, Chang PP, Solomon SD. American Heart Association's Life's Simple 7: Avoiding Heart Failure and Preserving Cardiac Structure and Function. *Am J Med* 2015; 128: 970-6
40. Sitlani CM, Rice KM, Lumley T, McKnight B, Cupples A, **Avery CL**, Whitsel EA, Noordam R, Psaty BM. GEE for GWAS using longitudinal phenotype data. *Statistics in Medicine* 2015; 34(1):118-30.
39. Postmus I, Trompet S, Deshmukh HA, Barnes MR, Li X, Warren HR, Chasman D, Zhou K, Arsenault BJ, Donnelly LA, Wiggins KL, **Avery CL**, Griffin P, Feng Q, Taylor KD, Li G, Evans DS, Smith AV, de Keyser CE, Johnson AD, de Craen AJ, Stott DJ, Buckley BM, Ford I, Westendorp RG, Slagboom PE, Sattar N, Munroe PB, Sever P, Poulter N, Stanton A, Shields DC, O'Brien E, Shaw-Hawkins S, Chen YD, Nickerson DA, Smith JD, Dubé MP, Boekholdt SM, Hovingh GK, Kastelein JJ, McKeigue PM, Betteridge J, Neil A, Durrington PN, Doney A, Carr F, Morris A, McCarthy MI, Groop L, Ahlqvist E; Welcome Trust Case Control Consortium, Bis JC, Rice K, Smith NL, Lumley T, Whitsel EA, Stürmer T, Boerwinkle E, Ngwa JS, O'Donnell CJ, Vasani RS, Wei WQ, Wilke RA, Liu CT, Sun F, Guo X, Heckbert SR, Post W, Sotoodehnia N,

Arnold AM, Stafford JM, Ding J, Herrington DM, Kritchevsky SB, Eiriksdottir G, Launer LJ, Harris TB, Chu AY, Giulianini F, MacFadyen JG, Barratt BJ, Nyberg F, Stricker BH, Uitterlinden AG, Hofman A, Rivadeneira F, Emilsson V, Franco OH, Ridker PM, Gudnason V, Liu Y, Denny JC, Ballantyne CM, Rotter JI, Adrienne Cupples L, Psaty BM, Palmer CAN, Tardif J-C, Colhoun HM, Hitman G, Krauss RM, Jukema JW, Caulfield MJ. Pharmacogenetic meta-analysis of genome-wide association studies of LDL cholesterol response to statins. *Nature Comm* 2014; Oct 28;5:5068

38. Caughey MC, Loehr LR, Cheng S, **Avery CL**, Solomon SD, Hinderliter AL. Associations between echocardiographic arterial compliance and incident cardiovascular disease in African Americans: The Atherosclerosis Risk in Communities Study. *Am J Hypertension* 2014; 114(12):1850-4

37. Caughey MC, **Avery CL**, Ni H, Astor B, Matsushita K, Wruck L, Rosamond W, Loehr LR. Outcomes of patients with anemia and acute decompensated heart failure with preserved versus reduced ejection fraction (from the ARIC study community surveillance). *American Journal of Cardiology* 2014; 114(12): 1850-1854

†36. Seyerle AA, Young AM, Jeff JM, Melton PE, Jorgensen NW, Lin Y, Carty C, Deelman E, Heckbert SR, Hindorff LA, Jackson RD, Martin LW, Perez MV, Psaty BM, Whitsel EA, North KE, Laston S, Kooperberg C, **Avery CL**. Evidence of heterogeneity in genetic determinants of QT interval across global populations: the Population Architecture using Genomics and Epidemiology (PAGE) Study. *Epidemiology* 2014; 25(6):790-8. PMID: PMC4380285

35. Harmon QE, Engel SM, Olshan AF, Wu MC, Moran T, Luo J, Stuebe A, **Avery CL**. Polymorphisms in inflammatory genes associated with Term SGA and Preeclampsia. *Am J Reprod Immunol* 2014; 71(5):472-84. PMID: PMC4040534

34. Cheng I, Kocarnik J, Dumitrescu L, Lindor N, Chang-Claude J, **Avery CL**, Love S-A, Slattery M, Chan A, Baron J, Hindorff L, Park S, Schumacher F, Hoffmeister M, Kraft P, Butler A, Junkins H, Duggan D, Hou L, Carlson C, Monroe K, Lin Y, Carty C, Mann S, Ma J, Giovannucci E, Jenkins M, Hopper J, Haile R, Conti D, Campbell P, Potter J, Caan B, Schoen R, Hayes R, Chanock S, Berndt S, Kury S, Bezieau S, Ambite JL, Kumaraguruparan G, Richardson D, Goodloe R, Dilks H, Baker P, Zanke B, Lemire M, Gallinger S, Hsu L, Jiao S, Harrison T, Seminara D, Haiman C, Kooperberg C, Wilkens L, Hutter C, White E, Crawford D, Heiss G, Hudson T, Brenner H, Bush W, Casey G, LeMarchand L, Peters U. Pleiotropic effects of genetic risk variants for other cancers on colorectal cancer risk: PAGE, GECCO, and CCFR Consortia. *Gut*, 2014; 63(5):800-7. PMID: PMC3918490

