

# Stanford

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## Thomas Quertermous, MD

William G. Irwin Professor in Cardiovascular Medicine  
Medicine - Cardiovascular Medicine

### CONTACT INFORMATION

- **Alternate Contact**

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### Bio

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#### ACADEMIC APPOINTMENTS

- Professor, Medicine - Cardiovascular Medicine
- Member, Bio-X
- Member, Cardiovascular Institute
- Member, Maternal & Child Health Research Institute (MCHRI)

#### LINKS

- Quertermous Lab homepage: <http://quertermous.stanford.edu>

### Research & Scholarship

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#### CURRENT RESEARCH AND SCHOLARLY INTERESTS

The Quertermous laboratory employs two basic approaches of study to better understand the genetic basis of atherosclerotic heart disease. One approach uses basic molecular biology methodology, primarily working with cellular and genetic mouse models, and is focused on the recently identified apelin-APJ pathway. A second approach employs the power of modern human genetics. Informative cohorts have been collected that allow investigation of risk factors such as hypertension and insulin resistance as well as coronary heart disease. Initial studies have employed the candidate gene approach, and more recently whole genome association studies, to identify allelic variation that is associated with risk factor and disease susceptibility.

#### CLINICAL TRIALS

- Permission to Collect Blood Over Time for Research, Not Recruiting

### Teaching

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#### STANFORD ADVISEES

**Doctoral Dissertation Reader (AC)**

Stephen Chang

**Postdoctoral Faculty Sponsor**

Laeya Abdoli Najmi, Paul Cheng, Hyun-Jung Kim, Jiehan Li, Huitong Shi, Stanislao Travisano, Quanyi Zhao

**Postdoctoral Research Mentor**

Paul Cheng, Huitong Shi, Stanislao Travisano, Quanyi Zhao

**Publications**

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**PUBLICATIONS**

- **Coronary Disease Associated Gene TCF21 Inhibits Smooth Muscle Cell Differentiation by Blocking the Myocardin-Serum Response Factor Pathway.** *Circulation research*  
Nagao, M., Lyu, Q., Zhao, Q., Wirka, R. C., Bagga, J., Nguyen, T., Cheng, P., Kim, J. B., Pjanic, M., Miano, J. M., Quertermous, T.  
2019
- **Atheroprotective roles of smooth muscle cell phenotypic modulation and the TCF21 disease gene as revealed by single-cell analysis.** *Nature medicine*  
Wirka, R. C., Wagh, D., Paik, D. T., Pjanic, M., Nguyen, T., Miller, C. L., Kundu, R., Nagao, M., Collier, J., Koyano, T. K., Fong, R., Woo, Y. J., Liu, et al  
2019
- **Stanford Cardiovascular Institute At the Forefront of Cardiovascular Research** *CIRCULATION RESEARCH*  
Wu, J. C., Woo, Y., Mayerle, M., Harrington, R. A., Quertermous, T.  
2019; 124 (10): 1420–24
- **TCF21 and AP-1 interact through epigenetic modifications to regulate coronary artery disease gene expression** *GENOME MEDICINE*  
Zhao, Q., Wirka, R., Trieu Nguyen, Nagao, M., Cheng, P., Miller, C. L., Kim, J., Pjanic, M., Quertermous, T.  
2019; 11
- **Stanford Cardiovascular Institute.** *Circulation research*  
Wu, J. C., Woo, Y. J., Mayerle, M., Harrington, R. A., Quertermous, T.  
2019; 124 (10): 1420–24
- **Advances in Transcriptomics: Investigating Cardiovascular Disease at Unprecedented Resolution.** *Circulation research*  
Wirka, R. C., Pjanic, M., Quertermous, T.  
2018; 122 (9): 1200–1220
- **Genetic Regulatory Mechanisms of Smooth Muscle Cells Map to Coronary Artery Disease Risk Loci.** *American journal of human genetics*  
Liu, B., Pjanic, M., Wang, T., Nguyen, T., Gloudemans, M., Rao, A., Castano, V. G., Nurnberg, S., Rader, D. J., Elwyn, S., Ingelsson, E., Montgomery, S. B., Miller, et al  
2018
- **Coronary artery disease genes SMAD3 and TCF21 promote opposing interactive genetic programs that regulate smooth muscle cell differentiation and disease risk.** *PLoS genetics*  
Iyer, D., Zhao, Q., Wirka, R., Naravane, A., Nguyen, T., Liu, B., Nagao, M., Cheng, P., Miller, C. L., Kim, J. B., Pjanic, M., Quertermous, T.  
2018; 14 (10): e1007681
- **Circulating peptide prevents preeclampsia** *SCIENCE*  
Wirka, R. C., Quertermous, T.  
2017; 357 (6352): 643–44
- **TCF21 and the environmental sensor aryl-hydrocarbon receptor cooperate to activate a pro-inflammatory gene expression program in coronary artery smooth muscle cells.** *PLoS genetics*  
Kim, J. B., Pjanic, M., Nguyen, T., Miller, C. L., Iyer, D., Liu, B., Wang, T., Sazonova, O., Carcamo-Orive, I., Matic, L. P., Maegdefessel, L., Hedin, U., Quertermous, et al  
2017; 13 (5)
- **Analysis of Transcriptional Variability in a Large Human iPSC Library Reveals Genetic and Non-genetic Determinants of Heterogeneity** *CELL STEM CELL*  
Carcamo-Orive, I., Hoffman, G. E., Cundiff, P., Beckmann, N. D., D'Souza, S. L., Knowles, J. W., Patel, A., Papatsenko, D., Abbasi, F., Reaven, G. M., Whalen, S., Lee, P., Shahbazi, et al

