Pharmacogenetics of Asthma Treatment (PHAT)

Scott T. Weiss, MD, Brigham and Women's Hospital

Augusto A. Litonjua, MD, Brigham and Women's Hospital Kelan G. Tantitsira, MD, Brigham and Women's Hospital Elliot Israel, MD, Brigham and Women's Hospital Ross Lazarus, MPH, Brigham and Women's Hospital Thomas J. Mariani, PhD, Brigham and Women's Hospital Steve D. Shapiro, MD, Brigham and Women's Hospital Michael E. Wechsler, MD, Brigham and Women's Hospital Jin Song Xu, PhD, Brigham and Women's Hospital Eugene R. Bleecker, MD, Wake Forest University Gregory Hawkins, PhD, Wake Forest University Deborah A. Meyers, PhD, Wake Forest University Nan M. Laird, PhD, Harvard School of Public Health Christoph Lange, PhD, Harvard School of Public Health Dennis W. McGraw, MD, University of Cincinnati Kersten M. Small, PhD, University of Cincinnati Marco F. Ramoni, PhD, Children's Hospital Stephen B. Liggett, MD, University of Maryland

Using genetics to predict responders to asthma therapy, based upon understanding of asthma pathways and clinically significant variation

Funded by NHLBI

Pharmacogenomics and Risk of Cardiovascular Disease (PARC)

Ronald M. Krauss, MD, Children's Hospital Oakland Research Institute

Deborah A. Nickerson, PhD, University of Washington Jerome I. Rotter, MD, Cedars-Sinai Medical Center Xiuqing Guo, PhD, Cedars-Sinai Medical Center Dai Vang, PhD, Cedars-Sinai Medical Center Paul T. Williams, PhD, Cedars-Sinai Medical Center Daniel I. Chasman, MD, Brigham and Women's Hospital Paul T. Williams, PhD, Brigham and Women's Hospital Nader R. Gherbranious, PhD, Marshfield Clinic Catherine A. McCarty, PhD, Marshfield Clinic Russell A. Wilkie, MD, PhD, Marshfield Clinic Patricia J. Blanche, BS, Children's Hosp. Oakland Res. Inst. Lara Mangravite, PhD, Children's Hosp. Oakland Res. Inst. Mark J. Rieder, PhD, University of Washington Joshua D. Smith, BS, University of Washington

Defining genetic contributions to differences among individuals in their responses to statin drugs and cardiovascular disease risk Funded by NHLBI

Amish Pharmacogenomics of Antiplatelet Intervention Study (PAPI)

Alan R. Shuldiner, MD, University of Maryland

Stuart T. Haines, PharmD, University of Maryland Braxton D. Mitchell, Jr., PhD, University of Maryland Jeffrey R. O'Connell, PhD, University of Maryland Toni I. Pollin, PhD, University of Maryland Nauder Faraday, MD, Johns Hopkins University William R. Herzog, MD, Johns Hopkins University Wendy S. Post, MD, Johns Hopkins University

Understanding genetic determinants of responses that vary among people taking anti-platelet agents used to treat and prevent cardiovascular disease

Funded by NIGMS

PharmGKB: Catalyzing Research in Pharmacogenetics

Russ B. Altman, MD, PhD, Stanford University

Teri E. Klein, PhD, Stanford University Tina M. Boussard, PhD, Stanford University Michelle W. Carrillo, PhD, Stanford University Winston Gor, MS, Stanford University Wark Kiuchi, MBA, Stanford University Feng Liu, MS, Stanford University Andrew MacBride, BA, Stanford University Caroline F. Thorn, PhD, Stanford University TC Truong, BS, Stanford University Mark Woon, BSE, Stanford University Tina Zhou, MSE, MSP, Stanford University

Designing a knowledge base to link phenotypes to genotypes in pharmacogenetics and pharmacogenomics, to be used as a tool to enable future research efforts

Funded by NIGMS, NHLBI, NHGRI, and NLM

A goal of the PGRN and PharmGKB is to aid all researchers in understanding how genes vary among individuals, and how that affects drug safety and efficacy. The network hopes to set standards for future research studies, and make scientific recommendations that will ultimately impact the clinical use of drugs.

Pharmacogenetics Research Network

National Institutes of Health U.S. Department of Health & Human Services

The NIH Pharmacogenetics Research Network (PGRN) is

a nationwide collaboration of researchers studying the contribution of genetics to predicting responses to a wide variety of medicines. The overall aims of the groups are to discover important drug pathways, to identify sequence variants in relevant genes, and to establish biological relationships to clinical drug responses. Since its inception in 2000, scientists have studied genes and medications given for a range of diseases, including asthma, cancer, heart disease, and depression.

www.nigms.nih.gov/pharmacogenetics

PharmG_{кв}

The Pharmacogenetics and Pharmacogenomics Knowledge Base (PharmGKB) is an integrated knowledge base for pharmacogenetics linking phenotypes and genotypes. It is available for the entire scientific community to make deposits and to use the data.

