

Stanford



Philip S. Tsao, PhD

Professor (Research) of Medicine (Cardiovascular Medicine)
Medicine - Cardiovascular Medicine

Bio

ACADEMIC APPOINTMENTS

- Professor (Research), Medicine - Cardiovascular Medicine
- Member, Bio-X
- Member, Cardiovascular Institute
- Member, Wu Tsai Human Performance Alliance
- Member, Maternal & Child Health Research Institute (MCHRI)
- Member, Wu Tsai Neurosciences Institute

ADMINISTRATIVE APPOINTMENTS

- Executive Committee, Stanford Cardiovascular Institute, (2010- present)
- Member, Stanford Diabetes Research Center, (2018- present)
- Co-Director, Cardiovascular Pulmonary Sciences Application, (2005-2018)

HONORS AND AWARDS

- Department of Medicine Teaching Award, Stanford (2003)
- Fellow, Arteriosclerosis, Thrombosis, and Vascular Biology Council of the American Heart Association (2003)
- Established Investigator Award, American Heart Association (2008)

PROFESSIONAL EDUCATION

- PhD, Thomas Jefferson University , Cardiovascular Physiology (1991)

Research & Scholarship

CURRENT RESEARCH AND SCHOLARLY INTERESTS

Our primary interests are in understanding the molecular underpinnings of vascular disease as well as assessing disease risk. We use a wide range of biochemical, molecular and physiological techniques to make primary observations in cell systems as well as preclinical models. Furthermore, we continue to extend our findings to human subjects in order to confirm their clinical applicability. Current research projects include:

Mechanisms regulating atherosclerosis and abdominal aortic aneurysm disease: While single genes can have dramatic effects in cellular biology, it is becoming increasingly clear that vascular disease (and health) is regulated by the coordinated expression of gene cassettes or pathways. By monitoring expression patterns of the entire genome simultaneously, we can begin to identify networks of genes that work in concert to affect disease progression.

Moreover, this approach can often implicate specific nexus genes that are at the center of larger networks and/or participate in multiple pathways. Additionally, we are investigating the role microRNAs, a newly discovered class of small RNA molecules, in orchestrating the activity of multiple genes during the course of disease.

Role of insulin resistance: Reduced activity of the endogenous hormone, insulin, is now recognized as a cardinal feature of type 2 diabetes and an independent risk factor for cardiovascular disease. We have investigated the effects of insulin resistance in several tissues and have recently focused our attention on adipose tissue biology and how it relates to CVD. Long known as a storage vehicle for excess calories, the fat cell is now recognized to be a factory of different products that can not only affect local activity, but can circulate in the blood as hormones and regulate many biological processes. For example, we have recently reported that the novel hormone, apelin, is produced by fat tissue and has important effects upon insulin resistance, obesity and diabetes, all of which have significant implications for cardiovascular disease.

Biomarkers for risk assessment: In addition to target identification, we are applying transcriptional profiling and pathway analysis for another important aspect of cardiovascular disease management—biomarker discovery. As the name connotes, a biomarker should be a good indication of the disease state and thereby allow for early detection as well as monitoring disease progression and, hopefully, efficacy of an applied therapy. Biomarkers can encompass a wide range of molecules including DNA variants, RNA, proteins, as well as lipids. They can even encompass modalities such as molecular imaging. We are engaged in not only identifying novel biomarkers for cardiovascular disease, but also in producing algorithms that combine multiple biomarkers to optimally assess risk.

CLINICAL TRIALS

- Predicting DVT Risk in GI Cancer Patients Using Plasma Biomarkers, Not Recruiting

Teaching

STANFORD ADVISEES

Postdoctoral Faculty Sponsor

Yuri Kono, Agnesa Mazrekaj

GRADUATE AND FELLOWSHIP PROGRAM AFFILIATIONS

- Bioengineering (Phd Program)
- Cardiovascular Medicine (Fellowship Program)
- Medicine (Masters Program)
- Molecular and Genetic Medicine (Fellowship Program)