2017; 20 (4): 518-?

- **Enhancer connectome in primary human cells identifies target genes of disease-associated DNA elements.** *Nature genetics*  
Mumbach, M. R., Satpathy, A. T., Boyle, E. A., Dai, C., Gowen, B. G., Cho, S. W., Nguyen, M. L., Rubin, A. J., Granja, J. M., Kazane, K. R., Wei, Y., Nguyen, T., Greenside, et al  
2017
- **Identification of new susceptibility loci for type 2 diabetes and shared etiological pathways with coronary heart disease.** *Nature genetics*  
Zhao, W., Rasheed, A., Tikkanen, E., Lee, J. J., Butterworth, A. S., Howson, J. M., Assimes, T. L., Chowdhury, R., Orho-Melander, M., Damrauer, S., Small, A., Asma, S., Imamura, et al  
2017
- **Analysis of Transcriptional Variability in a Large Human iPSC Library Reveals Genetic and Non-genetic Determinants of Heterogeneity.** *Cell stem cell*  
Carcamo-Orive, I., Hoffman, G. E., Cundiff, P., Beckmann, N. D., D'Souza, S. L., Knowles, J. W., Patel, A., Papatsenko, D., Abbasi, F., Reaven, G. M., Whalen, S., Lee, P., Shahbazi, et al  
2016
- **Nat1 Deficiency Is Associated with Mitochondrial Dysfunction and Exercise Intolerance in Mice** *CELL REPORTS*  
Chennamsetty, I., Coronado, M., Contrepolis, K., Keller, M. P., Carcamo-Orive, I., Sandin, J., Fajardo, G., Whittle, A. J., Fathzadeh, M., Snyder, M., Reaven, G., Attie, A. D., Bernstein, et al  
2016; 17 (2): 527-540
- **Genetics and Genomics of Coronary Artery Disease.** *Current cardiology reports*  
Pjanic, M., Miller, C. L., Wirka, R., Kim, J. B., Direnzo, D. M., Quertermous, T.  
2016; 18 (10): 102-?
- **Epigenetic response to environmental stress: Assembly of BRG1-G9a/GLP-DNMT3 repressive chromatin complex on Myh6 promoter in pathologically stressed hearts** *BIOCHIMICA ET BIOPHYSICA ACTA-MOLECULAR CELL RESEARCH*  
Han, P., Li, W., Yang, J., Shang, C., Lin, C., Cheng, W., Hang, C. T., Cheng, H., Chen, C., Wong, J., Xiong, Y., Zhao, M., Drakos, et al  
2016; 1863 (7): 1772-1781
- **Integrative functional genomics identifies regulatory mechanisms at coronary artery disease loci.** *Nature communications*  
Miller, C. L., Pjanic, M., Wang, T., Nguyen, T., Cohain, A., Lee, J. D., Perisic, L., Hedin, U., Kundu, R. K., Majmudar, D., Kim, J. B., Wang, O., Betsholtz, et al  
2016; 7: 12092-?
- **Coronary Artery Disease and Its Risk Factors: Leveraging Shared Genetics to Discover Novel Biology.** *Circulation research*  
Quertermous, T., Ingelsson, E.  
2016; 118 (1): 14-16
- **Coronary Artery Disease Associated Transcription Factor TCF21 Regulates Smooth Muscle Precursor Cells that Contribute to the Fibrous Cap.** *Genomics data*  
Nurnberg, S. T., Cheng, K., Raiesdana, A., Kundu, R., MILLER, C. L., Kim, J. B., Arora, K., Carcamo-Orive, I., Xiong, Y., Tellakula, N., Nanda, V., Murthy, N., Boisvert, et al  
2015; 5: 36-37
- **Characterization of TCF21 Downstream Target Regions Identifies a Transcriptional Network Linking Multiple Independent Coronary Artery Disease Loci** *PLOS GENETICS*  
Sazonova, O., Zhao, Y., Nuernberg, S., Miller, C., Pjanic, M., Castano, V. G., Kim, J. B., Salfati, E. L., Kundaje, A. B., Bejerano, G., Assimes, T., Yang, X., Quertermous, et al  
2015; 11 (5)
- **Coronary Artery Disease Associated Transcription Factor TCF21 Regulates Smooth Muscle Precursor Cells That Contribute to the Fibrous Cap** *PLOS GENETICS*  
Nurnberg, S. T., Cheng, K., Raiesdana, A., Kundu, R., Miller, C. L., Kim, J. B., Arora, K., Carcamo-Orive, I., Xiong, Y., Tellakula, N., Nanda, V., Murthy, N., Boisvert, et al  
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- **Coronary Artery Disease Associated Transcription Factor TCF21 Regulates Smooth Muscle Precursor Cells That Contribute to the Fibrous Cap.** *PLoS genetics*  
Nurnberg, S. T., Cheng, K., Raiesdana, A., Kundu, R., Miller, C. L., Kim, J. B., Arora, K., Carcamo-Orive, I., Xiong, Y., Tellakula, N., Nanda, V., Murthy, N., Boisvert, et al  
2015; 11 (5)