33. **Avery CL**, Sitlani C, Arking DE, Arnett DK, Bis J, Boerwinkle E, Buckley BM, Chen YDI, de Craen AJM, Enquohabarie D, Evans DS, Ford I, Garcia M, Harris TB, Heckbert S, Hochner H, Hsueh W-C, Jukema JW, Knekt P, Kristiansson K, Laaksonen M, Liu Y, Li X, MacFarlane P, Newton-Cheh C, Nieminen MS, Peloso GM, Porthan K, Rice K, Rotter JI, Sattar N, Siscovick D, Slagboom PE, Salomaa V, Sotoodehnia N, Stott DJ, Stürmer T, Trompet S, Westendorp RGJ, Whitsel EA, Psaty BM. Drug-gene interactions and the search for missing heritability: a cross-sectional pharmacogenomics study of the QT interval. *The Pharmacogenomics Journal* 2014; 14(1):6-13. PMID: PMC3766418

32. Wyss AB, Herring AH, **Avery CL**, Weissler MC, Bensen JT, Barnholtz-Sloan JS, Funkhouser WK, Olshan AF. Single-nucleotide Polymorphisms in Nucleotide Excision Repair Genes, Cancer Treatment, and Head and Neck Cancer Survival. *Cancer Causes and Control* 2014; 25(4): 437-50. PMID: PMC3766549
31. **Avery CL**, Der J, Whitsel EA, Stürmer T. Comparison of study designs used to detect and characterize pharmacogenomic interactions in non-experimental studies: a simulation study. *Pharmacogenetics and Genomics* 2014; 24(3):146-55. PMID: PMC3946643
30. Holliday KM, **Avery CL**, Poole C, McGraw KA, Williams R, Liao D, Smith RL, and Whitsel EA. Estimating Risk Associated with Meta-Analytically Informed Interpolation of Personal from Ambient Particulate Matter Exposures. *Epidemiology*, 2014; 25(1):35-43.
- †29. Seyerle AA and **CL Avery**. Understanding genetic epidemiology: The potential benefits and challenges of genetics for improving human health. [Invited Commentary]. *NC Med J*, 2013; 74(6): 505-8. PMID: PMC4041482
28. He Q, **Avery CL**, Lin D-Y. A general framework for association tests with multivariate traits in large-scale genomics studies. *Genetic Epidemiology*, 2013; 37(8): 759-97.
27. Harmon QE, Engel SM, Olshan AF, Moran T, Stuebe A, Luo J, Wu MC, **Avery CL**. Polymorphisms in Natural Kill Cell Related Genes are Associated with Preterm Birth. *American Journal of Epidemiology*, 2013;178(8):1208-1218. PMID: PMC3792727
26. Wyss AB, Herring AH, **Avery CL**, Weissler MC, Bensen JT, Barnholtz-Sloan JS, Funkhouser WK, Olshan AF. Single Nucleotide Polymorphisms in Nucleotide Excision Repair Genes, Cigarette Smoking, and the Risk of Head and Neck Cancer. *Cancer Epidemiology, Biomarkers & Prevention*, 2013;22(8):1428-1445.
- *25. Deo R, Nalls MA, **Avery CL**, Smith JG, Evans DS, Keller MF, Butler AM, Buxbaum SG, Quibrera M, Smith EN, Tanaka T, Akylbekova, E, Alonso A, Arking DE, Benjamin E, Berenson GS, Chen LY, Chen Y, Curb JD, Ellinor P, Evans MK, Ferrucci L, Fox E, Heckbert S, Heiss G, Hsueh W-C, Kerr K, Limacher MC, Liu Y, Lubitz S, Magnani J, Mehra R, Marcus GM, Murray SS, Newman AB, North KE, Paltoo DN, Reiner AP, Robinson JG, Samdarshi T, Schnabel RB, Schork NJ, Singleton AB, Soliman EZ, Sotoodehnia NS, Srinivasan SR, Taylor HA, Trevisan M, Zhang Z-M, Zonderman AB, Newton-Cheh C, Whitsel EA. *GJA1* is associated with heart rate in African Americans: a genome-wide association study of 13,372 participants. *Heart Rhythm* 2013;10(3):401-8. PMID: PMC3718037
24. Pendergrass SA, Brown-Gentry K, Dudek S, Frase A, Torstenson ES, Goodloe R, Ambite JL, **Avery CL**, Buyske S, Buzkova P, Deelman E, Fesinmeyer MD, Haiman C, Heiss G, Hindorff LA, Hsu C-N, Jackson RD, Koerber C, Le Marchand L, Park L, Lin Y, Matise TC, Wilkens LR, Monroe KR, Moreland L, Reiner A, Wallace R, Wilkens LR, Crawford DC, Ritchie MD. Phenotype-Wide Association Study (PheWAS) for Detection of Pleiotropy within the Multi-Ethnic Studies of the Population Architecture Using Genomics and Epidemiology (PAGE) Network. *PLoS Genetics* 2013; 9(1): e1003087. PMID: PMC3561060