- A web-based format for pharmacogenetics knowledge
- Curated, linked genotypes and phenotypes
- Genomic, molecular and cellular, and clinical datasets
- Annotated, interactive, consensus drug pathways
- Automated methods for identifying relationships
- Community-based literature submissions
- Access to the entire research community

www.pharmgkb.org

Research Groups

Pharmacogenetics of Membrane Transporters (PMT)

Kathleen M. Giacomini, PhD, University of California, San Francisco

Deanna L. Kroetz, PhD, University of California, San Francisco Leslie Z. Benet, PhD, University of California, San Francisco Claire M. Brett, MD. University of California, San Francisco Esteban G. Burchard, MD, University of California, San Francisco Richard M. Castro, MD, University of California, San Francisco Robert H. Edwards, MD, University of California, San Francisco Thomas E. Ferrin, PhD, University of California, San Francisco Charles E. Glatt, MD, PhD, University of California, Los Angeles Pui-Yan Kwok, MD, PhD, University of California, San Francisco Emil T. Lin, PhD, University of California, San Francisco Victor I. Reus, MD, University of California, San Francisco Neil J. Risch, PhD, University of California, San Francisco Andrei Sali, PhD, University of California, San Francisco Mark R. Segal, PhD, University of California, San Francisco John S. Witte, PhD, University of California, San Francisco Charles P. Quesenberry, Jr., PhD, University of California, Berkeley Jasper J. Rine, PhD, University of California, Berkeley Catherine A. Schaefer, PhD, Kaiser Permanente Richard M. Myers, PhD, Stanford University Wolfgang Sadee, PhD, Ohio State University

Studying membrane transporter genes in ethnically diverse samples, determining cellular phenotypes, and correlating variants with the clinical response to antidepressants Funded by NIGMS, NIMH, and NCI

Comprehensive Research on Expressed Alleles in Therapeutic Intervention (CREATE)

Howard L. McLeod, PharmD, Washington University

Michael A. Province, PhD, Washington University Mark Watson, MD, PhD, Washington University Sharon Marsh, PhD, Washington University Rakesh Nagarajan, PhD, Washington University Gary Stormo, PhD, Washington University Alan Templeton, PhD, Washington University Barbara Zehnbauer, PhD, Washington University Pui-Yan Kwok, MD, PhD, University of California, San Francisco

Identifying pathways of anti-cancer drugs, taking novel approaches to pathway dissection, and functionally assessing variants

Funded by NIGMS and NCI

Pharmacogenomics of Arrhythmia Therapy (PAT)

Dan M. Roden, MD, Vanderbilt University

Nancy J. Brown, MD. Vanderbilt University Dawood Darber, MD, PhD, Vanderbilt University Brian S. Donahue, MD, PhD, Vanderbilt University Alfred L. George, Jr., MD, Vanderbilt University Dario A. Giuse, PhD, Vanderbilt University Jonathan L. Haines, PhD, Vanderbilt University Prince J. Kannankeril, MD, Vanderbilt University Richard B. Kim, MD, PhD, Vanderbilt University Peter J. Mohler, PhD, Vanderbilt University Marylyn D. Ritchie, PhD, Vanderbilt University C. Michael Stein, MD, Vanderbilt University Janey Wang, MS, Vanderbilt University Ping Yang, PhD, Vanderbilt University Tao Yang, PhD, Vanderbilt University Patrick T. Ellinor, MD, PhD, Massachusetts General Hospital Calum A. MacRae, MD, PhD, Massachusetts General Hospital Theodore R. Addai, MD, Meharry Medical College Michael D. Floyd, MD, Meharry Medical College David J. Milan, MD, Brigham and Women's Hospital

Examining genes that modulate cardiac rhythm, accruing patients with QT responses to therapy, and evaluating drug responses in atrial fibrillation

Funded by NHLBI

Consortium on Breast Cancer Pharmacogenomics (COBRA)

David A. Flockhart, MD, PhD, Indiana University

Daniel F. Hayes, MD, University of Michigan Vered Stearns, MD, Johns Hopkins University Janet S. Carpenter, RN, PhD, Indiana University Zeruesenay Desta, PhD, Indiana University Stephen D. Hall, PhD, Indiana University Lang Li, PhD, Indiana University Sean D. Mooney, PhD, Indiana University Anne Nguyen, BS, Indiana University Todd C. Skaar, PhD, Indiana University Anna Maria Storniolo, MD, Indiana University James M. Rae, PhD, University of Michigan Nancy E. Davidson, MD, Johns Hopkins University Steffi Oestereich, PhD, Baylor University

Describing normal functions of estrogen, and how genetic variation contributes to efficacy and toxicity of endocrine treatments for breast cancer

Funded by NIGMS, NIEHS, and ORWH

Pharmacogenomic Evaluation of the Antihypertensive Response (PEAR)