- **Identification and validation of N-acetyltransferase 2 as an insulin sensitivity gene** *JOURNAL OF CLINICAL INVESTIGATION*  
Knowles, J. W., Xie, W., Zhang, Z., Chennemsetty, I., Assimes, T. L., Paananen, J., Hansson, O., Pankow, J., Goodarzi, M. O., Carcamo-Orive, I., Morris, A. P., Chen, Y. I., Maekinen, et al  
2015; 125 (4): 1739-1751
- **A long noncoding RNA protects the heart from pathological hypertrophy.** *Nature*  
Han, P., Li, W., Lin, C., Yang, J., Shang, C., Nurnberg, S. T., Jin, K. K., Xu, W., Lin, C., Lin, C., Xiong, Y., Chien, H., Zhou, et al  
2014; 514 (7520): 102-106
- **Dissecting the causal genetic mechanisms of coronary heart disease.** *Current atherosclerosis reports*  
Miller, C. L., Assimes, T. L., Montgomery, S. B., Quertermous, T.  
2014; 16 (5): 406-?
- **Clinical interpretation and implications of whole-genome sequencing.** *JAMA*  
Dewey, F. E., Grove, M. E., Pan, C., Goldstein, B. A., Bernstein, J. A., Chaib, H., Merker, J. D., Goldfeder, R. L., Enns, G. M., David, S. P., Pakdaman, N., Ormond, K. E., Caleshu, et al  
2014; 311 (10): 1035-1045
- **Coronary heart disease-associated variation in TCF21 disrupts a miR-224 binding site and miRNA-mediated regulation.** *PLoS genetics*  
Miller, C. L., Haas, U., Diaz, R., Leeper, N. J., Kundu, R. K., Patlolla, B., Assimes, T. L., Kaiser, F. J., Perisic, L., Hedin, U., Maegdefessel, L., Schunkert, H., Erdmann, et al  
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- **Disease-Related Growth Factor and Embryonic Signaling Pathways Modulate an Enhancer of TCF21 Expression at the 6q23.2 Coronary Heart Disease Locus** *PLOS GENETICS*  
Miller, C. L., Anderson, D. R., Kundu, R. K., Raiesdana, A., Nuernberg, S. T., Diaz, R., Cheng, K., Leeper, N. J., Chen, C., Chang, I., Schadt, E. E., Hsiung, C. A., Assimes, et al  
2013; 9 (7)
- **Loss of CDKN2B Promotes p53-Dependent Smooth Muscle Cell Apoptosis and Aneurysm Formation** *ARTERIOSCLEROSIS THROMBOSIS AND VASCULAR BIOLOGY*  
Leeper, N. J., Raiesdana, A., Kojima, Y., Kundu, R. K., Cheng, H., Maegdefessel, L., Toh, R., Ahn, G., Ali, Z. A., Anderson, D. R., Miller, C. L., Roberts, S. C., Spin, et al  
2013; 33 (1): E1-?
- **CDKN2B Regulates Cell Fate Decisions in Human Vascular Smooth Muscle Cells**  
Leeper, N. J., Raiesdana, A., Cheng, H., Kundu, R. K., Kojima, Y., Cheng, K., Schadt, E., Quertermous, T.  
LIPPINCOTT WILLIAMS & WILKINS.2010
- **Hundreds of variants clustered in genomic loci and biological pathways affect human height** *NATURE*  
Allen, H. L., Estrada, K., Lettre, G., Berndt, S. I., Weedon, M. N., Rivadeneira, F., Willer, C. J., Jackson, A. U., Vedantam, S., Raychaudhuri, S., Ferreira, T., Wood, A. R., Weyant, et al  
2010; 467 (7317): 832-838
- **Apelin signaling antagonizes Ang II effects in mouse models of atherosclerosis** *JOURNAL OF CLINICAL INVESTIGATION*  
Chun, H. J., Ali, Z. A., Kojima, Y., Kundu, R. K., Sheikh, A. Y., Agrawal, R., Zheng, L., Leeper, N. J., Pearl, N. E., Patterson, A. J., Anderson, J. P., Tsao, P. S., Lenardo, et al  
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- **PCSK6 Is a Key Protease in the Control of Smooth Muscle Cell Function in Vascular Remodeling.** *Circulation research*  