†23. Butler AM, Yin X, Evans DS, Nalls MA, Smith EN, Tanaka T, Li G, Buxbaum SG, Whitsel EA, Alonso A, Arking DE, Benjamin EJ, Berenson GS, Bis JC, Chen W, Deo R, Ellinor PT, Heckbert SR, Heiss G, Hsueh W-C, Keating BJ, Kerr KF, Li Y, Limacher MC, Liu Y, Lubitz SA, Marcianti KD, Mehra R, Meng YA, Newman AB, Newton-Cheh C, North KE, Palmer CD, Psaty BM, Quibrera PM, Reiner AP, Rotter JI, Schnabel RB, Schork NJ, Singleton AB, Smith JG, Soliman EZ, Srinivasan SR, Zhang Z-M, Zonderman AB, Ferrucci L, Murray SS, Evans MK, Sotoodehnia N, Magnani JW, **Avery CL**. Novel Loci Associated with PR Interval in a Genome-wide Association Study of Ten African American Populations. *Circ Cardiovasc Genet* 2012;5(6):639-46. PMID: PMC3560365

*22. Smith JG, **Avery CL**, Evans DS, Nalls MA, Meng YA, Smith EN, Palmer C, Tanaka T, Mehra R, Butler AM, Young T, Buxbaum SG, Kerr KF, Berenson GS, Schnabel RB, Marcianti KD, Ellinor PT, Magnani JW, Chen W, Curb JD, Limacher MC, North KE, Reiner AP, Quibrera PM, Schork NJ, Singleton AB, Soliman EZ, Solomon AJ, Charkravarti A, Srinivasan SR, Alonso A, Wallace R, Redline S, Zhang Z-M, Post WS, Zonderman AB, Taylor HA, Murray SS, Ferrucci L, Arking DE, Evans MK, Fox ER, Sotoodehnia N, Heckbert SR, Whitsel EA, Newton-Cheh C. The impact of ancestry and common genetic variants on QT interval in African Americans. *Circ Cardiovasc Genet* 2012; 5(6):647-55. PMID: PMC3568265

21. **Avery CL**, Kucharska-Newton A, Monda KL, Sharrett AR, Mosley TH, Klein BH, Cotch MF, Wong TY, Klein R. Impact of long-term measures of glucose and blood pressure on retinal microvascular characteristics: The Atherosclerosis Risk in Communities (ARIC) Carotid MRI Study. *Atherosclerosis* 2012; 225:412-417. PMID: PMC3513355

20. **Avery CL**, Loehr LR, Baggett C, Chang PP, Kucharska-Newton A, Matsushita K, Rosamond WD, Heiss G. The population burden of heart failure attributable to modifiable risk factors: The Atherosclerosis Risk in Communities (ARIC) study. *Journal of the American College of Cardiology* 2012; 60(17): 1640-6. PMID: PMC3653309.

19. **Avery CL**, Sethupathy P, Buyske S, He QC, Lin DY, Arking DE, Carty CL, Duggan D, Fesinmeyer M, Hindorff LA, Jeff JM, Klein L, Patton K, Peters U, Shohet RV, Sotoodehnia N, Young AM, Kooperberg C, Haiman CA, Mohlke KL, Whitsel EA, North KE. Fine-mapping and initial characterization of QT interval loci in African Americans. *PLoS Genetics* 2012; 8(8): e1002870. PMID: PMC3415454

18. **Avery CL**, He QC, North KE, Ambite JL, Boerwinkle E, Fornage M, Hindorff LA, Kooperberg C, Meigs JB, Pankow JS, Pendergrass SA, Psaty BM, Ritchie MD, Rotter JI, Taylor KD, Wilkens LR, Heiss G, Lin DY. A Phenomics-Based Strategy Identifies Loci on *APOC1*, *BRAP*, and *PLCG1* Associated with Metabolic Syndrome Phenotype Domains. *PLoS Genet* 2011; 7(10): e1002322. PMID: PMC3192835

17. Pendergrass SA, Brown-Gentry K, Dudek S, Ambite JL, **Avery CL**, Buyske S, Cai C, Heiss G, Hindorff LA, Kooperberg C, Lin Y, Jackson RD, Wallace R, Reiner A, Moreland L, Manolio TA, Matisse TC, Wilkens LR, Fesinmeyer MD, Hsu CN, Crawford DC, Ritchie MD. Phenotype-Wide Association Study (PheWAS) for Exploration of Novel Genotype-Phenotype Relationships within the PAGE network. *Genetic Epidemiology* 2011; 35(5): 410-22. PMID: PMC3116446

