

Date Prepared: August 26, 2014

Name: Christopher Holmes Newton-Cheh, MD, MPH, FAHA

Office Address: [REDACTED]

Home Address: [REDACTED]

Work Phone: [REDACTED]

Work Email: [REDACTED]

Work FAX: [REDACTED]

Place of Birth: [REDACTED]

Education

1986-1991	B.A.	English	Dartmouth College
1992-1996	M.D.		Columbia College of Physicians & Surgeons
2002-2004	M.P.H	Clinical Effectiveness	Harvard School of Public Health

Postdoctoral Training

07/96-06/99	Resident	Medicine	Massachusetts General Hospital
07/99-06/01,	Clinical Fellow	Cardiology	Massachusetts General Hospital
07/02-06/03			
07/01-06/02	Chief Resident	Medicine	Massachusetts General Hospital
07/02-06/04	Postdoctoral Fellow	Epidemiology Christopher O'Donnell, mentor	Framingham Heart Study
07/02-06/07	Postdoctoral Fellow	Genomics Joel Hirschhorn, mentor	Broad Institute of Harvard and MIT

Faculty Academic Appointments

07/01-07/08	Instructor	Medicine	Harvard Medical School
08/08-	Assistant Professor	Medicine	Harvard Medical School

Appointments at Hospitals/Affiliated Institutions

11/98-09/03	Hospitalist	Medicine	Emerson Hospital
10/02-11/04	Intensivist	Medicine	Newton-Wellesley Hospital
07/03-06/04	Graduate Assistant	Cardiology	Massachusetts General Hospital
07/04-06/09	Assistant in Medicine	Cardiology	Massachusetts General Hospital
07/09-	Assistant Physician	Cardiology	Massachusetts General Hospital

Other Professional Positions

2004-	Affiliate Scientist	Framingham Heart Study
2007-	Faculty	Center for Human Genetic Research, MGH
2007-	Faculty	Cardiovascular Research Center, MGH
2007-2013	Affiliate Scientist	Broad Institute of Harvard and MIT
2009-	Faculty	Biological and Biomedical Sciences, Harvard University
2009-	Scientific Advisory Board	FDA Endocrinologic and Metabolic Drugs Committee
2012-	Scientific Advisory Board	Arrhythmogenic Potential of Drugs (ARITMO, Erasmus Medical Center, Rotterdam, Netherlands)
2013-	Associate Member	Broad Institute of Harvard and MIT

Major Administrative Leadership Positions

Local

2000-2001	Administrative Fellow	Cardiology, Massachusetts General Hospital
2001-2002	Chief Resident	Medicine, Massachusetts General Hospital
2005-	Course Director, Primer in Medical and Population Genetics	Broad Institute of Harvard and MIT
2007-	Medical Director, Electrocardiography Laboratory	Massachusetts General Hospital
2007-	Conference organizer, A Primer on Complex Trait Genetics: Basic Principles for the Beginning Investigator	Massachusetts General Hospital
2007-	Co-director, Human Cardiovascular Genetics Program	Cardiovascular Research Center, Massachusetts General Hospital
2008-	Co-course director, Genetics in Medicine: Bench to Bedside	Harvard Medical School

Committee Service

Local

2005-2006	Medical and Population Genetics Planning Committee	Broad Institute of Harvard and MIT Member
-----------	--	---

National

2009-	Cardiac Safety Research Consortium	FDA/Pharma/Academic roundtable Member
2013-	QT Drug Lists Advisory Board	Academic review of literature to guide clinicians (qtdrugs.org)

Professional Societies

1996-	American Medical Association	
1996-	Massachusetts Medical Society	
1999-	American College of Cardiology	
2006-		Fellow
2003-	American Heart Association	
2007-		Fellow, AHA Council on Epidemiology & Prevention; Council on Basic Cardiovascular Sciences; Council on Functional Genomics and Translational Biology

2006-	AHA Functional Genomics and Translational Biology Council	
2006-		Leadership Committee Member
2009-2011		Vice Chair, Program Committee
2011-		Chair, Program Committee
2006-	Heart Rhythm Society	
2010-		Member

Grant Review Activities

2005-	Subcommittee on Review of Research Proposals, Executive Committee on Research	Massachusetts General Hospital Reviewer
2009-	Endocrinologic & Metabolic Drugs Advisory Committee	Food and Drug Administration Ad Hoc Reviewer
2010-	Genomics & Translational Biology Epidemiology and Observational Epidemiology study section	American Heart Association Reviewer
2010-	Clinical and Integrative Cardiovascular Science study section	National Institutes of Health Ad Hoc Reviewer
2012-	Genetics of Health and Disease	National Institutes of Health Ad Hoc Reviewer

Editorial Activities

Ad hoc Reviewer

New England Journal of Medicine
 Journal of the American Medical Association
 Nature Genetics
 Circulation
 Circulation Research
 Science
 American Journal of Human Genetics
 PLoS Genetics
 Human Molecular Genetics
 European Journal of Human Genetics
 Journal of the American College of Cardiology
 Nature Medicine

Other Editorial Roles

2007-2012	Section Editor	Genetics Section, Current Cardiovascular Risk Reports
2008-	Member, Editorial Board	Circulation: Cardiovascular Genetics

Honors and Prizes

1995	Alpha Omega Alpha	Columbia University College of Physicians & Surgeons	Medical Honor Society
2002, 2009	Partners in Excellence Award	Massachusetts General Hospital	
2008, 2011	Excellence in Tutoring Award	Harvard Medical School	Human Genetics, First Year Medical Students
2009	Best Translational	Massachusetts General	Selected from 256 abstracts

	Research Abstract, MGH Clinical Research Day Top Ten Cardiovascular Discoveries of 2009	Hospital American Heart Association	
2009			“GWAS identifies eight loci associated with blood pressure” (Newton-Cheh C et al <i>Nature Genetics</i> 2009)
2014	Top Original Paper of 2013 Award, Popular Science Category	Hypertension	“A Blood Pressure Genetic Risk Score Is a Significant Predictor of Incident Cardiovascular Events in 32 669 Individuals”

Funding Information

Past

2002-2005	Candidate gene association with QT interval (Young Investigator Award) GlaxoSmithKline Young Investigator Award PI (\$150,000)		
2002-2005	Study focused on candidate genes involved in Long QT Syndrome Training Program in the Epidemiology of Cardiovascular Disease NIH/T32 (HL07575) Trainee (PI: P. Ridker) Support for training in epidemiology and statistics		
2005 (returned)	Candidate gene and genome-wide association of QT interval American Heart Association Scientist Development Grant PI (\$236,364) Returned for overlap when K23 funded		
2005-2010	Genetic determinants of cardiac repolarization NIH/K23 (HL080025) PI (\$149,500/year) Candidate gene, genome-wide association and large-scale resequencing study of QT interval		
2006-2010	Genetic determinants of QT interval, sudden death Doris Duke Clinical Scientist Development Award PI (NCE) Career development award supporting large-scale replication of QT interval genome-wide association studies		
2009-2010	Effect of Moxifloxacin on the QT interval: Pilot NIH/CTSC PI (\$50,000/year) Principal investigator on project examining the QT interval response to oral moxifloxacin as pilot for large-scale pharmacogenetic study.		
2009-2012	Cardiology and Metabolic Patient Cohort Pfizer Co-I (\$9,985/year) The goal of this project is to recruit 3000 patients from the MGH Heart Center, to store serum and plasma for future studies, and to genotype DNA for genes that confer susceptibility to T2D and cardiovascular disease.		
2007-2012	Genome-wide association study of sudden cardiac arrest NIH/R01 (HL088456)		

- PI- subcontract (PI: N. Sotoodehnia)
 PI of subcontract providing study design and analysis methods for study of genetics of sudden cardiac death
- 2007-2013 Genomic dissection of QT interval duration and sudden death
 Burroughs Wellcome Career Award for Medical Scientists
 PI (\$151,000/year TDC)
 Career development award supporting QT interval genome-wide association and extension to sudden cardiac death cohorts.
- 2010-2013 Physiologic effects of natriuretic peptide genetic variation
 NIH/R01 (HL098283)
 PI (PIs: Newton-Cheh, Wang) (\$178,795/year TDC)
 Study to fine map and identify novel genetic factors influencing plasma natriuretic peptide concentration through resequencing and to characterize the physiologic effects of genetic variants through chronic and acute sodium challenge.

Current

- 2013-2017 Physiologic profiling of sGC genetic variants
 NIH/R01 (HL113933)
 PI (\$333,000/year TDC)
 Study to characterize the role of soluble guanylate cyclase as the mediator of a blood pressure association at the locus using genotype-directed translational studies in humans and to elucidate the causal variant and its mechanism of action using cellular models of human genetic variants.

Teaching of Students in Courses

2001-2002	Core II, medical sub-internship HMS Fourth year students	Massachusetts General Hospital 2 hours per week (4 months)
2004-2006	Program in Clinical Effectiveness: Critique of the Literature seminar MPH Students	Harvard School of Public Health 2 hours
2004-2006	Core I Electrocardiography Series, Massachusetts General Hospital: Introduction to electrocardiography including leads, rates, intervals, determination of axis. Medical Students	Harvard Medical School 1 hour
2006-2010 & 2012	First Year Human Genetics Small group session of medical students	Harvard Medical School 6 hours per week (3 weeks)
2007-	Genetics in Medicine: Bench to Bedside PhD Students	Harvard Medical School 3 hours
2008-	Program in Clinical Effectiveness: Genetic epidemiology MPH Students	Harvard School of Public Health 2 hours

Clinical Supervisory and Training Responsibilities

2001-2002	Chief Medical Resident Massachusetts General Hospital	Full-time, one year
2004-2006	Attending, Ellison general medicine service Massachusetts General Hospital	2-3 hours, twice weekly, one month
2004-2005,	Attending, Cardiology Consultation service	4 hours, 7 days per week, two weeks

2007	Massachusetts General Hospital	
2004-	Attend in outpatient fellow's clinic	4 times per year
	Massachusetts General Hospital	
2005-2013	Attend on inpatient Heart Failure and Transplant service	4 hours, 7 days per week, one half month; 3-4 times per year
	Massachusetts General Hospital	
2005	Attend on inpatient Cardiac Arrhythmia service	4 hours, 7 days per week, one half month
	Massachusetts General Hospital	
2006-2013	Attend on inpatient Coronary Care Unit	4 hours, 7 days per week, one half month
	Massachusetts General Hospital	
2008-2013	Attend on inpatient Heart Failure and Transplant service	4 hours, 7 days per week, one half month; 2 times per year