Publications

PUBLICATIONS

- **Genome-wide association meta-analysis identifies risk loci for abdominal aortic aneurysm and highlights PCSK9 as a therapeutic target** *NATURE GENETICS*
Roychowdhury, T., Klarin, D., Levin, M. G., Spin, J. M., Rhee, Y., Deng, A., Headley, C. A., Tsao, N. L., Gellatly, C., Zuber, V., Shen, F., Hornsby, W. E., Laursen, et al
2023
- **Genome-wide association meta-analysis identifies risk loci for abdominal aortic aneurysm and highlights PCSK9 as a therapeutic target.** *Nature genetics*
Roychowdhury, T., Klarin, D., Levin, M. G., Spin, J. M., Rhee, Y. H., Deng, A., Headley, C. A., Tsao, N. L., Gellatly, C., Zuber, V., Shen, F., Hornsby, W. E., Laursen, et al

2023

- **E-cigarette exposure augments murine abdominal aortic aneurysm development: role of Chil1.** *Cardiovascular research*
Mulorz, J., Spin, J. M., Mulorz, P., Wagenhauser, M., Deng, A., Mattern, K., Rhee, Y. H., Toyama, K., Adam, M., Schelzig, H., Maegdefessel, L., Tsao, P. S.
2022
- **The power of genetic diversity in genome-wide association studies of lipids.** *Nature*
Graham, S. E., Clarke, S. L., Wu, K. H., Kanoni, S., Zajac, G. J., Ramdas, S., Surakka, I., Ntalla, I., Vedantam, S., Winkler, T. W., Locke, A. E., Marouli, E., Hwang, et al
2021
- **Genetic Architecture of Abdominal Aortic Aneurysm in the Million Veteran Program.** *Circulation*
Klarin, D. n., Verma, S. S., Judy, R. n., Dikilitas, O. n., Wolford, B. N., Paranjpe, I. n., Levin, M. G., Pan, C. n., Tcheandjieu, C. n., Spin, J. M., Lynch, J. n., Assimes, T. L., Nyrønning, et al
2020
- **Genome-wide association study of peripheral artery disease in the Million Veteran Program.** *Nature medicine*
Klarin, D., Lynch, J., Aragam, K., Chaffin, M., Assimes, T. L., Huang, J., Lee, K. M., Shao, Q., Huffman, J. E., Natarajan, P., Arya, S., Small, A., Sun, et al
2019
- **Decoding the Genomics of Abdominal Aortic Aneurysm.** *Cell*
Li, J., Pan, C., Zhang, S., Spin, J. M., Deng, A., Leung, L. L., Dalman, R. L., Tsao, P. S., Snyder, M.
2018; 174 (6): 1361
- **Segmental Aortic Stiffening Contributes to Experimental Abdominal Aortic Aneurysm Development** *CIRCULATION*
Raaz, U., Zoellner, A. M., Schellinger, I. N., Toh, R., Nakagami, F., Brandt, M., Emrich, F. C., Kayama, Y., Eken, S., Adam, M., Maegdefessel, L., Hertel, T., Deng, et al
2015; 131 (20): 1783-1795
- **miR-24 limits aortic vascular inflammation and murine abdominal aneurysm development** *NATURE COMMUNICATIONS*
Maegdefessel, L., Spin, J. M., Raaz, U., Eken, S. M., Toh, R., Azuma, J., Adam, M., Nagakami, F., Heymann, H. M., Chernugobova, E., Jin, H., Roy, J., Hultgren, et al
2014; 5
- **Pathogenesis of Abdominal Aortic Aneurysms: MicroRNAs, Proteases, Genetic Associations.** *Annual review of medicine*
Maegdefessel, L., Dalman, R. L., Tsao, P. S.
2014; 65: 49-62
- **MicroRNA-21 Blocks Abdominal Aortic Aneurysm Development and Nicotine-Augmented Expansion** *SCIENCE TRANSLATIONAL MEDICINE*
Maegdefessel, L., Azuma, J., Toh, R., Deng, A., Merk, D. R., Raiesdana, A., Leeper, N. J., Raaz, U., Schoelmerich, A. M., McConnell, M. V., Dalman, R. L., Spin, J. M., Tsao, et al
2012; 4 (122)
- **Inhibition of microRNA-29b reduces murine abdominal aortic aneurysm development** *JOURNAL OF CLINICAL INVESTIGATION*
Maegdefessel, L., Azuma, J., Toh, R., Merk, D. R., Deng, A., Chin, J. T., Raaz, U., Schoelmerich, A. M., Raiesdana, A., Leeper, N. J., McConnell, M. V., Dalman, R. L., Spin, et al
2012; 122 (2): 497-506
- **A narrative review of recent literature of circulating biomarkers of abdominal aortic aneurysm.** *JVS-vascular science*
Rhee, Y. H., Spin, J. M., Tsao, P. S.
2026; 7: 100399
- **Carriage of rare APOB variants predisposes to severe steatotic liver disease and hepatocellular carcinoma.** *The Journal of clinical investigation*
Mureddu, M., Pelusi, S., Jamialahmadi, O., Vujkovic, M., Miano, L., Eidgah Torghabehei, H., Ronzoni, L., Malvestiti, F., Saracino, M., Periti, G., Moretti, V., Teerlink, C. C., Lynch, et al
2026