16. Agarwal SK, **Avery CL**, Catellier D, Chang P, Rhodes CE, Shahar E, Ballantyne CM, Heiss G, Hoogeveen RC. Variability in the measurement of high sensitivity troponin T (hsTNT): The Atherosclerosis Risk in Communities (ARIC) Study. *Clinical Chemistry* 2011; 57(6): 891-7
15. Whitsel EA and **Avery CL**. The Environmental Epidemiology of Atrial Arrhythmogenesis: A Research Agenda. *Journal of Epidemiology and Community Health* 2010; 64(7): 587-90.
14. **Avery CL**, Mills KT, Williams R, McGraw KA, Poole C, Smith RL, Whitsel EA. Estimating error in using residential outdoor PM_{2.5} concentrations as proxies for personal exposures: a meta-analysis. *Environmental Health Perspectives* 2010; 118(5): 673-8. PMID: PMC2866684.
13. **Avery CL**, Mills KT, Chambless LE, Chang PP, Folsom AR, Mosley TH, Ni H, Rosamond WD, Wagenknecht L, Wood J, Heiss G. Long-term association of heart failure hospitalizations with self-reported signs and symptoms: The Atherosclerosis Risk in Communities (ARIC) Study. *European Journal of Heart Failure* 2010; 12(3): 232-8. PMID: PMC2821798.
12. **Avery CL**, Mills KT, Williams R, McGraw KA, Poole C, Smith RL, Whitsel EA. Estimating error in using ambient PM_{2.5} concentrations as proxies for personal exposures. *Epidemiology* 2010; 21(2): 215-223.
11. North KE, Franceschini N, **Avery CL**, Baird L, Leppert M, Chung J, Hanis C, Boerwinkle E, Volcik K, Grove-Caona M, Gu C, Heiss G, Pankow J, Couper D, Ballantyne C, Kao L, Weder A, Cooper R, Ehert G, Chakravarti A, Hunt SC. Variation in the checkpoint kinase 2 gene is associated with type 2 diabetes in multiple populations. *Acta Diabetologica* 2010; 47 (Suppl 1): 199-207. PMID: PMC2965317.
10. **Avery CL**, Monda KL, and North KE. Genetic association studies and the effect of misclassification and selection bias in putative confounders. *BMC Proceedings* 2009, 3(Suppl 7):S48. PMID: PMC2795947.
9. Whitsel EA and **Avery CL**. The association between long-term particulate matter exposure and cardiovascular disease morbidity. United States Environmental Protection Agency. Integrated Science Assessment for Particulate Matter (Second External Review Draft). U.S. EPA, Washington, DC, EPA/600/R-08/139-B; 2009. Available at: <http://cfpub.epa.gov/ncea/cfm/recordisplay.cfm?deid=210586>.
8. Martin LJ, Woo JG, **Avery CL**, Chen H, North KE. Multiple testing in the genomics era: Findings from Genetic Analysis Workshop 15, Group 15. *Genetic Epidemiology* 2007; 31(S1): S124-S131.
7. Lee CR, North KE, Bray MS, **Avery CL**, Mosher MJ, Couper DJ, Coresh J, Folsom AR, Boerwinkle E, Heiss G, Zeldin DC. Genetic variation in endothelial nitric oxide synthase (*NOS3*), cigarette smoking, and cardiovascular disease risk: an Atherosclerosis Risk in Communities (ARIC) study. *Pharmacogenetics and Genomics* 2006;16: 891-899. PMID: PMC1978174.
6. Zeng D, Lin DY, **Avery CL**, North KE, Bray MS. Efficient semiparametric estimation of haplotype-disease associations in case-cohort and nested case-control studies. *Biostatistics* 2006;7:486-502

5. **Avery CL**, Freedman BI, Kraja A, Borecki IB, Miller MB, Pankow JS, Arnett D, Lewis CE, Myers RH, Hunt SC, North KE. Genotype-by-sex interaction in the etiology of diabetes: Support for sex-specific QTLs in HyperGEN participants. *Diabetologia* 2006;49:2329-36.
4. **Avery CL**, Martin LJ, Williams JT, North KE. Accuracy of haplotype estimation in a region of low linkage disequilibrium. *BMC Genetics* 2005; 6(Suppl 1):S80. PMID: PMC1866700.
3. Martin LJ, **Avery CL**, Williams JT, North KE. Possible QTL-specific genotype by alcoholism interaction for evoked EEG oscillations. *BMC Genetics* 2005; 6(Suppl 1):S123. PMID: PMC1866814.
2. **Avery CL**, Freedman BI, Heiss G, Kraja A, Rice T, Arnett D, Miller MB, Pankow JS, Lewis CE, Myers RH, Hunt SC, Almasy L, North KE. Linkage analysis of diabetes status among hypertensive families: the HyperGEN Study. *Diabetes*. 2004;53:3307-12.
1. Chen C, Bharucha V, Chen Y, Westenbroek R, Brown A, Malhotra J, Jones D, **Avery C**, Gillespie P, Kazen-Gillespie K, Saunders T, Macdonald R, Catterall W, LL Isom. Reduced Sodium Channel Density, Altered Voltage Dependence of Inactivation, and Increased Susceptibility to Seizures in Mice Lacking Sodium Channel $\beta 2$ Subunits. *Proc Nat Acad Sci USA*. 2002;99:17072-7. PMID: PMC139271.