Rykaczewska, U., Suur, B. E., Rohl, S., Razuvaev, A., Lengquist, M., Sabater-Lleal, M., van der Laan, S. W., Miller, C. L., Wirka, R. C., Kronqvist, M., Gonzalez Diez, M., Vesterlund, M., Gillgren, et al  
2020
- **FAM13A affects body fat distribution and adipocyte function.** *Nature communications*  
Fathzadeh, M., Li, J., Rao, A., Cook, N., Chennamsetty, I., Seldin, M., Zhou, X., Sangwung, P., Gloudemans, M. J., Keller, M., Attie, A., Yang, J., Wabitsch, et al  
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- **Genomic profiling of human vascular cells identifies TWIST1 as a causal gene for common vascular diseases.** *PLoS genetics*  
Nurnberg, S. T., Guerraty, M. A., Wirka, R. C., Rao, H. S., Pjanic, M., Norton, S., Serrano, F., Perisic, L., Elwyn, S., Pluta, J., Zhao, W., Testa, S., Park, et al  
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- **Effect of Common Genetic Variants of Growth Arrest-Specific 6 Gene on Insulin Resistance, Obesity and Type 2 Diabetes in an Asian Population.** *PLoS one*  
Hsieh, C. H., Chung, R. H., Lee, W. J., Lin, M. W., Chuang, L. M., Quertermous, T., Assimes, T., Hung, Y. J., Yu, Y. W.  
; 10 (8): e0135681
- **Adiponectin Receptor 3 is Associated With Endothelial Nitric Oxide Synthase Dysfunction and Predicts Insulin Resistance in South Asians**  
Chandy, M., Sayed, N., Lau, E., Liu, C., Wei Tzu-Tang, Chen, I. Y., Thomas, D., Rhee, J., Oh, B., Pepic, L., Husain, M., Quertermous, T., Nallamshetty, S., et al  
LIPPINCOTT WILLIAMS & WILKINS.2019
- **The role of insulin as a key regulator of seeding, proliferation, and mRNA transcription of human pluripotent stem cells.** *Stem cell research & therapy*  
Shahbazi, M., Cundiff, P., Zhou, W., Lee, P., Patel, A., D'Souza, S. L., Abbasi, F., Quertermous, T., Knowles, J. W.  
2019; 10 (1): 228
- **IGF1 gene is associated with triglyceride levels in subjects with family history of hypertension from the SAPHIRE and TWB projects**  
Wang, W., Chiu, Y., Chung, R., Hwu, C., Lee, I., Lee, C., Chang, Y., Hung, K., Quertermous, T., Chen, Y. I., Hsiung, C. A.  
NATURE PUBLISHING GROUP.2019: 163
- **Opportunities and challenges for transcriptome-wide association studies** *NATURE GENETICS*  
Wainberg, M., Sinnott-Armstrong, N., Mancuso, N., Barbeira, A. N., Knowles, D. A., Golan, D., Ermel, R., Ruusalepp, A., Quertermous, T., Hao, K., Björkegren, J. M., Im, H., Pasaniuc, et al  
2019; 51 (4): 592–99
- **TCF21 and AP-1 interact through epigenetic modifications to regulate coronary artery disease gene expression.** *Genome medicine*  
Zhao, Q., Wirka, R., Nguyen, T., Nagao, M., Cheng, P., Miller, C. L., Kim, J. B., Pjanic, M., Quertermous, T.  
2019; 11 (1): 23
- **CRISPR-Cas9-mediated knockout of SPRY2 in human hepatocytes leads to increased glucose uptake and lipid droplet accumulation.** *BMC endocrine disorders*  
Cook, N. L., Pjanic, M., Emmerich, A. G., Rao, A. S., Hetty, S., Knowles, J. W., Quertermous, T., Castillejo-López, C., Ingelsson, E.  
2019; 19 (1): 115