- **Genetic and Social Determinants of Renin-Angiotensin-Aldosterone System Inhibitor-Induced Angioedema: A Precision Medicine Health Equity Study.** *medRxiv : the preprint server for health sciences*
Toda, N., Haldar, T., Teerlink, C. C., Hu, D., Danilov, P., Huntsman, S., Lu, M., Tsao, P. S., Tcheandjieu, C., Iribarren, C., Bress, A., Lynch, J. A., Ziv, et al
2026
- **Harnessing iPSCs, 3D organoids, and multiomics to model rare vascular diseases: Emerging new approach methodologies.** *Vascular medicine (London, England)*
Liu, L., Wu, D., Tsao, P. S., Leeper, N. J., Sayed, N.
2026: 1358863X251394285
- **Opportunistic Genomic Screening for Familial Hypercholesterolemia to Improve Low-Density Lipoprotein Cholesterol: A Randomized Clinical Trial.** *JAMA network open*
Vassy, J. L., Brunette, C. A., Yi, T., Assimes, T. L., Christensen, K. D., Knowles, J. W., Sturm, A. C., Sun, Y. V., Alexander, N., Cardellino, M. P., Harrison, A., Gerety, H. L., Pyatt, et al
2026; 9 (1): e2549664
- **Leveraging Large-Scale Biobanks for Therapeutic Target Discovery.** *HGG advances*
Ferolito, B. R., Dashti, H., Giambartolomei, C., Peloso, G. M., Golden, D. J., Gravel-Pucillo, K., Rasooly, D., Horimoto, A. R., Matty, R., Gaziano, L., Liu, Y., Smit, I. A., Zdravil, et al
2025: 100556
- **Stem cell-based therapies for treatment of abdominal aortic aneurysm: development, application, and future potential.** *npj biomedical innovations*
Badawy, S., Anand, S., Marini, A. X., Mulorz, J., Tsao, P. S., Huang, N. F.
2025; 2 (1): 41
- **Multipopulation Genome-Wide Association Study Identifies Novel Loci for Bicuspid Aortic Valve and Reveals Shared Genetic Architecture With Aortopathies.** *Circulation. Genomic and precision medicine*
Bermudez, F., Shakt, G., Bowles, A., Alba, P., Dinatale, T., Chang, K. M., Tsao, P. S., Lynch, J., Dochtermann, D., Pyarajan, S., Small, A., Levin, M. G., Damrauer, et al
2025: e005491
- **Abstract 4367273: Epigenetic Architecture of Abdominal Aortic Aneurysm and Its Augmented Risk in Smokers: Insights from Methylome-Genome Integration**
Yuan, S., Shakt, G., Hartmann, K., Levin, M., Tsao, P., Voight, B., Jones, G., Damrauer, S.
LIPPINCOTT WILLIAMS & WILKINS.2025
- **Genomic Insights into Incretin-Based Therapeutics for Peripheral Arterial Disease**
Sorondo, S., Cabot, J., Tsao, P., Klarin, D., Leeper, N., Adkar, S.
LIPPINCOTT WILLIAMS & WILKINS.2025
- **GREGoR: accelerating genomics for rare diseases.** *Nature*
Dawood, M., Heavner, B., Wheeler, M. M., Ungar, R. A., LoTempio, J., Wiel, L., Berger, S., Bernstein, J. A., Chong, J. X., Délot, E. C., Eichler, E. E., Lupski, J. R., Shojaie, et al
2025; 647 (8089): 331-342
- **Clinical validation of a statin-benefit polygenic score using real-world cohorts of primary prevention participants.** *medRxiv : the preprint server for health sciences*
Haldar, T., Lee, K. M., Teerlink, C. C., Iyer, K. R., Voorhees, A. L., Najera, J. A., Toda, N., Lu, M., Douglas, M. P., Popejoy, A. B., Bress, A., Tcheandjieu, C., Tsao, et al
2025
- **Mendelian Randomization Suggests a Causal Link Between Glycemic Traits and Thoracic Aortic Structures and Diseases.** *JACC. Basic to translational science*
Daria, T., Iyer, K., Alkhalro, H., Kho, P. F., Suzuki, K., Hatzikotoulas, K., Southam, L., Taylor, H. J., Yin, X., Mandla, R., Huerta-Chagoya, A., Rayner, W. N., Levin, et al
2025: 101390
- **Polygenic Susceptibility in Peripartum, Alcohol-Induced, and Cancer Therapy-Related Cardiomyopathies.** *JAMA cardiology*
Maamari, D. J., Biddinger, K. J., Jurgens, S. J., Rämö, J. T., Gaziano, L., Zheng, A., Challa, S. P., Hayes, D., Gongora, C. A., Choi, S. H., Chang, K. M., Tsao, P. S., Arany, et al