Julie A. Johnson, PharmD, University of Florida

Eric Boerwinkle, PhD, University of Texas, Houston Stephen T. Turner, MD, Mayo Clinic

Rhonda Cooper-DeHoff, PharmD, University of Florida R. Whit Curry, MD, University of Florida Yan Gong, PhD, University of Florida John G. Gums, PharmD, University of Florida Laurence Kennedy, MD, University of Florida Taimour Langee, PhD, University of Florida Issam Zineh, PharmD, University of Florida Kent R. Bailey, PhD, Mayo Clinic Gary L. Schwartz, MD, Mayo Clinic Evelyn Baranco, MD, Emory University Arlene B, Chapman, MD, Emory University Andrei Rodin, PhD, University of Texas, Houston

Identifying gene variants involved in responses to drugs commonly used for hypertension, a beta-blocker and a thiazide diuretic, to predict therapeutic and adverse responses

Funded by NIGMS

Pharmacogenetics of Nicotine Addiction and Treatment (PNAT)

Neal L. Benowitz, MD, University of California, San Francisco and Huijun Z. Ring, PhD, SRI International

Stanley P. Azen, PhD, University of Southern California David V. Conti, PhD, University of Southern California Daniel O. Stram, PhD, University of Southern California Puter D. Karp, PhD, SRI International Christina N. Lessov, PhD, SRI International Yannick Pouliot, PhD, SRI International Gary E. Swan, PhD, SRI International Delia A. Dempsey, MD, University of California, San Francisco Peyton Jacob III, PhD, University of California, San Francisco Jennifer B. McClure, PhD, Group Health Cooperative Caryn E. Lerman, PhD, University of Pennsylvania Stephen I. Rennard, MD, University of Nebraska Rachel T. Tyndale, PhD, University of Toronto

Investigating the genetic basis of nicotine addiction and influences on responses to drug therapies used for smoking cessation

Funded by NIDA, NIGMS, and NCI

Pharmacogenetics of Anticancer Agents Research Group (PAAR) Mark J. Ratain, MD, University of Chicago and Mary V. Relling, PharmD, St. Jude

Children's Research Hospital

Judith A. Badner, MD, University of Chicago Nancy J. Cox, PhD, University of Chicago Soma Das, PhD, University of Chicago Christopher K. Daughtery, MD, University of Chicago Apurva A. Desai, MD, University of Chicago Anna DiRienzo, PhD, University of Chicago M. Eileen Dolan, PhD, University of Chicago Federico Innocenti, MD, PhD, University of Chicago Wanging Liu, PhD, University of Chicago Amittha Wickrema, PhD, University of Chicago Xiaolin Wu, PhD, University of Chicago William E. Evans, PharmD., St. Jude Children's Research Hospital Cheng Cheng, PhD, St. Jude Children's Research Hospital Wei Liu, PhD, St. Jude Children's Research Hospital Clavton W. Naeve, PhD, St. Jude Children's Research Hospital Geoffrey A. M. Neale, PhD, St. Jude Children's Research Hospital Deging Pei, MS, St. Jude Children's Research Hospital Ching-Hon Pui, MD, St. Jude Children's Research Hospital Erin G. Schuetz, PhD, St. Jude Children's Research Hospital John D. Schuetz, PhD, St. Jude Children's Research Hospital Wenjian Yang, PhD, St. Jude Children's Research Hospital Stephen C. Strom, MD, University of Pittsburgh Gary L. Rosner, ScD, MD Anderson Cancer Center Christopher I. Amos, PhD, MD Anderson Cancer Center Edwin H. Cook, PhD, University of Illinois at Chicago

Elucidating the impact of germ-line variants on the efficacy and adverse effects of anticancer drugs used to treat colorectal cancer and childhood leukemias

Funded by NIGMS and NCI

Pharmacogenetics of Phase II Drug Metabolizing Enzymes (PPII)

Richard M. Weinshilboum, MD, Mayo Clinic

Eric D. Wieben, PhD, Mayo Clinic

Alex A. Adjei, MD, Mayo Clinic John L. Black, III, MD, Mayo Clinic Christopher G. Chute, MD, DrPH, Mayo Clinic Jalie M. Cunningham, PhD, Mayo Clinic James N. Ingle, MD, Mayo Clinic David A. Mrazek, MD, FRC, Mayo Clinic Janet E. Olson, PhD, Mayo Clinic Daniel J. Schaid, PhD, Mayo Clinic Celine M. Vachon, PhD, Mayo Clinic Liewei Wang, MD, PhD, Mayo Clinic Aman Buzdar, MD, MD Anderson Cancer Center Vivien C. Yee, PhD, Case Western Reserve

Discovering variants and identifying mechanisms involved in phase II conjugating enzymes controlling biotransformation of drugs, hormones, and neurotransmitters

Funded by NIGMS and NCI