- **Detailed Functional Characterization of a Waist-Hip Ratio Locus in 7p15.2 Defines an Enhancer Controlling Adipocyte Differentiation.** *iScience*  
Castillejo-Lopez, C., Pjanic, M., Pirona, A. C., Hetty, S., Wabitsch, M., Wadelius, C., Quertermous, T., Arner, E., Ingelsson, E.  
2019; 20: 42–59
- **Opportunities and challenges for transcriptome-wide association studies.** *Nature genetics*  
Wainberg, M., Sinnott-Armstrong, N., Mancuso, N., Barbeira, A. N., Knowles, D. A., Golan, D., Ermel, R., Ruusalepp, A., Quertermous, T., Hao, K., Björkegren, J. L., Im, H. K., Pasaniuc, et al  
2019; 51 (4): 592–99
- **Genetic variation of SORBS1 gene is associated with glucose homeostasis and age at onset of diabetes: A SAPHIRE Cohort Study** *SCIENTIFIC REPORTS*  
Chang, T., Wang, W., Hsiung, C. A., He, C., Lin, M., Sheu, W., Chang, Y., Quertermous, T., Chen, Y., Rotter, J. I., Chuang, L., SAPHIRE Study Grp  
2018; 8: 10574
- **Large-Scale Single-Cell RNA-Seq Reveals Molecular Signatures of Heterogeneous Populations of Human Induced Pluripotent Stem Cell-Derived Endothelial Cells.** *Circulation research*  
Paik, D. T., Tian, L., Lee, J., Sayed, N., Chen, I. Y., Rhee, S., Rhee, J., Kim, Y., Wirka, R. C., Buikema, J. W., Wu, S. M., Red-Horse, K., Quertermous, et al  
2018
- **Apelin and APJ orchestrate complex tissue-specific control of cardiomyocyte hypertrophy and contractility in the hypertrophy-heart failure transition.** *American journal of physiology. Heart and circulatory physiology*  
Parikh, V. N., Liu, J., Shang, C., Woods, C., Chang, A. C., Zhao, M., Charo, D. N., Grunwald, Z., Huang, Y., Seo, K., Tsao, P. S., Bernstein, D., Ruiz-Lozano, et al  
2018
- **Functional Assays to Screen and Dissect Genomic Hits: Doubling Down on the National Investment in Genomic Research.** *Circulation. Genomic and precision medicine*  
Musunuru, K., Bernstein, D., Cole, F. S., Khokha, M. K., Lee, F. S., Lin, S., McDonald, T. V., Moskowitz, I. P., Quertermous, T., Sankaran, V. G., Schwartz, D. A., Silverman, E. K., Zhou, et al  
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- **IGF1 Gene Is Associated With Triglyceride Levels In Subjects With Family History Of Hypertension From The SAPHIRE And TWB Projects** *INTERNATIONAL JOURNAL OF MEDICAL SCIENCES*  
Wang, W., Chiu, Y., Chung, R., Hwu, C., Lee, I., Lee, C., Chang, Y., Hung, K., Quertermous, T., Chen, Y., Hsiung, C. A.  
2018; 15 (10): 1035–42
- **Functional regulatory mechanism of smooth muscle cell-restricted LMOD1 coronary artery disease locus.** *PLoS genetics*  
Nanda, V., Wang, T., Pjanic, M., Liu, B., Nguyen, T., Matic, L. P., Hedin, U., Koplev, S., Ma, L., Franzén, O., Ruusalepp, A., Schadt, E. E., Björkegren, et al  
2018; 14 (11): e1007755
- **Load-dependent effects of apelin on murine cardiomyocytes** *PROGRESS IN BIOPHYSICS & MOLECULAR BIOLOGY*  
Peyronnet, R., Bollensdorff, C., Capel, R. A., Rog-Zielinska, E. A., Woods, C. E., Charo, D. N., Lookin, O., Fajardo, G., Ho, M., Quertermous, T., Ashley, E. A., Kohl, P.  
2017; 130: 333–43
- **CRP-level-associated polymorphism rs1205 within the CRP gene is associated with 2-hour glucose level: The SAPHIRE study** *SCIENTIFIC REPORTS*  