2025

- **Polygenic risk score for type 2 diabetes shows context-dependent effects across populations.** *Nature communications*
Guo, B., Cai, Y., Kim, D., Smit, R. A., Wang, Z., Iyer, K. R., Hilliard, A. T., Haessler, J., Tao, R., Broadaway, K. A., Wang, Y., Pozdeyev, N., Stæger, et al
2025; 16 (1): 8632
- **Germline Variants Influence Chronic Liver Disease Progression through Distinct Pathways.** *medRxiv : the preprint server for health sciences*
Vujkovic, M., Kaplan, D. E., Ghouse, J., Loza, B. L., Brancale, J., Lewis, A., Zhang, D. Y., Levin, M. G., Veatch, O. J., Johnson, J. P., Schneider, C. V., Verma, A., Wangenstein, et al
2025
- **Single-cell polygenic risk scores dissect cellular and molecular heterogeneity of complex human diseases.** *Nature biotechnology*
Zhang, S., Shu, H., Zhou, J., Rubin-Sigler, J., Yang, X., Liu, Y., Cooper-Knock, J., Monte, E., Zhu, C., Tu, S., Li, H., Tong, M., Ecker, et al
2025
- **Multi-ancestry polygenic risk scores for the prediction of type 2 diabetes and complications in diverse ancestries.** *medRxiv : the preprint server for health sciences*
Huerta-Chagoya, A., Kim, J., Mandla, R., Lu, Y., Suzuki, K., Petty, L. E., Ng, H. K., Choi, J., Lee, S., Rout, M., Lin, K., Adair, L. S., Adeyemo, et al
2025
- **Circulating Protein Mediators Linking Genetically Predicted Smoking to Abdominal Aortic Aneurysm: A Genomic-Proteomic Analysis.** *Arteriosclerosis, thrombosis, and vascular biology*
Yuan, S., Khodursky, S., Geng, J., Sharma, P., Spin, J. M., Tsao, P., Levin, M. G., Damrauer, S. M.
2025
- **A genome-wide association study of hidradenitis suppurativa from the VA's Million Veteran Program.** *medRxiv : the preprint server for health sciences*
Wendland, Z., Teerlink, C. C., Pridgen, K. M., Lo, S., Sayed, C., Van Straalen, K. R., Tcheandjieu, C., Tsao, P. S., Chang, K. M., Li, Y., Mohlke, K. L., Sun, Q., Lynch, et al
2025
- **MicroRNA-15a-5p mediates abdominal aortic aneurysm progression and serves as a potential diagnostic and prognostic circulating biomarker.** *Communications medicine*
Winski, G., Chernogubova, E., Busch, A., Eken, S. M., Jin, H., Lindquist Liljeqvist, M., Khan, T., Bäcklund, A., Paloschi, V., Roy, J., Hultgren, R., Brostjan, C., de Borst, et al
2025; 5 (1): 218
- **CYP2C19 Polymorphisms and Clinical Outcomes Following Percutaneous Coronary Intervention in the Million Veteran Program.** *Clinical pharmacology and therapeutics*
Chanfreau-Coffinier, C., Friede, K. A., Plomondon, M. E., Lee, K. M., Lu, Z., Dinatale, T., DuVall, S. L., Vassy, J. L., Waldo, S. W., Cleator, J. H., Maddox, T. M., Rader, D. J., Assimes, et al
2025
- **Genome-wide Association Study of Lower Extremity Aneurysms Reveals Shared Genetic Architecture With Abdominal Aortic Aneurysms**
Cabot, J., Sorondo, S., Adkar, S., Klarin, D., Tsao, P., Levin, M., Damrauer, S.
MOSBY-ELSEVIER.2025
- **Plasma proteome-wide Mendelian randomization reveals the association of extracellular matrix proteins with abdominal aortic aneurysm.** *JVS-vascular science*
Khodursky, S., Yuan, S., Spin, J. M., Tsao, P. S., Levin, M. G., Damrauer, S. M.
2025; 6: 100290
- **Plasma proteomic signatures for type 2 diabetes and related traits in the UK Biobank cohort.** *Diabetes research and clinical practice*
Gupte, T. P., Azizi, Z., Kho, P. F., Zhou, J., Nzenkue, K., Chen, M. L., Panyard, D. J., Guarischi-Sousa, R., Hilliard, A. T., Sharma, D., Watson, K., Abbasi, F., Tsao, et al
2025: 112194
- **Dissecting the Genetic Architecture of Intracranial Aneurysms.** *Circulation. Genomic and precision medicine*
Adkar, S. S., Lynch, J., Choi, R. B., Roychowdhury, T., Judy, R. L., Paruchuri, K., Go, D. C., Bamezai, S., Cabot, J., Sorondo, S., Levin, M. G., Milewicz, D. M., Willer, et al
2025: e004626