Laboratory and Other Research Supervisory and Training Responsibilities

2006	Supervision of technician (Ripal Shah) who subsequently went into industry as a consultant	Weekly mentorship for 12 months
2006	Supervision of technician (Alim Hirji) as a medical student who subsequently completed pediatric residency at the Hospital for Sick Kids of Toronto	Weekly mentorship for 12 months
2006-2007	Supervision of technician (Sirisha Kovvali) who subsequently matriculated to the Oregon Health & Science University, MPH	Weekly mentorship for 24 months
2006-2007	Supervision of technician (Aarti Surti) who subsequently matriculated to the University of Michigan, MD	Weekly mentorship for 24 months
2007	Supervision of technician (Marie-Louise Meng) who subsequently matriculated to Harvard Medical School, MD	Weekly mentorship for 12 months
2007, 2008	Supervision of technician (Sara Tribune) who subsequently matriculated to Brown University, MD	Summer mentorship for 3 months each summer
2009-2012	Supervision of clinical research coordinator (Allicia Ryan) who subsequently matriculated to a school of osteopathy	Weekly mentorship for 36 months
2010	Supervision of technician (Kimberly Kummer), now a college student at Dartmouth	Summer mentorship for 3 months
2010-2012	Supervision of clinical research coordinator (Andrew Martinez)	Weekly mentorship for 24 months
2011, 2012	Supervision of technician (Samuel Kim) who subsequently matriculated to Harvard Medical School, MD	Summer mentorship for 3 months each summer
2011-2012	Supervision of research analyst (Sara Pulit) who subsequently matriculated to University of Utrecht, Netherlands, MS	Weekly mentorship for 12 months

2013	Supervision of clinical research coordinator (Jillian Pickett)	Summer mentorship for 3 months
2013-	Supervision of clinical research coordinator (Courtney Karol)	Weekly mentorship

Formally Supervised Trainees

2006-	Peter Noseworthy, MD/ Electrophysiologist/Physician-scientist, Mayo Clinic Supervision of postdoctoral fellow in studies of the genetic basis of cardiac arrhythmias and cardiotoxic drug responses. Peter authored a review of the genetics of sudden cardiac death published in <i>Circulation</i> , has participated in a genome-wide association study of QT interval published in <i>Nature Genetics</i> is conducting large-scale studies of association of genetic variants altering QT interval in the Finrisk study, and a physiologic study of QT interval response to exposure to moxifloxacin. Peter was awarded two post-doctoral fellowship grants on which I am the primary sponsor for this work from the American Heart Association (returned) and the Heart Rhythm Society (current).	
2007-2012	J. Gustav Smith, Faculty, Lund University, Sweden	
2014-	Supervision of medical student (and subsequently PhD student) in genome-wide association studies of electrocardiographic measures and biomarkers of cardiovascular disease Gustav published 14 articles on which we were both co-authors, on 9 of which he was the first author. I was one of three thesis mentors on his PhD at Lund University, Sweden. He is currently completing a second post-doc in my laboratory for 2 years.	
2008-2013	Pankaj Arora, MD/ Cardiology Fellow, University of Alabama Supervision of postdoctoral fellow in resequencing and association studies of blood pressure genes in the Framingham Heart Study and in the Malmö Diet and Cancer Study. Pankaj worked on several studies of the genetic basis of blood pressure regulation. Pankaj authored a review of the genetic basis of hypertension at <i>Current Opinion in Cardiology</i> and an original research article in the <i>Journal of Clinical Investigation</i> .	
2009-2010	Abigail Khan, MD/ Cardiology Fellow, University of Pennsylvania Supervision of postdoc in characterizing the physiologic response in humans of genetic variation at the natriuretic peptide locus. Abigail was a postdoctoral fellow in my lab and worked on several studies of the genetic basis of blood pressure regulation.	
2010-2011	Sandosh Padmanabhan, MD/ Faculty, University of Glasgow Supervision of postdoctoral fellow, on one-year sabbatical from University of Glasgow where he is on the Genetics Faculty, using targeted exonic resequencing to characterize novel blood pressure loci. Sandosh is supported by an early faculty award from the British Heart Foundation on which I am the primary sponsor.	
2010-2012	Erin Coglianese, MD/ Staff Cardiologist, Loyola University Medical Center Supervision of postdoctoral research fellow initiating and completing a pilot study of the role of common genetic variation in the response to inhaled nitric oxide to examine the physiologic basis of an association with blood pressure.	
2013-	Gulum Kosova, PhD/ Postdoctoral fellow, Massachusetts General Hospital Supervision of post-doctoral research fellow, focused on blood pressure genetics studies and computational biology. She is using next generation sequence data to identify novel genetic variation that influences blood pressure regulation.	
2013-	Michael Rosenberg, MD/ Postdoctoral fellow, Massachusetts General Hospital Supervision of post-doctoral fellow (trained as an electrophysiologist) focused on studies of sudden cardiac death and arrhythmias in response to drug exposure. Mike is leading several	

translational studies and runs a multi-center consortium of torsade and drug response to anti-arrhythmics.

Formal Teaching of Peers (e.g., CME and other continuing education courses)

2005	Cardiac Case Review Harvard Review of Internal Medicine, Massachusetts General Hospital	One Talk Cambridge, Massachusetts
2006, 2007, 2008, 2009, 2010	Heart Failure Management Harvard Primary Care Internal Medicine, Massachusetts General Hospital	One Talk Cambridge, Massachusetts
2011	EPI Lecture Harvard School of Public Health	One Talk Boston, Massachusetts
2011	Heart Failure/Transplant Services Lecture Massachusetts General Hospital	One Talk Boston, Massachusetts

Invited Presentations and Courses

Local

2002	The Long QT Syndrome: Implications for the General Population / Grand Rounds Department of Medicine, Massachusetts General Hospital, Boston, MA
2003	Genetics of QT Interval Variation in the General Population / Research Conference Division of Preventative Medicine, Brigham and Women’s Hospital, Boston, MA
2004	Dissecting the Genetic Determinants of Variation in Cardiac Repolarization / Research Conference Division of Preventive Medicine, Brigham and Women’s Hospital, Boston, MA
2004	Genetic Determinants of Variation in Electrocardiographic Measures / Research Conference Endocrine Program, Broad Institute of Harvard and MIT, Cambridge, MA
2004	Genetic Dissection of QT Interval Variation in the General Population / Research Conference Program in Medical and Population Genetics, Broad Institute of Harvard and MIT, Cambridge, MA
2005	Approaches to the Genetic Dissection of Complex Traits / Epidemiology Core Curriculum Cardiology Fellowship, Massachusetts General Hospital, Boston, MA
2005	Clinical and Genetic Correlates of the Aldosterone to Renin Ratio in the Framingham Heart Study / Research Conference Altshuler Laboratory, Department of Molecular Biology, Massachusetts General Hospital, Boston, MA
2005	Common variation in CAPON is reproducibly associated with QT interval variation in 3 cohorts / Research Conference Altshuler Laboratory, Department of Molecular Biology, Massachusetts General Hospital, Boston, MA
2006	Genetic Determinants of the Electrocardiographic QT Interval: Insights from Population Studies / Medicine Grand Rounds Department of Medicine, Massachusetts General Hospital, Boston, MA
2006	Identifying Novel Repolarization Genes: Genetic Dissection of QT Interval in Population Studies / Cardiology Grand Rounds Department of Medicine, Massachusetts General Hospital, Boston, MA

- 2006 Approaches to the Genetic Dissection of Complex Traits / Epidemiology Core Curriculum
Cardiology Fellowship, Massachusetts General Hospital, Boston, MA
- 2006 Genetic dissection of QT interval variation: defining the genetic determinants of sudden
cardiac death & drug-induced arrhythmias / Lab Group Retreat
Altshuler, Burt, Daly, Hirschhorn lab group, Harvard Medical School
- 2008 Defining the genetic basis of QT interval duration: implications for drug toxicity and
sudden cardiac death / Anesthesia Grand Rounds
Anesthesiology Department, Massachusetts General Hospital, Boston, MA
- 2009 Harnessing genetics to understand blood pressure physiology / Medicine Grand Rounds
Department of Medicine, Massachusetts General Hospital, Boston, MA
- 2009 Harnessing genetics to understand blood pressure physiology / Cardiology Grand Rounds
Cardiology Division, Massachusetts General Hospital, Boston, MA
- 2010 Pharmacogenomics / One-Day Primer on Human Genetics Symposium
Clinical Research Program, Massachusetts General Hospital, Boston, MA
- 2010 Insights into the genetic determinants of blood pressure / Research Seminar
Center for Human Genetic Research, Massachusetts General Hospital, Boston, MA
- 2010 Harnessing human genetics to understand myocardial repolarization and cardiotoxic drug
response / Research Conference
Center for Life Sciences, Beth Israel Deaconess Medical Center, Boston, MA
- 2010 Harnessing human genetics to identify novel physiologic determinants of hypertension /
Renal Grand Rounds
Brigham and Women's Hospital, Boston, MA
- 2011 Translating Genetic Variants Influencing Blood Pressure to the Patient / Invited Speaker
Heart Failure Research Conference, Massachusetts General Hospital, Boston, MA
- 2011 Torsade de pointes and polymorphic ventricular tachycardia / Invited Speaker
ECG Session: TdP and PMVT, Massachusetts General Hospital, Boston, MA
- 2011 Lessons from the Study of Blood Pressure Genetics / Invited Speaker
A Primer on Complex Trait Genetics: Basic Principles for the Beginning Investigator,
Massachusetts General Hospital, Boston, MA
- 2011 Harnessing human genetics to identify novel blood pressure pathways / Invited Speaker
Cardiology Translational Research Seminar, Massachusetts General Hospital, Boston, MA
- 2012 Harnessing genetics to improve our understanding of blood pressure regulation / Invited
Speaker
Center for Human Genetic Research, Massachusetts General Hospital, Boston, MA
- 2012 Harnessing Genetics to Improve Our Understanding of Blood Pressure / Invited Speaker
Cardiovascular Research Center Retreat, American Academy of Arts & Sciences,
Cambridge, MA
- 2013 Genetics establishes a significant role for cGMP pathways in human blood pressure
regulation / Invited Speaker
Anesthesia Research Conference, Massachusetts General Hospital, Boston, MA
- 2013 Harnessing genetics to improve our understanding of blood pressure regulation / Invited
Speaker
Center for Human Genetic Research Retreat, Broad Institute, Cambridge, MA
- 2013 Torsade de pointes and polymorphic ventricular tachycardia / Invited Speaker
Cardiology Fellows Seminar, Massachusetts General Hospital, Boston, MA
- 2014 Genetic studies of blood pressure regulation converge on cGMP-regulating pathways /
Invited Speaker
Heart Failure/Transplant Seminar, Massachusetts General Hospital, Boston, MA