Sheu, W., Wang, W., Wu, K., He, C., Hwu, C., Quertermous, T., Hsieh, W., Lee, W., Ting, C., Chen, Y. I., Hsiung, C. A.  
2017; 7: 7987
- **Genome-wide copy number variation analysis identified deletions in SFMBT1 associated with fasting plasma glucose in a Han Chinese population** *BMC GENOMICS*  
Chung, R., Chiu, Y., Hung, Y., Lee, W., Wu, K., Chen, H., Lin, M., Chen, Y. I., Quertermous, T., Hsiung, C. A.  
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- **Trans-ethnic fine-mapping of genetic loci for body mass index in the diverse ancestral populations of the Population Architecture using Genomics and Epidemiology (PAGE) Study reveals evidence for multiple signals at established loci** *HUMAN GENETICS*  
Fernandez-Rhodes, L., Gong, J., Haessler, J., Franceschini, N., Graff, M., Nishimura, K. K., Wang, Y., Highland, H. M., Yoneyama, S., Bush, W. S., Goodloe, R., Ritchie, M. D., Crawford, et al  
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- **Fifteen new risk loci for coronary artery disease highlight arterial-wall-specific mechanisms.** *Nature genetics*  
Howson, J. M., Zhao, W., Barnes, D. R., Ho, W., Young, R., Paul, D. S., Waite, L. L., Freitag, D. F., Fauman, E. B., Salfati, E. L., Sun, B. B., Eicher, J. D., Johnson, et al  
2017
- **Association analyses of East Asian individuals and trans-ancestry analyses with European individuals reveal new loci associated with cholesterol and triglyceride levels** *HUMAN MOLECULAR GENETICS*  
Spracklen, C. N., Chen, P., Kim, Y. J., Wang, X., Cai, H., Li, S., Long, J., Wu, Y., Wang, Y. X., Takeuchi, F., Wu, J., Jung, K., Hu, et al  
2017; 26 (9): 1770-1784
- **Alternative Progenitor Cells Compensate to Rebuild the Coronary Vasculature in Elabela- and Apj-Deficient Hearts.** *Developmental cell*  
Sharma, B., Ho, L., Ford, G. H., Chen, H. I., Goldstone, A. B., Woo, Y. J., Quertermous, T., Reversade, B., Red-Horse, K.  
2017
- **Induced Pluripotent Stem Cell-Derived Endothelial Cells in Insulin Resistance and Metabolic Syndrome.** *Arteriosclerosis, thrombosis, and vascular biology*  
Carcamo-Orive, I., Huang, N. F., Quertermous, T., Knowles, J. W.  
2017; 37 (11): 2038–42
- **Endothelial APLNR regulates tissue fatty acid uptake and is essential for apelin's glucose-lowering effects.** *Science translational medicine*  
Hwangbo, C., Wu, J., Papangelis, I., Adachi, T., Sharma, B., Park, S., Zhao, L., Ju, H., Go, G. W., Cui, G., Inayathullah, M., Job, J. K., Rajadas, et al  
2017; 9 (407)
- **Fine-mapping of lipid regions in global populations discovers ethnic-specific signals and refines previously identified lipid loci** *HUMAN MOLECULAR GENETICS*  
Zubair, N., Graff, M., Ambite, J. L., Bush, W. S., Kichaev, G., Lu, Y., Manichaikul, A., Sheu, W. H., Absher, D., Assimes, T. L., Bielinski, S. J., Bottinger, E. P., Buzkova, et al  
2016; 25 (24): 5500-5512
- **Targeting LOXL2 for cardiac interstitial fibrosis and heart failure treatment** *NATURE COMMUNICATIONS*  
Yang, J., Savvatis, K., Kang, J. S., Fan, P., Zhong, H., Schwartz, K., Barry, V., Mikels-Vigdal, A., Karpinski, S., Korniyev, D., Adamkewicz, J., Feng, X., Zhou, et al  
2016; 7