GRANTS

Current Research Support

Project # R01HL147853 Avery NHLBI 01/01/2020-12/31/2023 (NCE)
 Title: Inflammatory mediators of cardiometabolic risk in Latinos
 Description: This grant will measure eicosanoid metabolites in HCHS/SOL participants and examine their association with diabetes and cardiometabolic risk factors.
 Role: Principal investigator (20% effort)
 Total costs: \$3,875,000

Project #R01HL143885 Avery, Gordon-Larsen, North, Sumner (MPI) 04/01/2019-03/31/2023 (NCE)
 Title: Leveraging multi-omics approaches to examine metabolic challenges of obesity in relation to cardiovascular diseases
 Description: This study will test molecular mechanisms underlying obesity-associated CVD
 Role: Principal Investigator (with Penny Gordon-Larsen, Kari North, and Susan Sumner)
 Total costs: \$ 8,894,348

Project #R01HL142825 Avery NHLBI 02/10/2019-01/31/2024 (NCE)
 Title: Characterizing pleiotropy in cardiometabolic phenotypes among diverse populations
 Description: This study will examine the role of shared genetic effects in the etiology of cardiometabolic traits that span blood pressure, glucose, obesity, electrocardiographic, and lipid measurements.
 Role: Principal Investigator
 Total costs: \$3,231,039

Project #R01HG010297 Matisse, Gignoux (MPI) NHGRI 06/01/2019-05/31/2023
Title: PAGE III: Population Architecture using Genomics and Epidemiology
Description: This grant will test molecular mechanisms underlying established obesity genetic variants using metabolic markers via metabolomics.
Role: Co-investigator
Total costs: \$6,800,000

Project #R01HL151152 Kooperberg, Gignoux, North (MPI) NHLBI 08/2020-07/2024
Title: Polygenic Risk Scores for Diverse Populations - Bridging Research and Clinical Care
Description: We address major limitations in the science of polygenic risk scores (PRS) in multi-ethnic populations.
Role: Co-investigator
Total costs: \$9,617,000

Project #T32HL129982 Mohlke, North (MPD) NHLBI 05/01/16 – 04/31/26
Title: The Genetic Epidemiology of Heart, Lung, and Blood Traits Training Grant (GenHLB)
Description: Training grant in heart, lung, and blood genetic epidemiology for pre- and post-doctoral fellows in the University of North Carolina's Department of Epidemiology.
Role: HLB Genetic Epidemiology training dimension lead
Total costs: \$1,207,760

Pending Research Support

Project # R01HL172887 Avery, Graff (MPI) NHLBI 12/01/2024-11/31/2026
Title: Phenotypic consequences of lipoprotein(a) in diverse populations
Total costs: \$5,000,000

Project # Avery, Muntner (MPI) Amgen 06/01/24 – 05/30/25
Title: Development and validation of an algorithm to identify adults with elevated lipoprotein(a).
Total costs: \$250,000

Project # Welch (PI) NIA 12/01/2024-11/31/2026
Title: Inflammatory mediators of maternal preconception health and later-life CVD risk
Role: Subcontract PI
Total costs: \$300,000

Completed Research Support

Project #R01HL142825S1 Avery NHLBI 07/19/2019-01/31/2024
Title: Characterizing pleiotropy in cardiometabolic phenotypes among diverse populations-Admin Suppl
Description: This administrative supplement will support Ms. Sofia Dimos's training in the genetic epidemiology of cardiovascular diseases.
Role: Principal Investigator and mentor
Total costs: \$102,012

Project #2011561720 Avery Amgen 02/01/2019-12/31/2021
Title: Phenome-wide association study of Lp(a) variants
Description: This study will assess the wide-ranging phenotypic consequences of genetically inferred Lp(a)
Role: Principal investigator
Total costs: \$ 411,530

Project #R03HD097630 Avery, Harris (MPI) NICHD 09/20/2018-08/31/20
Title: Add Health GWAS Data: User Support and Research Tools Enable Widespread Access
Description: The study outlines a plan to develop tools to enable widespread access to and use of genome-wide association study (GWAS) data among respondents in the National Longitudinal Study of Adolescent to Adult Health (Add Health).
Role: Principal Investigator (with Kathleen Mullan Harris)
Total cost: \$155,500

Project #U01HG007416 North (PI) NHGRI 07/01/13- 05/31/19
Title: Genetic epidemiology of causal variants across the life course (CALiCo) Phase II
Description: The Genetic Epidemiology of Causal Variants Across the Life Course (CALiCo) is a member study of Population Architecture using Genomics and Epidemiology (PAGE). PAGE was established to investigate the epidemiologic architecture of mature genetic variants associated with complex diseases.
Role: Co-investigator
Total costs: \$3,057,956

Project # R01HD73342 Harris (PI) NICHD 07/01/12-03/01/19
Title: Add Health: A Genotype and Phenotype Resource for the Global Scientific Community
Description: This proposal will complete genome-wide genotyping on the remaining 6,000 Add Health samples using current Illumina chip technology, prepare genotype and phenotype data and documentation for dissemination, and conduct GWAS
Role: Co-Investigator
Total costs: \$2,864,536