Regional

- 2003 Candidate Gene Association Study of Risk of Myocardial Infarction / Journal Club
Framingham Heart Study, Framingham, MA
- 2003 Genetics of QT Interval Variation in the General Population / Research Conference
Program in Medical and Population Genetics, Whitehead Institute Center for Genome
Research, Cambridge, MA
- 2003 Heritability of Electrocardiographic and Echocardiographic Measures of Left Ventricular
Hypertrophy / Research Conference
Framingham Heart Study, Framingham, MA
- 2004 The role of 5-Lipoxygenase Activating Protein and Risk of Myocardial Infarction and
Stroke / Journal Club
Framingham Heart Study, Framingham, MA
- 2005 Association of Genetic Variation in KCNH2 with QT Interval Variation / Research
Conference
Framingham Heart Study, Framingham, MA
- 2005 Conserved non-coding regions represent functional genetic sequence under negative
selection / Research Conference
Boston University School of Public Health, Program in Statistical Genetics,
- 2010 Harnessing human genetics to understand cardiovascular physiology: QT interval duration
/ Research Conference
Cardiovascular Research Center Seminars, Brown University, Providence, RI
- 2012 GWAS of the Electrocardiogram: Implications for SCD Syndromes / Invited Speaker
Heart Rhythm Society Scientific Sessions, Boston, MA
- 2012 Harnessing genetics to improve our understanding of blood pressure regulation / Invited
Speaker
Framingham Heart Study Research Conference, Framingham, MA
- 2013 The Genetic Basis of Drug-induced Arrhythmia / Invited Speaker
Beth Israel Deaconess Medical Center Electrophysiology Section Seminar, Boston, MA
- 2014 What have genome-wide association studies taught us about cGMP and blood pressure
regulation? / Invited Speaker
Ironwood Pharmaceuticals, Cambridge, MA
- 2014 What has human genetics taught us about the fundamental mechanisms underlying heart
failure? / Invited Speaker
Novartis Institutes for BioMedical Research, Cambridge, MA

National

- 2004 Genetic Determinants of Variation in Electrocardiographic Measures / Research
Conference
Laboratory of Biochemical Genetics and Metabolism, Rockefeller University, New York,
NY
- 2005 Genomic studies in the Framingham population / Research Seminar
Cardiovascular Division, Johns Hopkins School of Medicine/Reynolds Center, Baltimore,
MD
- 2006 Genetic determinants of the QT interval / Research Seminar
Center for Human Genomics, Wake Forest University School of Medicine, Charlotte, NC
- 2007 Defining the genetic basis of electrocardiographic QT interval duration and sudden cardiac
death / Research Conference

- Seminars in Clinical Research, Rockefeller University, New York, NY
- 2007 Genetic dissection of QT interval variation and sudden cardiac death / Research Seminar
Jackson Heart Study, University of Mississippi Medical Center, Jackson, MI
- 2007 The genetic dissection of QT interval duration: implications for drug toxicity and sudden death risk assessment / Research Seminar
Seminars in Integrative Genomics, Vanderbilt University Medical Center, Nashville, TN
- 2007 The genetic dissection of myocardial repolarization and sudden cardiac death / Research Seminar
Department of Preventive Medicine, Feinberg School of Medicine, Northwestern University, Chicago, IL
- 2008 Genetic determinants of QT interval and sudden death / Grand Rounds
Cardiology Division, MD Anderson
- 2008 QTGEN: Meta-analysis of genome-wide association studies of QT interval duration in 13,109 individuals / Research Conference
CHARGE Consortium Conference, Department of Epidemiology, University of Washington, Seattle
- 2009 Late-Breaking Developments in the Genetics of Blood Pressure Regulation and Hypertension / Late-breaking Research Conference
American Society of Hypertension, San Francisco, CA
- 2009 The Genetic Determinants of QT-interval Variation / International Congress on Genetics/Genomics
American Heart Association
- 2009 Integrative Genomic Studies in Humans at the FHS / Scientific Sessions
American Heart Association, Orlando, FL
- 2010 What can we learn from common genetic variants about sudden cardiac death risk? / Scientific Sessions
Heart Rhythm Society, Denver, CO
- 2011 Pharmacogenetics of QT Interval and Potential Clinical Applications to Reduce Off-Target Drug Toxicity / Invited Speaker
NHLBI and PMC New Frontiers in Personalized Medicine: Cardiovascular Research and Clinical Care Meeting, George Washington University, Washington, DC
- 2011 Genome Wide Association Studies - Hope or Hype / Invited Speaker
Heart Rhythm Society Scientific Sessions, San Francisco, CA
- 2011 Genetics establishes a significant role for cGMP pathways in human blood pressure regulation / Invited Speaker
Merck Research Laboratories Seminar, Rathway, NJ
- 2011 Whole exome sequencing of blood pressure extremes / Invited Speaker
American Heart Association Scientific Sessions, Orlando, FL
- 2011 Genetic markers of prognosis: What have we learned from studies of blood pressure? / Invited Speaker
American Heart Association Scientific Sessions, Orlando, FL
- 2012 Harnessing genetics to improve our understanding of blood pressure regulation / Invited Speaker
Northwestern University, Cardiology Grand Rounds, Chicago, IL
- 2012 The genetic basis of arrhythmias / Invited Speaker
University of Rochester, Cardiology Grand Rounds, Rochester, NY
- 2012 From Disease Variants to Disease Mechanism: Progress and Challenges / Invited Speaker
American Heart Association Scientific Sessions, Los Angeles, CA

- 2013 Harnessing genetics to identify novel therapeutic targets and cardiotoxic off-target effects/
Invited Speaker
Regeneron Pharmaceuticals, Tarrytown, NY
- 2013 Genomic Studies of the Electrocardiogram: New Insights into Arrhythmia Mechanisms /
Invited Speaker
Heart Rhythm Society Scientific Sessions, Denver, CO
- 2013 Genomics of Sudden Cardiac Death / Invited Speaker
Heart Rhythm Society Scientific Sessions, Denver, CO

International

- 2006 Genetic dissection of the QT interval / Invited Seminar
Department of Epidemiology and Biostatistics Erasmus Medical Centre, Rotterdam,
Netherlands
- 2006 Genetic dissection of QT interval variation / Invited Seminar
Folkhälsan-Östanlid Hospital, Jakobstad, Finland
- 2006 Genetic dissection of QT interval variation and sudden cardiac death / Invited Seminar
Diabetes and Endocrinology Research Unit, Lund University, Malmö, Sweden
- 2008 The genetic determinants of electrocardiographic QT interval variation and ventricular
arrhythmias / Invited Seminar
Department of Epidemiology, Erasmus Medical Centre, Rotterdam, Netherlands
- 2009 The genetic dissection of complex cardiovascular traits: blood pressure and
electrocardiographic QT interval / Invited Speaker
William Harvey Research Institute at Barts and the London School of Medicine and
Dentistry, London, England
- 2009 The genetic basis of QT interval and blood pressure variation / Invited Seminar
University Medical Center, Groningen, Netherlands
- 2010 Genome-wide association of blood pressure and hypertension / Invited Talk
Toronto Heart Summit, University of Toronto, Canada
- 2010 Harnessing genetics to find determinants of sudden cardiac death and cardiotoxic drug
response / Invited seminar
National Institute of Health and Welfare, Helsinki, Finland
- 2011 Genetics establishes a significant role for cGMP pathways in human blood pressure
regulation/ Invited speaker
5th International Conference on cGMP, Halle, Germany
- 2012 Harnessing genetics to improve our understanding of blood pressure and arrhythmias /
Invited Speaker
University of Glasgow, Glasgow, Scotland
- 2013 What have genome-wide association studies taught us about cGMP and blood pressure
regulation? / Invited Speaker
6th International Conference on cGMP, Erfurt, Germany
- 2013 Harnessing genetics to identify novel therapeutic targets and cardiotoxic off-target effects /
Invited Speaker
Nature Genetics/Wellcome Trust Genomics of Common Disease conference, Oxford, UK

Current Licensure and Certification

- 1998 Board of Registration in Medicine, Massachusetts (Full license)
- 2003 ABIM-Cardiovascular Disease (Certified)
- 2012 ABIM-Advanced Heart Failure/Transplant Cardiology (Certified)

Practice Activities

2004	Attending Physician, Cardiology	Heart Failure and Transplantation Section, Massachusetts General Hospital	Heart failure clinic and inpatient care of advanced heart failure patients
------	---------------------------------	---	--

Technological and Other Scientific Innovations

MGH 3691: EP 08166038.3	Use of Adrenomedullin for Cardiovascular Risk Prediction, filed October 07, 2008; licensed to Brahms AG (now ThermoFischer) In this patent my colleagues and I are co-inventors described the use of adrenomedullin as a predictive biomarker of future cardiovascular events at the population level.
MGH 20275: EP 08268096.9	Arginine Vasopressin Pro-Hormone as Predictive Biomarker for Diabetes, filed October 31, 2008; licensed to Brahms AG (now ThermoFischer) In this patent, my colleagues and I are co-inventors of the use of Arginine vasopressin pro-hormone as a predictive biomarker of future diabetes status at the population level.
MGH provisional	INHIBITORS OF MICRORNAs THAT REGULATE PRODUCTION OF ATRIAL NATUERETIC PEPTIDE (ANP) AS THERAPEUTICS AND USES THEREOF In this patent, my colleagues and I are co-inventors of of the use of microRNA inhibitors to augment atrial natriuretic peptide activity in hypertension and heart failure.