- **Heterogeneous effects of genetic variants and traits associated with fasting insulin on cardiometabolic outcomes.** *Nature communications*
Sevilla-González, M., Smith, K., Wang, N., Jensen, A. E., Litkowski, E. M., Kim, H., DiCorpo, D. A., Hsu, S., Cui, J., Liu, C. T., Yu, C., McNeil, J. J., Lacaze, et al
2025; 16 (1): 2569
- **CXCL12 drives natural variation in coronary artery anatomy across diverse populations.** *Cell*
Rios Coronado, P. E., Zhou, J., Fan, X., Zanetti, D., Naftaly, J. A., Prabala, P., Martínez Jaimes, A. M., Farah, E. N., Kundu, S., Deshpande, S. S., Evergreen, I., Kho, P. F., Ma, et al
2025
- **Identifying Circulating Protein Mediators in the Link Between Smoking and Abdominal Aortic Aneurysm: An Integrated Analysis of Human Proteomic and Genomic Data.** *medRxiv : the preprint server for health sciences*
Yuan, S., Khodursky, S., Geng, J., Sharma, P., Spin, J. M., Tsao, P., Levin, M. G., Damrauer, S. M.
2025
- **Investigation of Genomic and Transcriptomic Risk Factors of Clopidogrel Response in African Americans.** *Clinical pharmacology and therapeutics*
Yang, G., Alarcon, C., Chanfreau, C., Lee, N. H., Friedman, P., Nutescu, E., Tuck, M., O'Brien, T., Gong, L., Klein, T. E., Chang, K. M., Tsao, P. S., Meltzer, et al
2025
- **Plasma proteome-wide Mendelian randomization reveals the association of extracellular matrix proteins with abdominal aortic aneurysm** *JVS-VASCULAR SCIENCE*
Khodursky, S., Yuan, S., Spin, J. M., Tsao, P. S., Levin, M. G., Damrauer, S. M.
2025; 6
- **Critical genes in genitourinary embryogenesis are related to the development of adult hydrocele.** *Scientific reports*
Roberson, J. L., Neylan, C. J., Judy, R., Walker, V., Tsao, P. S., Damrauer, S. M., Maguire, L. H.
2024; 14 (1): 30314
- **Publisher Correction: Genome-wide association study reveals mechanisms underlying dilated cardiomyopathy and myocardial resilience.** *Nature genetics*
Jurgens, S. J., Rämö, J. T., Kramarenko, D. R., Wijdeveld, L. F., Haas, J., Chaffin, M. D., Garnier, S., Gaziano, L., Weng, L. C., Lipov, A., Zheng, S. L., Henry, A., Huffman, et al
2024
- **Genome-wide association analyses identify distinct genetic architectures for age-related macular degeneration across ancestries.** *Nature genetics*
Gorman, B. R., Voloudakis, G., Igo, R. P., Kinzy, T., Halladay, C. W., Bigdeli, T. B., Zeng, B., Venkatesh, S., Cooke Bailey, J. N., Crawford, D. C., Markianos, K., Dong, F., Schreiner, et al
2024
- **Genome-wide association study reveals mechanisms underlying dilated cardiomyopathy and myocardial resilience.** *Nature genetics*
Jurgens, S. J., Rämö, J. T., Kramarenko, D. R., Wijdeveld, L. F., Haas, J., Chaffin, M. D., Garnier, S., Gaziano, L., Weng, L. C., Lipov, A., Zheng, S. L., Henry, A., Huffman, et al
2024
- **Polygenic Risk Scores and Risk of Drug-induced Liver Injury following initiation of antiretroviral therapy in people living with HIV**
Cindi, Z., Cardone, K. M., Bradford, Y., Kim, H., Daar, E. S., Gulick, R., Riddler, S. A., Vujkovic, M., Chang, K., Tsao, P. S., Maartens, G., Sinxadi, P., Haas, et al
WILEY.2024: 351
- **NOVEL GENETIC ARCHITECTURE AND METABOLIC INTERACTIONS FOR ALL-CAUSE CIRRHOSIS AND HEPATOCELLULAR CARCINOMA DEFINED IN MULTI-ANCESTRY META-ANALYSES**
Vujkovic, M., Kaplan, D., Ghouse, J., Levin, M., Brancale, J., Lewis, A., Veatch, O., Schneider, C., Chang, B., Verma, A., Wangensteen, K., Gill, D., Konkwo, et al
LIPPINCOTT WILLIAMS & WILKINS.2024: S1992-S1993
- **Plasma proteomics reveals the potential causal impact of extracellular matrix proteins on abdominal aortic aneurysm.** *medRxiv : the preprint server for health sciences*
Khodursky, S., Yuan, S., Spin, J. M., Tsao, P. S., Levin, M. G., Damrauer, S. M.