Project # R01HL103612 Psaty (PI) NHLBI 08/10/11-07/31/17 (NCE)
Title: Prospective meta-analyses of drug-gene interactions: CHARGE GWAS consortium
Description: The investigators will use GWAS methods to identify potential drug-gene interactions related to unintended drug effects and to identify genetic variants that may modify the intended effects of treatments for lipids and blood pressure on change in risk factor levels, carotid atherosclerosis and incidence of cardiovascular events.
Role: Co-Investigator
Total costs: \$439,755

Project # 15GPSPG239 Avery (PI) AHA 02/1/2015- 01/31/2017
Title: Pharmacogenomics of risk factors for cardiac arrhythmias in global populations
Description: This study will evaluate common SNPs influencing arrhythmogenic adverse drug reactions in European American, African American, and Hispanics/Latino populations.
Role: Principal Investigator

Direct costs: \$500,000

Project # R21HL121580 Avery (PI) NHLBI 09/15/13- 07/31/16 (NCE)

Title: The natural history of cardiovascular health in U.S. populations

Description: This study will characterize the stages and age epochs at which African American, Hispanic/Latino, American Indian, and Caucasian U.S. populations transition between ideal, intermediate, and poor levels of cardiovascular health metrics.

Role: Principal Investigator

Total costs: \$210,486

Project #PPHF 2014 Hoerger (PI) CDC 10/01/2014-09/30/16

Title: Developing Online Toolkits to Evaluate Diabetes Burdens and Impacts of Diabetes
Preventions for States

Description: This project will compute and report diabetes-attributable morbidity, mortality, and economic costs as well as health and economic impacts of diabetes prevention programs for use by U.S. states

Role: Consultant

Project # R00HL098458 Avery (PI) NHLBI 08/15/11-05/31/15

Title: Enumerating the community burden of heart failure

Description: This study is designed to facilitate enumeration of heart failure in outpatient settings, accomplished by developing and calibrating simple but accurate heart failure classification tools in outpatient settings and applying these tools in the Atherosclerosis Risk in Communities Study.

Role: Principal Investigator

Total costs: \$450,000

Project # R01ES017794 Whitsel (PI) NIEHS 08/11/10-05/30/15

Title: Modification of PM-mediated arrhythmogenesis in populations

Description: This application will investigate susceptibility to the arrhythmogenic effects of particulate matter (PM) air pollution contributed by common genetic variation in the Women's Health Initiative Clinical Trial and Atherosclerosis Risk in Communities studies.

Role: Co-investigator

Total costs: \$2,289,037

Project # A10-0347-001 North (PI) NHLBI 08/01/2009-06/01/2011

Title: GWAS Analysis for the Women's Health Initiative Study

Description: The goal of this study is to detect, map, and identify polymorphic genes that influence variation in complex phenotypes in women.

Role: Co-investigator

Total costs: \$708,762

Project: # U01HG004803 North (PI) NHGRI 06/01/11-05/31/14

Title: Genetic epidemiology of causal variants across the life course (CALiCo)

Description: The Genetic Epidemiology of Causal Variants Across the Life Course (CALiCo) is a member study of Population Architecture using Genomics and Epidemiology (PAGE). PAGE was established to investigate the epidemiologic architecture of mature genetic variants associated with

complex diseases.
Role: Co-investigator
Total costs: \$3,275,184

Project: # AS405 Avery (PI) WHI SE-RC RFA 11/01/12-9/30/13
Title: Pharmacogenomics of risk factors for cardiac arrhythmias in global populations
Description: Pilot study to include Women’s Health Initiative Clinical Trial participants in a ten-study consortium examining pharmacogenomic influences on ECG traits.
Role: Principal Investigator
Total costs: \$20,000

Project # K99HL098458 Avery (PI) NHLBI 04/01/10–08/14/11
Title: Enumerating the community burden of heart failure
Description: This study is designed to facilitate enumeration of heart failure in outpatient settings, accomplished by developing and calibrating simple but accurate classification tools in outpatient settings and applying these tools in the Atherosclerosis Risk in Communities Study cohort.
Role: Principal Investigator
Total costs: \$200,000

Project # 0615310U Avery (PI) AHA 07/01/06–08/30/07
Title: Tobacco exposure, DNA repair genes, and atherosclerosis: the Atherosclerosis Risk in Communities (ARIC) study
Description: This project evaluated how common DNA repair gene variants, in combination with exposure to tobacco smoke, modified the risk of atherosclerosis and its clinical sequelae.
Role: Principal Investigator
Total costs: \$25,950

TEACHING RECORD

University of North Carolina at Chapel Hill, Department of Epidemiology

Lead instructor EPID 716: Epidemiologic Analysis of Binary Data, 2013-22, 24-
Number of students: 33-69

EPID 889: Readings in CVD Epidemiology, 2021, 2022, 2023
Number of students: 20

EPID 889: OMICs of HLB traits, 2019
Number of students: 19

EPID 889: Population Burden Measures in Epidemiology, 2012-16
Number of students: 10-15

Developer MPH@UNC online course: Epidemiologic Analysis of Binary Data
(2021-22)