Education of Patients and Service to the Community

Activities

1996	Native Alaskan/Indian Health Service rotations / Medical Student Two month rotation in Indian Health Service hospital in rural Alaska
2007-	Lexington, MA Medical Reserve Corps, Department of Public Health / Executive Committee Contribute to disaster planning and resource allocation at town level

Publications - Peer reviewed publications in print or other media

Note: surname changed from Newton to Newton-Cheh in 1999

1. Polanczyk CA, **Newton C**, Dec GW, Di Salvo TG. Quality of care and hospital readmission in congestive heart failure: An explicit review process. *J Card Fail.* 2001 Dec; 7(4): 289-98.
2. **Newton-Cheh C**, Larson MG, Corey DC, Benjamin EJ, Herbert AG, Levy D, D'Agostino RB, O'Donnell CJ. QT interval is a heritable quantitative trait with evidence of linkage to chromosome 3 in a genome-wide linkage analysis: The Framingham Heart Study. *Heart Rhythm.* 2005 Mar; 2(3): 277-84.
3. Kathiresan S, Larson MG, Vasani RS, Guo CY, Vita JA, Mitchell GF, Keyes MJ, **Newton-Cheh C**, Musone SL, Lochner AL, Drake JA, Levy D, O'Donnell CJ, Hirschhorn JN, Benjamin EJ. Common genetic variation at the endothelial nitric oxide synthase locus and relations to brachial artery

- vasodilator function in the community. *Circulation*. 2005 Sep 6; 112(10):1419-27.
4. Drake JA, Bird C, Nemesh J, Thomas DJ, **Newton-Cheh C**, Reymond A, Excoffier L, Attar H, Antonarakis SE, Dermitzakis ET, Hirschhorn JN. Conserved noncoding sequences are selectively constrained and not mutation cold spots. *Nature Genetics*. 2006 Feb; 38(2): 223-227.
 5. Kathiresan S, Gona P, Larson MG, Vita JA, Mitchell GF, Tofler GH, Levy D, **Newton-Cheh C**, Wang TJ, Benjamin EJ, Vasani RS. Cross-sectional relations of multiple biomarkers from distinct biological pathways to brachial artery endothelial function. *Circulation*. 2006 Feb 21; 113(7): 938-45.
 6. Kathiresan S, Larson MG, Vasani RS, Guo CY, Gona P, Keaney JF Jr, Wilson PW, **Newton-Cheh C**, Musone SL, Camargo AL, Drake JA, Levy D, O'Donnell CJ, Hirschhorn JN, Benjamin EJ. Contribution of Clinical Correlates and 13 C-reactive protein Gene Polymorphisms to Interindividual Variability in Serum C-Reactive Protein Level. *Circulation*. 2006 Mar 21; 113(11):1415-23.
 7. Arking DE, Pfeufer A, Post W, Kao WH, **Newton-Cheh C**, Ikeda M, West K, Kashuk C, Akyol M, Perz S, Jalilzadeh S, Illig T, Gieger C, Guo CY, Larson MG, Wichmann HE, Marban E, O'Donnell CJ, Hirschhorn JN, Kaab S, Spooner PM, Meitinger T, Chakravarti A. A common genetic variant in the NOS1 regulator NOS1AP modulates cardiac repolarization. *Nature Genetics*. 2006 Jun; 38(6): 644-651.
 8. Wang TJ, Gona P, Larson MG, Tofler GH, Levy D, **Newton-Cheh C**, Jacques PF, Rifai N, Selhub J, Robins SJ, Benjamin EJ, D'Agostino RB, Vasani RS. Multiple biomarkers for the prediction of first major cardiovascular events and death. *N Engl J Med*. 2006 Dec 21; 355(25): 2631-9.
 9. Cupples LA, Arruda HT, Benjamin EJ, D'Agostino RB Sr, Demissie S, Destefano AL, Dupuis J, Falls KM, Fox CS, Gottlieb DJ, Govindaraju DR, Guo CY, Heard-Costa NL, Hwang SJ, Kathiresan S, Kiel DP, Laramie JM, Larson MG, Levy D, Liu CY, Lunetta KL, Mailman MD, Manning AK, Meigs JB, Murabito JM, **Newton-Cheh C**, O'Connor GT, O'Donnell CJ, Pandey M, Seshadri S, Vasani RS, Wang ZY, Wilk JB, Wolf PA, Yang Q, Atwood LD. The Framingham Heart Study 100K SNP genome-wide association study resource: overview of 17 phenotype working group reports. *BMC Med Genet*. 2007 Sep 19;8 Suppl 1:S1. PMID: PMC1995613.
 10. Wang TJ, Gona P, Larson MG, Levy D, Benjamin EJ, Tofler GH, Jacques PF, Meigs JB, Rifai N, Selhub J, Robins SJ, **Newton-Cheh C**, Vasani RS. Multiple biomarkers and the risk of incident hypertension. *Hypertension*. 2007 Mar; 49(3): 432-8.
 11. **Newton-Cheh C**, Guo CY, Gona P, Larson MG, Benjamin EJ, Wang TJ, Kathiresan S, O'Donnell CJ, Musone SL, Camargo AL, Drake JA, Levy D, Hirschhorn JN, Vasani RS. Clinical and Genetic Correlates of Ambulatory Serum Aldosterone to Renin Ratio and Relations to Blood Pressure Progression and Incidence of Hypertension: The Framingham Heart Study. *Hypertension*. 2007 Apr; 49(4): 846-56.
 12. Diabetes Genetics Initiative of Broad Institute of Harvard and MIT, Lund University, and Novartis Institutes of BioMedical Research, Saxena R, Voight BF, Lyssenko V, Burt NP, de Bakker PI, Chen H, Roix JJ, Kathiresan S, Hirschhorn JN, Daly MJ, Hughes TE, Groop L, Altshuler D, Almgren P, Florez JC, Meyer J, Ardlie K, Bengtsson Bostrom K, Isomaa B, Lettre G, Lindblad U, Lyon HN, Melander O, **Newton-Cheh C**, Nilsson P, Orho-Melander M, Rastam L, Speliotes EK, Taskinen MR,

- Tuomi T, Guiducci C, Berglund A, Carlson J, Gianniny L, Hackett R, Hall L, Holmkvist J, Laurila E, Sjogren M, Sterner M, Surti A, Svensson M, Svensson M, Tewhey R, Blumenstiel B, Parkin M, Defelice M, Barry R, Brodeur W, Camarata J, Chia N, Fava M, Gibbons J, Handsaker B, Healy C, Nguyen K, Gates C, Sougnez C, Gage D, Nizzari M, Gabriel SB, Chirn GW, Ma Q, Parikh H, Richardson D, Ricke D, Purcell S. Genome-Wide Association Analysis Identifies Loci for Type 2 Diabetes and Triglyceride Levels. *Science*. 2007 Jun 1; 316(5829):1331-6.
13. Mitchell GF, Guo CY, Kathiresan S, Vasani RS, Larson MG, Vita JA, Keyes MJ, Vyas M, **Newton-Cheh C**, Musone SL, Camargo AL, Drake JA, Levy D, O'Donnell CJ, Hirschhorn JN, Benjamin EJ. Vascular Stiffness and Genetic Variation at the Endothelial Nitric Oxide Synthase Locus. The Framingham Heart Study. *Hypertension*. 2007 Jun; 49(6):1285-90.
 14. Ingelsson E, Larson MG, Vasani RS, O'Donnell CJ, Yin X, Hirschhorn JN, **Newton-Cheh C**, Drake JA, Musone SL, Heard-Costa NL, Benjamin EJ, Levy D, Atwood LD, Wang TJ, Kathiresan S. Heritability, linkage, and genetic associations of exercise treadmill test responses. *Circulation*. 2007 Jun 12; 115(23): 2917-24.
 15. Aarnoudse AJLH*, **Newton-Cheh C***, de Bakker PIW, Straus SMJM, Kors JA, Hofman A, Uitterlinden AG, Witteman JCM, Stricker BHC. Common NOS1AP variants are associated with a prolonged QTc interval in the Rotterdam Study. *Circulation*. 2007 Jul 3; 116(1):10-6.
 16. **Newton-Cheh C**, Guo CY, Larson MG, Musone SL, Surti A, Camargo AL, Drake JA, Benjamin EJ, Levy D, D'Agostino RB Sr, Hirschhorn JN, O'Donnell CJ. Common Genetic Variation in KCNH2 Is Associated With QT Interval Duration: The Framingham Heart Study. *Circulation*. 2007 Sep 4; 116(10):1128-36.
 17. Larson MG, Atwood LD, Benjamin EJ, Cupples LA, D'Agostino RB Sr, Fox CS, Govindaraju DR, Guo CY, Heard-Costa NL, Hwang SJ, Murabito JM, **Newton-Cheh C**, O'Donnell CJ, Seshadri S, Vasani RS, Wang TJ, Wolf PA, Levy D. Framingham Heart Study 100K project: genome-wide associations for cardiovascular disease outcomes. *BMC Med Genet*. 2007 Sep 19; 8 Suppl 1:S5. PMID: PMC1995607.
 18. Levy D, Larson MG, Benjamin EJ, **Newton-Cheh C**, Wang TJ, Hwang SJ, Vasani RS, Mitchell GF. Framingham Heart Study 100K Project: genome-wide associations for blood pressure and arterial stiffness. *BMC Med Genet*. 2007 Sep 19; 8 Suppl 1:S3. PMID: PMC1995621.
 19. Vasani RS, Larson MG, Aragam J, Wang TJ, Mitchell GF, Kathiresan S, **Newton-Cheh C**, Vita JA, Keyes MJ, O'Donnell CJ, Levy D, Benjamin EJ. Genome-wide association of echocardiographic dimensions, brachial artery endothelial function and treadmill exercise responses in the Framingham Heart Study. *BMC Med Genet*. 2007. Sep 19; 8 Suppl 1:S2. PMID: PMC1995617.
 20. **Newton-Cheh C**, Guo CY, Wang TJ, O'donnell CJ, Levy D, Larson MG. Genome-wide association study of electrocardiographic and heart rate variability traits: the Framingham Heart Study. *BMC Med Genet*. 2007 Sep 19;8 Suppl 1:S7. PMID: PMC1995612
 21. Parikh NI, Gona P, Larson MG, Wang TJ, **Newton-Cheh C**, Levy D, Benjamin EJ, Kannel WB, Vasani RS. Plasma renin and risk of cardiovascular disease and mortality: the Framingham Heart Study. *Eur Heart J*. 2007. Nov; 28(21): 2644-52.