2024

- **Exome wide association study for blood lipids in 1,158,017 individuals from diverse populations.** *medRxiv : the preprint server for health sciences*
Koyama, S., Yu, Z., Choi, S. H., Jurgens, S. J., Selvaraj, M. S., Klarin, D., Huffman, J. E., Clarke, S. L., Trinh, M. N., Ravi, A., Dron, J. S., Spinks, C., Surakka, et al
2024
- **A plasma proteomic signature for atherosclerotic cardiovascular disease risk prediction in the UK Biobank cohort.** *medRxiv : the preprint server for health sciences*
Gupte, T. P., Azizi, Z., Kho, P. F., Zhou, J., Chen, M., Panyard, D. J., Guarischi-Sousa, R., Hilliard, A. T., Sharma, D., Watson, K., Abbasi, F., Tsao, P. S., Clarke, et al
2024
- **Plasma proteomic signatures for type 2 diabetes mellitus and related traits in the UK Biobank cohort.** *medRxiv : the preprint server for health sciences*
Gupte, T. P., Azizi, Z., Kho, P. F., Zhou, J., Nzenkue, K., Chen, M., Panyard, D. J., Guarischi-Sousa, R., Hilliard, A. T., Sharma, D., Watson, K., Abbasi, F., Tsao, et al
2024
- **Genetically predicted lipoprotein(a) associates with coronary artery plaque severity independent of low-density lipoprotein cholesterol.** *European journal of preventive cardiology*
Clarke, S. L., Huang, R. D., Hilliard, A. T., Levin, M. G., Sharma, D., Thomson, B., Lynch, J., Tsao, P. S., Gaziano, J. M., Assimes, T. L.
2024
- **Association of Cardiac MRI-derived Aortic Stiffness with Early Stages and Progression of Heart Failure with Preserved Ejection Fraction.** *Radiology. Cardiothoracic imaging*
Schulz, A., Schellinger, I. N., Backhaus, S. J., Adler, A. S., Lange, T., Evertz, R., Kowallick, J. T., Hoffmann, A., Matek, C., Tsao, P. S., Hasenfuß, G., Raaz, U., Schuster, et al
2024; 6 (4): e230344
- **Mitochondrial Transplantation Promotes Protective Effector and Memory CD4+ T Cell Response During Mycobacterium Tuberculosis Infection and Diminishes Exhaustion and Senescence in Elderly CD4+ T cells.** *Advanced science (Weinheim, Baden-Wuerttemberg, Germany)*
Headley, C. A., Gautam, S., Olmo-Fontanez, A., Garcia-Vilanova, A., Dwivedi, V., Schami, A., Weintraub, S., Tsao, P. S., Torrelles, J. B., Turner, J.
2024: e2401077
- **Diversity and scale: Genetic architecture of 2068 traits in the VA Million Veteran Program.** *Science (New York, N.Y.)*