Co-instructor EPID 890: Perspectives in Epidemiology and Public Health, 2014-15
Number of students: 10

 EPID 715: Theory and Quantitative Methods in Epidemiology, 2012
Number of students: 45

Invited lecturer EPID 889: Genetic epidemiology data analysis, 2021
Number of students: 18

 EPID 735: Cardiovascular Disease Epidemiology, 2009-11, 15-
Number of students: ~15

 EPID 710: Fundamentals of Epidemiology, 2014, 16
Number of students: ~45

 EPID 743: Genetic Epidemiology, 2009-13
Number of students: ~15

 EPID 810: Design of Clinical Research, 2012
Number of students: ~10

 PATH 797: Molecular and Cellular Biology of CVD, 2013, 15, 16
Number of students: ~5

University of Ulm, Germany (Summer Institute)

Lead instructor Epidemiologic Analysis of Binary Data, 2018
Number of students: 14

 Genetic Epidemiology, 2011
Number of students: 30

POSTDOCTORAL MENTORING

Current: 1) Moa Park Lee (2018-)
 2) Mohanraj Krishnan (2021-)

Completed:
 1) Harold Lee (2021-2)
 2) Raul Mendez Giraldez (2015-17)

GRADUATE STUDENT MENTORING

Current:
Dissertation chair:
 1) Epidemiology: Moa Park Lee (expected graduation: 05/2024)
 2) Epidemiology: Sarah Koenigsberg (expected graduation: 08/2026)
 3) Epidemiology: Emily Drzymalla (expected graduation: 08/2027)

4) Epidemiology: Katherine Connors (expected graduation: 08/2028)

Dissertation committee member:

- 1) Epidemiology: Leah Sadinski
- 2) Epidemiology: Alvin Thomas
- 3) Sociology: Adam Lilly
- 4) Nutrition: Autumn Hullings

Completed:

Dissertation chair (year graduated):

- 1) Epidemiology: Carolina Downie (2023)
- 2) Epidemiology: Brooke Staley (2023)
- 3) Epidemiology: Chani Hodonsky (2019)
- 4) Epidemiology: Joseph Engeda (2019)
- 5) Epidemiology: Amanda Seyerle (2016)

Masters paper chair (year graduated):

- 1) Epidemiology: Anna Ballou (2023)
- 2) Epidemiology: Antoine Baldassari (2018)
- 3) Epidemiology: Amanda Seyerle (2013)

Dissertation committee member (year graduated):

- 1) Nutrition: Kyle Busse (2024)
- 2) Biostatistics: Yilun Li (2024)
- 3) Biostatistics: Yianchang Li (2024)
- 4) Epidemiology: Yifei Lu (2023)
- 5) Nutrition: Caitlyn Lowery (2023)
- 6) Nutrition: Sarah Frank (2023)
- 7) Epidemiology: Victoria Buchanan (2023)
- 8) Epidemiology: Poojan Shrestha (2023)
- 9) Epidemiology: Daeun Kim (2023)
- 10) Biostatistics: Richard Sizelove (2023)
- 11) Epidemiology: Audrey Renson (2021)
- 12) Nutrition: Natalia Rebolledo Fuentealba (2021)
- 13) Nutrition: Yiqing Wang (2020)
- 14) Nutrition: Tania Aburto Soto (2019)
- 15) Epidemiology: Erline Miller (2018)
- 16) Epidemiology: Shakia Hardy (2017)
- 17) Epidemiology: Aderonke Akinkugbe (2017)
- 18) Epidemiology: Anna Bauer (2017)
- 19) Health Policy: Jamie Jarmul (2017)
- 20) Epidemiology: Kapuaola Gellert (2016)
- 21) Epidemiology: Evan Busch (2015)
- 22) Epidemiology: Sara Jones (2015)
- 23) Epidemiology: Annah Wyss (2013)
- 24) Epidemiology: Quaker Harmon (2012)

25) Biostatistics: Qianchuan (Chad) He (2012)

Masters paper second reader (year graduated):

1) Epidemiology: Katelyn Holliday (2011)

UNDERGRADUATE STUDENT MENTORING

Current: None

Completed:

- 1) Allison Baumgartner (2022-2023)
- 2) Sofia Dimos (2018-2021)
- 3) Ashlyn Stackhouse (2017-2018)

UNDERGRADUATE, PREDOCTORAL, AND POSTDOCTORAL AWARDS*

**Restricted to awards where I am the primary mentor/senior author*

- 1) Moa Park Lee (2020-21)
Recipient, F32HL149256, "A Phenome-Wide Association Study to Identify and Characterize Phenotypic Effects of Lp(a)"
- 2) Sofia Dimos (2019-21)
Recipient, R01HL142825S1, Research Supplement to Promote Diversity in Health-Related Research
- 3) Katelyn Holliday (2014)
Recipient, Steven Blair Award for Excellence in Physical Activity Research, 2014 AHA EPI/Lifestyle conference
- 4) Shakia Hardy (2014)
Finalist, Jeremiah and Rose Stamler Research Award for New Investigators, 2014 AHA EPI/Lifestyle conference