22. Kathiresan S, Melander O, Guiducci C, Surti A, Burt NP, Rieder MJ, Cooper GM, Roos C, Voight BF, Havulinna AS, Wahlstrand B, Hedner T, Corella D, Tai ES, Ordovas JM, Berglund G, Vartiainen E, Jousilahti P, Hedblad B, Taskinen MR, **Newton-Cheh C**, Salomaa V, Peltonen L, Groop L, Altshuler DM, Orho-Melander M. Six new loci associated with blood low-density lipoprotein cholesterol, high-density lipoprotein cholesterol or triglycerides in humans. *Nature Genetics*. 2008. Feb; 40(2):189-97. PMID: PMC2682493.
23. Kathiresan S, Melander O, Anevski D, Guiducci C, Burt NP, Roos C, Hirschhorn JN, Berglund G, Hedblad B, Groop L, Altshuler DM, **Newton-Cheh C**, Orho-Melander M. Polymorphisms associated with cholesterol and risk of cardiovascular events. *N Engl J Med*. 2008. Mar 20; 358(12):1240-9.
24. Lehtinen AB*, **Newton-Cheh C***, Ziegler JT, Langefeld CD, Freedman BI, Daniel KR, Herrington DM, Bowden DW. Association of NOS1AP genetic variants with QT interval duration in families from the Diabetes Heart Study. *Diabetes*. 2008; 57(4): 1108-1114.
25. Ji W, Foo JN, O'Roak BJ, Zhao H, Larson MG, Simon DB, **Newton-Cheh C**, State MW, Levy D, Lifton RP. Rare independent mutations in renal salt handling genes contribute to blood pressure variation. *Nature Genetics*. 2008 May; 40(5): 592-9. PMID: PMC3766631
26. Schnabel R, Larson MG, Dupuis J, Lunetta KL, Lipinska I, Meigs JB, Yin X, Rong J, Vita JA, **Newton-Cheh C**, Levy D, Keaney JF, Jr., Vasan RS, Mitchell GF, Benjamin EJ. Relations of inflammatory biomarkers and common genetic variants with arterial stiffness and wave reflection. *Hypertension*. 2008; 51(6): 1651-1657. PMID: PMC2892983.
27. Becker ML, Aarnoudse AJ, **Newton-Cheh C**, Hofman A, Witteman JC, Uitterlinden AG, Visser LE, Stricker BH. Common variation in the NOS1AP gene is associated with reduced glucose-lowering effect and with increased mortality in users of sulfonylurea. *Pharmacogenet Genomics*. 2008 Jul; 18(7): 591-597.
28. Lieb W, Pencina MJ, Wang TJ, Larson MG, Lanier KJ, Benjamin EJ, Levy D, Tofler GH, Meigs JB, **Newton-Cheh C**, Vasan RS. Association of Parental Hypertension with Concentrations of Select Biomarkers in Nonhypertensive Offspring. *Hypertension*. 2008 Aug; 52(2): 381-6. PMID: PMC2574605.
29. Becker ML, Visser LE, **Newton-Cheh C**, Witteman JC, Hofman A, Uitterlinden AG, Stricker BH. Genetic variation in the NOS1AP gene is associated with the incidence of diabetes mellitus in users of calcium channel blockers. *Diabetologia*. 2008 Nov; 51(11): 2138-40. Epub 2008 Sep 3. PMID: PMC2668085.
30. Becker ML, Visser LE, **Newton-Cheh C**, Hofman A, Uitterlinden AG, Witteman JC, Stricker BH. A common NOS1AP genetic polymorphism is associated with increased cardiovascular mortality in users of dihydropyridine calcium channel blockers. *Br J Clin Pharmacol*. 2009 Jan; 67(1): 61-7. PMID: PMC2668085.
31. Marjamaa A*, **Newton-Cheh C***, Porthan K, Reunanen A, Lahermo P, Väänänen H, Jula A, Karanko H, Swan H, Toivonen L, Nieminen MS, Viitasalo M, Peltonen L, Oikarinen L, Palotie A, Kontula K, Salomaa V. Common candidate gene variants are associated with QT interval duration in the general

- population. *J Intern Med.* 2009 Apr; 265(4): 448-58. PMID: PMC2704397.
32. van Noord C, Straus SM, Sturkenboom MC, Hofman A, Aarnoudse AJ, Bagnardi V, Kors JA, **Newton-Cheh C**, Witteman JC, Stricker BH. Psychotropic drugs associated with corrected QT interval prolongation. *J Clin Psychopharmacol.* 2009 Feb; 29(1):9-15.
33. Lowe JK, Maller JB, Pe'er I, Neale BM, Salit J, Kenny EE, Shea JL, Burkhardt R, Smith JG, Ji W, Noel M, Foo JN, Blundell ML, Skilling V, Garcia L, Sullivan ML, Lee HE, Labek A, Ferdowsian H, Auerbach SB, Lifton RP, **Newton-Cheh C**, Breslow JL, Stoffel M, Daly MJ, Altshuler DM, Friedman JM. Genome-wide association studies in an isolated founder population from the Pacific Island of Kosrae. *PLoS Genet.* 2009 Feb; 5(2): e1000365. PMID: PMC2628735.
34. Schnabel RB, Sullivan LM, Levy D, Pencina MJ, Massaro JM, D'Agostino RB Sr, **Newton-Cheh C**, Yamamoto JF, Magnani JW, Tadros TM, Kannel WB, Wang TJ, Ellinor PT, Wolf PA, Vasani RS, Benjamin EJ. Development of a risk score for atrial fibrillation (Framingham Heart Study): a community-based cohort study. *Lancet.* 2009. Feb 28; 373(9665): 739-45. PMID: PMC2764235.
35. **Newton-Cheh C**, Larson MG, Ramachandran VS, Levy D, Bloch KD, Surti A, Guiducci C, Kathiresan S, Benjamin EJ, Struck J, Morgenthaler NG, Bergmann A, Blankenberg S, Kee F, Nilsson P, Yin X, Peltonen L, Vartiainen E, Salomaa V, Hirschhorn JN, Melander O, Wang TJ. Association of Common Variants in NPPA and NPPB with Circulating Natriuretic Peptides and Blood Pressure. *Nature Genetics.* 2009; 41(3): 348-53. PMID: PMC2664511.
36. **Newton-Cheh C***, Eijgelsheim M*, Rice KM*, de Bakker PI*, Yin X, Estrada K, Bis JC, Marciante K, Rivadeneira F, Noseworthy PA, Sotoodehnia N, Smith NL, Rotter JI, Kors JA, Witteman JC, Hofman A, Heckbert SR, O'Donnell CJ, Uitterlinden AG, Psaty BM, Lumley T, Larson MG, Ch Stricker BH. Common variants at ten loci influence QT interval duration in the QTGEN Study. *Nature Genetics.* 2009; 41(4): 399-406. PMID: PMC2701449.
37. Marjamaa A, Salomaa V, **Newton-Cheh C**, Porthan K, Reunanen A, Karanko H, Jula A, Lahermo P, Väänänen H, Toivonen L, Swan H, Viitasalo M, Nieminen MS, Peltonen L, Oikarinen L, Palotie A, Kontula K. High prevalence of four long QT syndrome founder mutations in the Finnish population. *Ann Med.* 2009; 41(3): 234-40. PMID: PMC2668713.
38. Kaab S, Darbar D, van Noord C, Dupuis J, Pfeufer A, **Newton-Cheh C**, Schnabel R, Makino S, Sinner MF, Kannankeril PJ, Beckmann BM, Choudry S, Donahue BS, Heeringa J, Perz S, Lunetta KL, Larson MG, Levy D, MacRae CA, Ruskin JN, Wacker A, Schomig A, Wichmann HE, Steinbeck G, Meitinger T, Uitterlinden AG, Witteman JC, Roden DM, Benjamin EJ, Ellinor PT. Large scale replication and meta-analysis of variants on chromosome 4q25 associated with atrial fibrillation. *Eur Heart J.* 2009; 30(7): 813-819. PMID: PMC2663727.
39. van Noord C, Aarnoudse AJ, Eijgelsheim M, Sturkenboom MC, Straus SM, Hofman A, Kors JA, **Newton-Cheh C**, Witteman JC, Stricker BH. Calcium channel blockers, NOS1AP, and heart-rate-corrected QT prolongation. *Pharmacogenet Genomics.* 2009 Apr; 19(4): 260-266.
40. Smith JG, Lowe JK, Kovvali S, Maller JB, Salit J, Daly MJ, Stoffel M, Altshuler DM, Friedman JM, Breslow JL, **Newton-Cheh C**. Genome-wide association study of electrocardiographic conduction measures in an isolated founder population: Kosrae. *Heart Rhythm.* 2009; 6(5): 634-641. PMID:

41. **Newton-Cheh C***, Johnson T*, Gateva V*, Tobin MD*, Bochud M, Coin L, Najjar SS, Zhao JH, Heath SC, Eyheramendy S, Papadakis K, Voight BF, Scott LJ, Zhang F, Farrall M, Tanaka T, Wallace C, Chambers JC, Khaw KT, Nilsson P, van der Harst P, Polidoro S, Grobbee DE, Onland-Moret NC, Bots ML, Wain LV, Elliott KS, Teumer A, Luan J, Lucas G, Kuusisto J, Burton PR, Hadley D, McArdle WL; Wellcome Trust Case Control Consortium, Brown M, Dominiczak A, Newhouse SJ, Samani NJ, Webster J, Zeggini E, Beckmann JS, Bergmann S, Lim N, Song K, Vollenweider P, Waeber G, Waterworth DM, Yuan X, Groop L, Orho-Melander M, Allione A, Di Gregorio A, Guarrera S, Panico S, Ricceri F, Romanazzi V, Sacerdote C, Vineis P, Barroso I, Sandhu MS, Luben RN, Crawford GJ, Jousilahti P, Perola M, Boehnke M, Bonnycastle LL, Collins FS, Jackson AU, Mohlke KL, Stringham HM, Valle TT, Willer CJ, Bergman RN, Morken MA, Döring A, Gieger C, Illig T, Meitinger T, Org E, Pfeufer A, Wichmann HE, Kathiresan S, Marrugat J, O'Donnell CJ, Schwartz SM, Siscovick DS, Subirana I, Freimer NB, Hartikainen AL, McCarthy MI, O'Reilly PF, Peltonen L, Pouta A, de Jong PE, Snieder H, van Gilst WH, Clarke R, Goel A, Hamsten A, Peden JF, Seedorf U, Syvänen AC, Tognoni G, Lakatta EG, Sanna S, Scheet P, Schlessinger D, Scuteri A, Dörr M, Ernst F, Felix SB, Homuth G, Lorbeer R, Reffelmann T, Rettig R, Völker U, Galan P, Gut IG, Hercberg S, Lathrop GM, Zelenika D, Deloukas P, Soranzo N, Williams FM, Zhai G, Salomaa V, Laakso M, Elosua R, Forouhi NG, Völzke H, Uiterwaal CS, van der Schouw YT, Numans ME, Matullo G, Navis G, Berglund G, Bingham SA, Kooner JS, Connell JM, Bandinelli S, Ferrucci L, Watkins H, Spector TD, Tuomilehto J, Altshuler D, Strachan DP, Laan M, Meneton P, Wareham NJ, Uda M, Jarvelin MR, Mooser V, Melander O, Loos RJ, Elliott P, Abecasis GR, Caulfield M, Munroe PB. Genome-wide association study identifies eight loci associated with blood pressure. *Nature Genetics*. 2009; 41: 666-676. PMID: PMC2891673.
42. Cheng S, Keyes MJ, Larson MG, McCabe EL, **Newton-Cheh C**, Levy D, Benjamin EJ, Vasani RS, Wang TJ. Long-term outcomes in individuals with prolonged PR interval or first-degree atrioventricular block. *JAMA*. 2009; 301: 2571-7. PMID: PMC2765917.
43. Melander O, **Newton-Cheh C**, Almgren P, Hedblad B, Berglund G, Engström G, Persson M, Smith JG, Magnusson M, Christensson A, Struck J, Morgenthaler NG, Bergmann A, Pencina MJ, Wang TJ. Novel and Conventional Biomarkers for Prediction of Incident Cardiovascular Events in the Community. *JAMA*. 2009; 302: 49-57. PMID: PMC3090639.
44. Nolte IM, Wallace C, Newhouse SJ, Waggott D, Fu J, Soranzo N, Gwilliam R, Deloukas P, Savelieva I, Zheng D, Dalageorgou C, Farrall M, Samani NJ, Connell J, Brown M, Dominiczak A, Lathrop M, Zeggini E, Wain LV; Wellcome Trust Case Control Consortium; DCCT/EDIC Research Group, **Newton-Cheh C**, Eijgelsheim M, Rice K, de Bakker PI; QTGEN consortium, Pfeufer A, Sanna S, Arking DE; QTSCD consortium, Asselbergs FW, Spector TD, Carter ND, Jeffery S, Tobin M, Caulfield M, Snieder H, Paterson AD, Munroe PB, Jamshidi Y. Common genetic variation near the phospholamban gene is associated with cardiac repolarisation: meta-analysis of three genome-wide association studies. *PLoS One*. 2009; 4(7): e6138. PMID: PMC2704957.
45. Akylbekova EL, Crow RS, Johnson WD, Buxbaum SG, Njemanze S, Fox E, Sarpong DF, Taylor HA, **Newton-Cheh C**. Clinical correlates and heritability of QT interval duration in blacks: the Jackson Heart Study. *Circ Arrhythmia Electrophysiol*. 2009; 2: 427-432. PMID: PMC2772163.
46. Smith JG, **Newton-Cheh C**, Hedblad B, Struck J, Morgenthaler NG, Bergmann A, Wang TJ,

Melander O. Distribution and Correlates of Midregional Proadrenomedullin in the General Population. *Clin Chem.* 2009; 55:1593-95.