- **Transcriptomic Profiling Maps Anatomically Patterned Subpopulations among Single Embryonic Cardiac Cells** *DEVELOPMENTAL CELL*  
Li, G., Xu, A., Sim, S., Priest, J. R., Tian, X., Khan, T., Quertermous, T., Zhou, B., Tsao, P. S., Quake, S. R., Wu, S. M.  
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Priest, J. R., Gawad, C., Kahlig, K. M., Yu, J. K., O'Hara, T., Boyle, P. M., Rajamani, S., Clark, M. J., Garcia, S. T., Ceresnak, S., Harris, J., Boyle, S., Dewey, et al  
2016; 113 (41): 11555-11560
- **The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals** *NATURE GENETICS*  
Ehret, G. B., Ferreira, T., Chasman, D. I., Jackson, A. U., Schmidt, E. M., Johnson, T., Thorleifsson, G., Luan, J., Donnelly, L. A., Kanoni, S., Petersen, A. -, Pihurl, V., Strawbridge, et al  
2016; 48 (10): 1171-1184
- **The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals.** *Nature genetics*  
Ehret, G. B., Ferreira, T., Chasman, D. I., Jackson, A. U., Schmidt, E. M., Johnson, T., Thorleifsson, G., Luan, J., Donnelly, L. A., Kanoni, S., Petersen, A., Pihur, V., Strawbridge, et al  
2016; 48 (10): 1171-1184
- **Pathological Ace2-to-Ace enzyme switch in the stressed heart is transcriptionally controlled by the endothelial Brg1-FoxM1 complex** *PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE UNITED STATES OF AMERICA*  
Yang, J., Feng, X., Zhou, Q., Cheng, W., Shang, C., Han, P., Lin, C., Chen, H. V., Quertermous, T., Chang, C.  
2016; 113 (38): E5628-E5635
- **CD47-blocking antibodies restore phagocytosis and prevent atherosclerosis.** *Nature*  
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