Verma, A., Huffman, J. E., Rodriguez, A., Conery, M., Liu, M., Ho, Y. L., Kim, Y., Heise, D. A., Guare, L., Panickan, V. A., Garcon, H., Linares, F., Costa, et al
2024; 385 (6706): eadj1182
- **Abdominal aortic aneurysm and cardiometabolic traits share strong genetic susceptibility to lipid metabolism and inflammation.** *Nature communications*
Zheng, S., Tsao, P. S., Pan, C.
2024; 15 (1): 5652
- **Clinical and genetic risk factors for progressive fibrosis in metabolic dysfunction-associated steatotic liver disease.** *Hepatology communications*
Kaplan, D. E., Teerlink, C. C., Schwantes-An, T. H., Norden-Krichmar, T. M., DuVall, S. L., Morgan, T. R., Tsao, P. S., Voight, B. F., Lynch, J. A., Vujković, M., Chang, K. M.
2024; 8 (7)
- **Gasdermin D Inhibitor Necrosulfonamide Alleviates Angiotensin II-Induced Abdominal Aortic Aneurysms in Apolipoprotein E-Deficient Mice.** *Biomolecules*
Guo, J., Zhang, Q., Li, Z., Qin, M., Shi, J., Wang, Y., Ai, W., Ju, J., Samura, M., Tsao, P. S., Xu, B.
2024; 14 (6)
- **Postdoctoral T32 training is correlated with obtaining an academic primarily research faculty position.** *PloS one*
Mueller, A. L., Schnirel, A., Kleppner, S., Tsao, P., Leeper, N. J.
2024; 19 (6): e0303792

- **Deconvolution of polygenic risk score in single cells unravels cellular and molecular heterogeneity of complex human diseases.** *bioRxiv : the preprint server for biology*
Zhang, S., Shu, H., Zhou, J., Rubin-Sigler, J., Yang, X., Liu, Y., Cooper-Knock, J., Monte, E., Zhu, C., Tu, S., Li, H., Tong, M., Ecker, et al
2024
- **Integrative common and rare variant analyses provide insights into the genetic architecture of liver cirrhosis.** *Nature genetics*
Ghouse, J., Sveinbjörnsson, G., Vujkovic, M., Seidelin, A. S., Gellert-Kristensen, H., Ahlberg, G., Tragante, V., Rand, S. A., Brancale, J., Vilarinho, S., Lundegaard, P. R., Sørensen, E., Erikstrup, et al
2024
- **Type 1 Diabetes Genetic Risk in 109,954 Veterans With Adult-Onset Diabetes: The Million Veteran Program (MVP).** *Diabetes care*
Yang, P. K., Jackson, S. L., Charest, B. R., Cheng, Y. J., Sun, Y. V., Raghavan, S., Litkowski, E. M., Legvold, B. T., Rhee, M. K., Oram, R. A., Kuklina, E. V., Vujkovic, M., Reaven, et al
2024
- **A multi-ancestry GWAS of Fuchs corneal dystrophy highlights the contributions of laminins, collagen, and endothelial cell regulation** *COMMUNICATIONS BIOLOGY*
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