47. Benjamin EJ, Rice KM, Arking DE, Pfeufer A, van Noord C, Smith AV, Schnabel RB, Bis JC, Boerwinkle E, Sinner MF, Dehghan A, Lubitz SA, D'Agostino Sr RB, Lumley T, Ehret GB, Heeringa J, Aspelund T, **Newton-Cheh C**, Larson MG, Marcianti KD, Soliman EZ, Rivadeneira F, Wang TJ, Eiriksdottir G, Levy D, Psaty BM, Li M, Chamberlain AM, Hofman A, Vasani RS, Harris TB, Rotter JI, Kao WH, Agarwal SK, Stricker BH, Wang K, Launer LJ, Smith NL, Chakravarti A, Uitterlinden AG, Wolf PA, Sotoodehnia N, Köttgen A, van Duijn CM, Meitinger T, Mueller M, Perz S, Steinbeck G, Wichmann HE, Lunetta KL, Heckbert SR, Gudnason V, Alonso A, Kääb S, Ellinor PT, Witteman JC. Variants in ZFHX3 are associated with atrial fibrillation in individuals of European ancestry. *Nature Genetics.* 2009; 41: 879-881. PMID: PMC2761746.
48. Body SC, Collard CD, Sherran SK, Fox AA, Liu K, Ritchie MD, Perry TE, Muehlschlegel JD, Aranki S, Donahue BS, Pretorius M, Estrada JC, Ellinor PT, **Newton-Cheh C**, Seidman CE, Herman DS, Lichtner P, Meitinger T, Pfeufer A, Kaab S, Brown NJ, Roden DM, Darbar D. Variation in the 4q25 Chromosomal Locus Predicts Atrial Fibrillation after Coronary Artery Bypass Graft Surgery. *Circulation: Cardiovascular Genetics.* 2009 Oct 1; 2(5): 499-506. PMID: PMC2801871.
49. Eijgelsheim M, **Newton-Cheh C**, Aarnoudse AL, van Noord C, Witteman JC, Hofman A, Uitterlinden AG, Stricker BH. Genetic variation in NOS1AP is associated with sudden cardiac death: evidence from the Rotterdam Study. *Hum Mol Genet.* 2009 Nov 1; 18(21): 4213-8. PMID: PMC2758139.
50. **Newton-Cheh C**, Cook NR, VanDenburgh M, Rimm EB, Ridker PM, Albert CM. A common variant at 9p21 is associated with sudden and arrhythmic cardiac death. *Circulation.* 2009 Nov 24; 120(21): 2062-8. PMID: PMC2785227.
51. Pfeufer A, van Noord C, Marcianti KD, Arking DE, Larson MG, Smith AV, Tarasov KV, Müller M, Sotoodehnia N, Sinner MF, Verwoert GC, Li M, Kao WH, Köttgen A, Coresh J, Bis JC, Psaty BM, Rice K, Rotter JI, Rivadeneira F, Hofman A, Kors JA, Stricker BH, Uitterlinden AG, van Duijn CM, Beckmann BM, Sauter W, Gieger C, Lubitz SA, **Newton-Cheh C**, Wang TJ, Magnani JW, Schnabel RB, Chung MK, Barnard J, Smith JD, Van Wagoner DR, Vasani RS, Aspelund T, Eiriksdottir G, Harris TB, Launer LJ, Najjar SS, Lakatta E, Schlessinger D, Uda M, Abecasis GR, Müller-Myhsok B, Ehret GB, Boerwinkle E, Chakravarti A, Soliman EZ, Lunetta KL, Perz S, Wichmann HE, Meitinger T, Levy D, Gudnason V, Ellinor PT, Sanna S, Kääb S, Witteman JC, Alonso A, Benjamin EJ, Heckbert SR. Genome-wide association study of PR interval. *Nature Genetics.* 2010 Feb; 42(2): 153-9. PMID: PMC2850197.
52. Ellinor PT, Lunetta KL, Glazer NL, Pfeufer A, Alonso A, Chung MK, Sinner MF, de Bakker PI, Mueller M, Lubitz SA, Fox E, Darbar D, Smith NL, Smith JD, Schnabel RB, Soliman EZ, Rice KM, Van Wagoner DR, Beckmann BM, van Noord C, Wang K, Ehret GB, Rotter JI, Hazen SL, Steinbeck G, Smith AV, Launer LJ, Harris TB, Makino S, Nelis M, Milan DJ, Perz S, Esko T, Köttgen A, Moebus S, **Newton-Cheh C**, Li M, Möhlenkamp S, Wang TJ, Kao WH, Vasani RS, Nöthen MM, Macrae CA, Ch Stricker BH, Hofman A, Uitterlinden AG, Levy D, Boerwinkle E, Metspalu A, Topol EJ, Chakravarti A, Gudnason V, Psaty BM, Roden DM, Meitinger T, Wichmann HE, Witteman JC, Barnard J, Arking DE, Benjamin EJ, Heckbert SR, Kääb S. Common variants in KCNN3 are associated with lone atrial fibrillation. *Nature Genetics.* 2010 Mar; 42(3): 240-4. PMID: PMC2871387.

53. Min SS, Turner JR, Nada A, DiMino TL, Hynie I, Kleiman R, Kowey P, Krucoff MW, Mason JW, Phipps A, **Newton-Cheh C**, Pordy R, Strnadova C, Targum S, Uhl K, Finkle J. Evaluation of ventricular arrhythmias in early clinical pharmacology trials and potential consequences for later development. *Am Heart J*. 2010 May; 159(5): 716-29.
54. Enhörning S, Wang TJ, Nilsson PM, Almgren P, Hedblad B, Berglund G, Struck J, Morgenthaler NG, Bergmann A, Lindholm E, Groop L, Lyssenko V, Orho-Melander M, **Newton-Cheh C***, Melander O*. Plasma Copeptin and the Risk of Diabetes Mellitus. *Circulation*. 2010; 121: 2102-2108. PMID: PMC3763235.
55. Albert CM, Macrae CA, Chasman DI, Vandenburgh M, Buring JE, Manson JE, Cook NR, **Newton-Cheh CH**. Common Variants in Cardiac Ion Channel Genes Are Associated with Sudden Cardiac Death. *Circ Arrhythm Electrophysiol*. 2010 Jun 1; 3(3): 222-9. PMID: PMC2891421.
56. Musunuru K, Lettre G, Young T, Farlow DN, Pirruccello JP, Ejebe KG, Keating BJ, Yang Q, Chen MH, Lapchyk N, Crenshaw A, Ziaugra L, Rachupka A, Benjamin EJ, Cupples LA, Fornage M, Fox ER, Heckbert SR, Hirschhorn JN, **Newton-Cheh CH**, Nizzari MM, Paltoo DN, Papanicolaou GJ, Patel SR, Psaty BM, Rader DJ, Redline S, Rich SS, Rotter JI, Taylor HA Jr, Tracy RP, Vasani RS, Wilson JG, Kathiresan S, Fabsitz RR, Boerwinkle E, Gabriel SB. Candidate Gene Association Resource (CARE): Design, Methods, and Proof of Concept. *Circ Cardiovasc Genet*. 2010 Jun 1; 3(3): 267-75. PMID: PMC3048024.
57. Porthan K, Marjamaa A, Viitasalo M, Väänänen H, Jula A, Toivonen L, Nieminen MS, **Newton-Cheh C**, Salomaa V, Kontula K, Oikarinen L. Relationship of common candidate gene variants to electrocardiographic T-wave peak to T-wave end interval and T-wave morphology parameters. *Heart Rhythm*. 2010 Jul; 7(7): 898-903. PMID: PMC2904845.
58. Cheng S, Larson MG, Keyes MJ, McCabe EL, **Newton-Cheh C**, Levy D, Benjamin EJ, Vasani RS, Wang TJ. Relation of QRS width in healthy persons to risk of future permanent pacemaker implantation. *Am J Cardiol*. 2010 Sep 1; 106(5): 668-72. PMID: PMC3012354.
59. Eijgelsheim M, **Newton-Cheh C**, Sotoodehnia N, **de Bakker PI**, Müller M, Morrison AC, Smith AV, Isaacs A, Sanna S, Dörr M, Navarro P, Fuchsberger C, Nolte IM, de Geus EJ, Estrada K, Hwang SJ, Bis JC, Rückert IM, Alonso A, Launer LJ, Hottenga JJ, Rivadeneira F, Noseworthy PA, Rice KM, Perz S, Arking DE, Spector TD, Kors JA, Aulchenko YS, Tarasov KV, Homuth G, Wild SH, Marroni F, Gieger C, Licht CM, Prineas RJ, Hofman A, Rotter JI, Hicks AA, Ernst F, Najjar SS, Wright AF, Peters A, Fox ER, Oostra BA, Kroemer HK, Couper D, Völzke H, Campbell H, Meitinger T, Uda M, Witteman JC, Psaty BM, Wichmann HE, Harris TB, Kääb S, Siscovick DS, Jamshidi Y, Uitterlinden AG, Folsom AR, Larson MG, Wilson JF, Penninx BW, Snieder H, Pramstaller PP, van Duijn CM, Lakatta EG, Felix SB, Gudnason V, Pfeufer A, Heckbert SR, Stricker BH, Boerwinkle E, O'Donnell CJ. Genome-wide association analysis identifies multiple loci related to resting heart rate. *Hum Mol Genet*. 2010 Oct 1; 19(19): 3885-94. doi: 10.1093/hmg/ddq303. Epub 2010 Jul 16. PMID: PMC3657480
60. Rodriguez I, Erdman A, Padhi D, Garnett CE, Zhao H, Targum SL, Balakrishnan S, Strnadova C, Viner N, Geiger MJ, **Newton-Cheh C**, Litwin J, Pugsley MK, Sager PT, Krucoff MW, Finkle JK. Electrocardiographic assessment for therapeutic proteins--scientific discussion. *Am Heart J*. 2010

Oct; 160(4): 627-34. doi: 10.1016/j.ahj.2010.07.001.