PROFESSIONAL SERVICE

Invited Grant Reviews

2021-	NIH NHLBI Mentored Transition to Independence Review Committee permanent member
2019-21	NIH NHLBI Mentored Transition to Independence Review Committee ad hoc reviewer
2017, 21	NIH CHS study section ad hoc reviewer
2020	05 ZRG1 PSE-E (02) S Chronic Diseases and Epidemiology ad hoc reviewer
2018	Mid-Atlantic NORC Pilot and Feasibility Grant ad hoc reviewer
2018	AHA Uncovering New Patterns ad hoc reviewer
2016	NIH/NHLBI K01 special emphasis panel ad hoc reviewer
2015	AHA Behavioral Science - Prevention/Intervention ad hoc reviewer
2014	NIH CASE/EPIC study section (ZRG1 PSEQ 90) ad hoc reviewer
2012	Center for Environmental Health and Susceptibility (CEHS), University of

North Carolina at Chapel Hill

Leadership

- 2020- Leader, UNC Cardiovascular Epidemiology Program Area (50 faculty, staff, and students)
- 2018- Coleader, UNC CVD Genetic Epidemiology computational lab (25 faculty, staff, and students)

Scientific and Advisory Committees/Councils

- 2023- Amgen, the Center for Observational Research (CfOR) Methods Council
- 2021-4 American Heart Association EPI Statistics Committee of the Council on Epidemiology and Prevention (Nominated member)
- 2012 Circulatory Device Advisory Committee panelist, U.S. Food and Drug Administration, Silver Spring, MD

University and Study Committees

- 2019,- University of North Carolina Postdoctoral Award for Research Excellence Selection Committee
- 2014- Atherosclerosis Risk in Communities Ancillary Studies
- 2013- HCHS/SOL ECG genomics working group (chair)
- 2012- Women's Health Initiative Publications & Presentations
- 2011- University of North Carolina Department of Epidemiology Graduate Studies

Editorial

Associate Editor
Heart (2022-)

Invited Manuscript Review

American Journal of Epidemiology (2015 – 1)
American Journal of Preventive Medicine (2016 -1)
American Heart Association abstract review (2012, 2016-)
Annals of Epidemiology (2018-1)
Circulation (2012 -1, 2013 – 2, 2014-2, 2015-1, 2019-1, 2020-1, 2021-1)
Circulation: Genomics and Precision Medicine (2012 -1, 2021-1)
Clinical Genetics (2017-1)
Current Pharmacogenomics and Personalized Medicine (2017-1)
Diabetes (2015 – 1)
Diabetes Care (2020 – 1)
Diabetes Research and Clinical Practice (2012 – 1)
Diabetic Research (2013 – 1)
Environmental Health Perspectives (2013- 2)
Epidemiology (2015 – 2; 2017-1)
Genetic Epidemiology (2012 -1, 2020 - 1)
Heart (2013 – 1, 2014-1)
JAMA Neurology (2020-1)
Journal of Internal Medicine (2018 -1)

North Carolina Medical Journal (2013, 1)
Pharmacoepidemiology and Drug Safety (2015, 1)
Pharmacy and Therapeutics (2014, 1)
PLoS Medicine (2020-1)
PLoS One (2013 – 1, 2014-1)
Scientific Reports (2016 -1)
 Society for Epidemiologic Research abstract review (2011, 2012)

Conference Contribution

2021-22 AHA Epidemiology|Lifestyle Organizing Committee
 2021 Chair, SER session “So much more than GWAS: How genetics can strengthen causal inference”
 2011 Co-chair, North American Congress of Epidemiology symposium: “What role does the epidemiologist play in genetic epidemiology?”

MEMBERSHIPS AND FELLOWSHIPS

2019- Member, Society for Epidemiologic Research
 2015- Fellow of the American Heart Association (FAHA)
 2003- Member, American Heart Association

INVITED SPEAKER

2023 Human Health Exposure Analysis Resource (HHEAR) Exposomics Webinar
 2022 Columbia University: The Exposome Boot Camp: Measuring Exposures on an Omic Scale
 2022 NIH/NHLBI: Epidemiology Pan-T32 Conference Series
 2022 University of Southern California Center for Genetic Epidemiology: Seminar Series
 2021 University of North Carolina Center for Pharmacoepidemiology: Methods Symposium
 2019 University of Alabama: CCVC 8th Annual Symposium New Horizons in CVD: A focus on Precision and Personalized Medicine
 2019 NIH/NHLBI: K to R01 Investigators meeting
 2019 Society for Epidemiology Research (SER)
 2015 North Carolina State University: PopMED Forum
 2012 University of North Carolina: Cardiovascular Grand Rounds

DIVERSITY, EQUITY, AND INCLUSION

2006- Research awards that prioritized inclusion of diverse populations
 2018 Mentored undergraduate diversity supplement to R01HL142825
 2014, 19 Ad hoc reviewer: NHLBI MOSAIC and K01 awards to promote diversity
 2019 Redesigned EPID 716 to integrate content measuring race and racism
 2019 Organized Departmental and CVD program area anti-racism trainings