61. Dey BR, Chung SS, Spitzer TR, Zheng H, Macgillivray TE, Seldin DC, McAfee S, Ballen K, Attar E, Wang T, Shin J, **Newton-Cheh C**, Moore S, Sanchorawala V, Skinner M, Madsen JC, Semigran MJ. Cardiac Transplantation Followed by Dose-Intensive Melphalan and Autologous Stem-Cell Transplantation for Light Chain Amyloidosis and Heart Failure. *Transplantation*. 2010 Oct 27; 90(8): 905-11. PMID: PMC2964067.
62. Padmanabhan S, Melander O, Johnson T, Di Blasio AM, Lee WK, Gentilini D, Hastie CE, Menni C, Monti MC, Delles C, Laing S, Corso B, Navis G, Kwakernaak AJ, van der Harst P, Bochud M, Maillard M, Burnier M, Hedner T, Kjeldsen S, Wahlstrand B, Sjögren M, Fava C, Montagnana M, Danese E, Torffvit O, Hedblad B, Snieder H, Connell JM, Brown M, Samani NJ, Farrall M, Cesana G, Mancina G, Signorini S, Grassi G, Eyheramendy S, Wichmann HE, Laan M, Strachan DP, Sever P, Shields DC, Stanton A, Vollenweider P, Teumer A, Völzke H, Rettig R, **Newton-Cheh C**, Arora P, Zhang F, Soranzo N, Spector TD, Lucas G, Kathiresan S, Siscovick DS, Luan J, Loos RJ, Wareham NJ, Penninx BW, Nolte IM, McBride M, Miller WH, Nicklin SA, Baker AH, Graham D, McDonald RA, Pell JP, Sattar N, Welsh P; Global BPgen Consortium, Munroe P, Caulfield MJ, Zanchetti A, Dominiczak AF. Genome-wide association study of blood pressure extremes identifies variant near UMOD associated with hypertension. *PLoS Genet*. 2010 Oct 28; 6(10): e1001177. PMID: PMC2965757.
63. Chen-Tournoux A, Khan AM, Baggish AL, Castro VM, Semigran MJ, McCabe EL, Moukarbel G, Reingold J, Durrani S, Lewis GD, **Newton-Cheh C**, Scherrer-Crosbie M, Kaplan LM, Wang TJ. Effect of weight loss after weight loss surgery on plasma N-terminal pro-B-type natriuretic peptide levels. *Am J Cardiol*. 2010 Nov 15; 106(10): 1450-5. doi: 10.1016/j.amjcard.2010.06.076. PMID: PMC3170817.
64. Smith JG, **Newton-Cheh C**, Almgren P, Struck J, Morgenthaler NG, Bergmann A, Platonov PG, Hedblad B, Engström G, Wang TJ, Melander O. Assessment of Conventional Cardiovascular Risk Factors and Multiple Biomarkers for the Prediction of Incident Heart Failure and Atrial Fibrillation. *J. Am. Coll. Cardiol*. 2010; 56: 1712-1719. PMID: PMC3005324.
65. Sotoodehnia N, Isaacs A, de Bakker PI, Dörr M, **Newton-Cheh C**, Nolte IM, van der Harst P, Müller M, Eijgelsheim M, Alonso A, Hicks AA, Padmanabhan S, Hayward C, Smith AV, Polasek O, Giovannone S, Fu J, Magnani JW, Marcianti KD, Pfeufer A, Gharib SA, Teumer A, Li M, Bis JC, Rivadeneira F, Aspelund T, Köttgen A, Johnson T, Rice K, Sie MP, Wang YA, Klopp N, Fuchsberger C, Wild SH, Mateo Leach I, Estrada K, Völker U, Wright AF, Asselbergs FW, Qu J, Chakravarti A, Sinner MF, Kors JA, Petersmann A, Harris TB, Soliman EZ, Munroe PB, Psaty BM, Oostra BA, Cupples LA, Perz S, de Boer RA, Uitterlinden AG, Völzke H, Spector TD, Liu FY, Boerwinkle E, Dominiczak AF, Rotter JI, van Herpen G, Levy D, Wichmann HE, van Gilst WH, Witteman JC, Kroemer HK, Kao WH, Heckbert SR, Meitinger T, Hofman A, Campbell H, Folsom AR, van Veldhuisen DJ, Schwenbacher C, O'Donnell CJ, Volpato CB, Caulfield MJ, Connell JM, Launer L, Lu X, Franke L, Fehrmann RS, te Meerman G, Groen HJ, Weersma RK, van den Berg LH, Wijmenga C, Ophoff RA, Navis G, Rudan I, Snieder H, Wilson JF, Pramstaller PP, Siscovick DS, Wang TJ, Gudnason V, van Duijn CM, Felix SB, Fishman GI, Jamshidi Y, Stricker BH, Samani NJ, Kääb S, Arking DE. Common variants in 22 loci are associated with QRS duration and cardiac ventricular conduction. *Nat Genet*. 2010 Dec; 42(12): 1068-76. doi: 10.1038/ng.716. Epub 2010 Nov 14. PMID: PMC3338195.

66. Min JL, Taylor JM, Richards JB, Watts T, Pettersson FH, Broxholme J, Ahmadi KR, Surdulescu GL, Lowy E, Gieger C, **Newton-Cheh C**, Perola M, Soranzo N, Surakka I, Lindgren CM, Ragoussis J, Morris AP, Cardon LR, Spector TD, Zondervan KT. The Use of Genome-Wide eQTL Associations in Lymphoblastoid Cell Lines to Identify Novel Genetic Pathways Involved in Complex Traits. *PLoS One*. 2011; 6(7):e22070. Epub 2011 Jul 15. PMID: PMC3137612.
67. Smith JG, Magnani JW, Palmer C, Meng YA, Soliman EZ, Musani SK, Kerr KF, Schnabel RB, Lubitz SA, Sotoodehnia N, Redline S, Pfeufer A, Müller M, Evans DS, Nalls MA, Liu Y, Newman AB, Zonderman AB, Evans MK, Deo R, Ellinor PT, Paltoo DN, **Newton-Cheh C**, Benjamin EJ, Mehra R, Alonso A, Heckbert SR, Fox ER; Candidate-gene Association Resource (CARE) Consortium. Genome-Wide Association Studies of the PR Interval in African Americans. *PLoS Genet*. 2011 Feb 10; 7(2): e1001304. PMID: PMC3037415.
68. Lettre G, Palmer CD, Young T, Ejebe KG, Allayee H, Benjamin EJ, Bennett F, Bowden DW, Chakravarti A, Dreisbach A, Farlow DN, Folsom AR, Fornage M, Forrester T, Fox E, Haiman CA, Hartiala J, Harris TB, Hazen SL, Heckbert SR, Henderson BE, Hirschhorn JN, Keating BJ, Kritchevsky SB, Larkin E, Li M, Rudock ME, McKenzie CA, Meigs JB, Meng YA, Mosley TH, Newman AB, **Newton-Cheh CH**, Paltoo DN, Papanicolaou GJ, Patterson N, Post WS, Psaty BM, Qasim AN, Qu L, Rader DJ, Redline S, Reilly MP, Reiner AP, Rich SS, Rotter JI, Liu Y, Shrader P, Siscovick DS, Tang WH, Taylor HA, Tracy RP, Vasani RS, Waters KM, Wilks R, Wilson JG, Fabsitz RR, Gabriel SB, Kathiresan S, Boerwinkle E. Genome-Wide Association Study of Coronary Heart Disease and Its Risk Factors in 8,090 African Americans: The NHLBI CARE Project. *PLoS Genet*. 2011 Feb 10; 7(2): e1001300. PMID: PMC3037413.
69. Kenny EE, Kim M, Gusev A, Lowe JK, Salit J, Smith JG, Kovvali S, Kang HM, **Newton-Cheh C**, Daly MJ, Stoffel M, Altshuler DM, Friedman JM, Eskin E, Breslow JL, Pe'er I. Increased power of mixed models facilitates association mapping of 10 loci for metabolic traits in an isolated population. *Hum Mol Genet*. 2011 Feb 15; 20(4): 827-39. PMID: PMC3024042.
70. Ellis KL, **Newton-Cheh C**, Wang TJ, Frampton CM, Doughty RN, Whalley GA, Ellis CJ, Skelton L, Davis N, Yandle TG, Troughton RW, Richards AM, Cameron VA. Association of genetic variation in the natriuretic peptide system with cardiovascular outcomes. *J Mol Cell Cardiol*. 2011 Apr; 50(4): 695-701.
71. **Newton-Cheh C**, Lin AE, Baggish AL, Wang H. Case records of the Massachusetts General Hospital. Case 11-2011. A 47-year-old man with systemic lupus erythematosus and heart failure. *N Engl J Med*. 2011 Apr 14; 364(15): 1450-1460.
72. Gavin MC, **Newton-Cheh C**, Gaziano JM, Cook NR, VanDenburgh M, Albert CM. A common variant in the β_2 -adrenergic receptor and risk of sudden cardiac death. *Heart Rhythm*. 2011 May; 8(5): 704-10. doi: 10.1016/j.hrthm.2011.01.003. Epub 2011 Jan 6. PMID: PMC3225286.
73. Johnson AD, **Newton-Cheh C**, Chasman DI, Ehret GB, Johnson T, Rose L, Rice K, Verwoert GC, Launer LJ, Gudnason V, Larson MG, Chakravarti A, Psaty BM, Caulfield M, van Duijn CM, Ridker PM, Munroe PB, Levy D; Cohorts for Heart and Aging Research in Genomic Epidemiology Consortium; Global BPgen Consortium; Women's Genome Health Study. Association of hypertension drug target genes with blood pressure and hypertension in 86,588 individuals. *Hypertension*. 2011

May; 57(5):903-10. Epub 2011 Mar 28. PMID: PMC3099407.

74. Noseworthy PA, Tikkanen JT, Porthan K, Oikarinen L, Pietilä A, Harald K, Peloso GM, Merchant FM, Jula A, Väänänen H, Hwang SJ, O'Donnell CJ, Salomaa V, **Newton-Cheh C**, Huikuri HV. The early repolarization pattern in the general population: clinical correlates and heritability. *J Am Coll Cardiol*. 2011 May 31; 57(22): 2284-9. doi: 10.1016/j.jacc.2011.04.003. PMID: PMC3183435.
75. Zhu X, Young JH, Fox E, Keating BJ, Franceschini N, Kang S, Tayo B, Adeyemo A, Sun YV, Li Y, Morrison A, **Newton-Cheh C**, Liu K, Ganesh SK, Kutlar A, Vasani RS, Dreisbach A, Wyatt S, Polak J, Palmas W, Musani S, Taylor H, Fabsitz R, Townsend RR, Dries D, Glessner J, Chiang CW, Mosley T, Kardia S, Curb D, Hirschhorn JN, Rotimi C, Reiner A, Eaton C, Rotter JI, Cooper RS, Redline S, Chakravarti A, Levy D. Combined admixture mapping and association analysis identifies a novel blood pressure genetic locus on 5p13: contributions from the CARE consortium. *Hum Mol Genet*. 2011 Jun 1; 20(11):2285-95. PMID: PMC3090198.
76. Arking DE, Juntila MJ, Goyette P, Huertas-Vazquez A, Eijgelsheim M, Blom MT, **Newton-Cheh C**, Reinier K, Teodorescu C, Uy-Evanado A, Carter-Monroe N, Kaikkonen KS, Kortelainen ML, Boucher G, Lagacé C, Moes A, Zhao X, Kolodgie F, Rivadeneira F, Hofman A, Wittteman JC, Uitterlinden AG, Marsman RF, Pazoki R, Bardai A, Koster RW, Dehghan A, Hwang SJ, Bhatnagar P, Post W, Hilton G, Prineas RJ, Li M, Köttgen A, Ehret G, Boerwinkle E, Coresh J, Kao WH, Psaty BM, Tomaselli GF, Sotoodehnia N, Siscovick DS, Burke GL, Marbán E, Spooner PM, Cupples LA, Jui J, Gunson K, Kesäniemi YA, Wilde AA, Tardif JC, O'Donnell CJ, Bezzina CR, Virmani R, Stricker BH, Tan HL, Albert CM, Chakravarti A, Rioux JD, Huikuri HV, Chugh SS. Identification of a sudden cardiac death susceptibility locus at 2q24.2 through genome-wide association in European ancestry individuals. *PLoS Genet*. 2011 Jun; 7(6):e1002158. PMID: PMC3128111
77. Noseworthy PA, Havulinna AS, Porthan K, Lahtinen AM, Jula A, Karhunen PJ, Perola M, Oikarinen L, Kontula KK, Salomaa V, **Newton-Cheh C**. Common genetic variants, QT interval, and sudden cardiac death in a Finnish population-based study. *Circ Cardiovasc Genet*. 2011 Jun; 4(3):305-11. Epub 2011 Apr 21. PMID: PMC3119024.
78. Noseworthy PA, Weiner R, Kim J, Keelara V, Wang F, Berkstresser B, Wood MJ, Wang TJ, Picard MH, Hutter AM Jr, **Newton-Cheh C**, Baggish AL. Early repolarization pattern in competitive athletes: clinical correlates and the effects of exercise training. *Circ Arrhythm Electrophysiol*. 2011 Aug; 4(4): 432-40. doi: 10.1161/CIRCEP.111.962852. Epub 2011 May 4. PMID: PMC3700366.
79. Lam CS, Cheng S, Choong K, Larson MG, Murabito JM, **Newton-Cheh C**, Bhasin S, McCabe EL, Miller KK, Redfield MM, Vasani RS, Coviello AD, Wang TJ. Influence of sex and hormone status on circulating natriuretic peptides. *J Am Coll Cardiol*. 2011 Aug 2; 58(6):618-26. PMID: PMC3170816.
80. International Consortium for Blood Pressure Genome-Wide Association Studies, Ehret GB, Munroe PB, Rice KM, Bochud M, Johnson AD, Chasman DI, Smith AV, Tobin MD, Verwoert GC, Hwang SJ, Pihur V, Vollenweider P, O'Reilly PF, Amin N, Bragg-Gresham JL, Teumer A, Glazer NL, Launer L, Zhao JH, Aulchenko Y, Heath S, Söber S, Parsa A, Luan J, Arora P, Dehghan A, Zhang F, Lucas G, Hicks AA, Jackson AU, Peden JF, Tanaka T, Wild SH, Rudan I, Igl W, Milaneschi Y, Parker AN, Fava C, Chambers JC, Fox ER, Kumari M, Go MJ, van der Harst P, Kao WH, Sjögren M, Vinay DG, Alexander M, Tabara Y, Shaw-Hawkins S, Whincup PH, Liu Y, Shi G, Kuusisto J, Tayo B, Seielstad

M, Sim X, Nguyen KD, Lehtimäki T, Matullo G, Wu Y, Gaunt TR, Onland-Moret NC, Cooper MN, Platou CG, Org E, Hardy R, Dahgam S, Palmen J, Vitart V, Braund PS, Kuznetsova T, Uiterwaal CS, Adeyemo A, Palmas W, Campbell H, Ludwig B, Tomaszewski M, Tzoulaki I, Palmer ND; CARDIoGRAM consortium; CKDGen Consortium; KidneyGen Consortium; EchoGen consortium; CHARGE-HF consortium, Aspelund T, Garcia M, Chang YP, O'Connell JR, Steinle NI, Grobbee DE, Arking DE, Kardina SL, Morrison AC, Hernandez D, Najjar S, McArdle WL, Hadley D, Brown MJ, Connell JM, Hingorani AD, Day IN, Lawlor DA, Beilby JP, Lawrence RW, Clarke R, Hopewell JC, Ongen H, Dreisbach AW, Li Y, Young JH, Bis JC, Kähönen M, Viikari J, Adair LS, Lee NR, Chen MH, Olden M, Pattaro C, Bolton JA, Köttgen A, Bergmann S, Mooser V, Chaturvedi N, Frayling TM, Islam M, Jafar TH, Erdmann J, Kulkarni SR, Bornstein SR, Grässler J, Groop L, Voight BF, Kettunen J, Howard P, Taylor A, Guarrera S, Ricceri F, Emilsson V, Plump A, Barroso I, Khaw KT, Weder AB, Hunt SC, Sun YV, Bergman RN, Collins FS, Bonnycastle LL, Scott LJ, Stringham HM, Peltonen L, Perola M, Vartiainen E, Brand SM, Staessen JA, Wang TJ, Burton PR, Soler Artigas M, Dong Y, Snieder H, Wang X, Zhu H, Lohman KK, Rudock ME, Heckbert SR, Smith NL, Wiggins KL, Doumatey A, Shriner D, Veldre G, Viigimaa M, Kinra S, Prabhakaran D, Tripathy V, Langefeld CD, Rosengren A, Thelle DS, Corsi AM, Singleton A, Forrester T, Hilton G, McKenzie CA, Salako T, Iwai N, Kita Y, Ogiwara T, Ohkubo T, Okamura T, Ueshima H, Umemura S, Eyheramendy S, Meitinger T, Wichmann HE, Cho YS, Kim HL, Lee JY, Scott J, Sehmi JS, Zhang W, Hedblad B, Nilsson P, Smith GD, Wong A, Narisu N, Stančáková A, Raffel LJ, Yao J, Kathiresan S, O'Donnell CJ, Schwartz SM, Ikram MA, Longstreth WT Jr, Mosley TH, Seshadri S, Shrine NR, Wain LV, Morken MA, Swift AJ, Laitinen J, Prokopenko I, Zitting P, Cooper JA, Humphries SE, Danesh J, Rasheed A, Goel A, Hamsten A, Watkins H, Bakker SJ, van Gilst WH, Janipalli CS, Mani KR, Yajnik CS, Hofman A, Mattace-Raso FU, Oostra BA, Demirkan A, Isaacs A, Rivadeneira F, Lakatta EG, Orru M, Scuteri A, Ala-Korpela M, Kangas AJ, Lyytikäinen LP, Soininen P, Tukiainen T, Würtz P, Ong RT, Dörr M, Kroemer HK, Völker U, Völzke H, Galan P, Hercberg S, Lathrop M, Zelenika D, Deloukas P, Mangino M, Spector TD, Zhai G, Meschia JF, Nalls MA, Sharma P, Terzic J, Kumar MV, Denniff M, Zukowska-Szczepowska E, Wagenknecht LE, Fowkes FG, Charchar FJ, Schwarz PE, Hayward C, Guo X, Rotimi C, Bots ML, Brand E, Samani NJ, Polasek O, Talmud PJ, Nyberg F, Kuh D, Laan M, Hveem K, Palmer LJ, van der Schouw YT, Casas JP, Mohlke KL, Vineis P, Raitakari O, Ganesh SK, Wong TY, Tai ES, Cooper RS, Laakso M, Rao DC, Harris TB, Morris RW, Dominiczak AF, Kivimäki M, Marmot MG, Miki T, Saleheen D, Chandak GR, Coresh J, Navis G, Salomaa V, Han BG, Zhu X, Kooner JS, Melander O, Ridker PM, Bandinelli S, Gyllenstein UB, Wright AF, Wilson JF, Ferrucci L, Farrall M, Tuomilehto J, Pramstaller PP, Elosua R, Soranzo N, Sijbrands EJ, Altshuler D, Loos RJ, Shuldiner AR, Gieger C, Meneton P, Uitterlinden AG, Wareham NJ, Gudnason V, Rotter JI, Rettig R, Uda M, Strachan DP, Witteman JC, Hartikainen AL, Beckmann JS, Boerwinkle E, Vasani RS, Boehnke M, Larson MG, Jarvelin MR, Psaty BM, Abecasis GR, Chakravarti A, Elliott P, van Duijn CM, **Newton-Cheh C**, Levy D, Caulfield MJ, Johnson T. Genetic variants in novel pathways influence blood pressure and cardiovascular disease risk. *Nature*. 2011 Sep 11; 478(7367): 103-9. doi: 10.1038/nature10405. PMID: PMC3340926.

81. Wain LV, Verwoert GC, O'Reilly PF, Shi G, Johnson T, Johnson AD, Bochud M, Rice KM, Henneman P, Smith AV, Ehret GB, Amin N, Larson MG, Mooser V, Hadley D, Dörr M, Bis JC, Aspelund T, Esko T, Janssens AC, Zhao JH, Heath S, Laan M, Fu J, Pistis G, Luan J, Arora P, Lucas G, Pirastu N, Pichler I, Jackson AU, Webster RJ, Zhang F, Peden JF, Schmidt H, Tanaka T, Campbell H, Igl W, Milaneschi Y, Hottenga JJ, Vitart V, Chasman DI, Trompet S, Bragg-Gresham JL, Alizadeh BZ, Chambers JC, Guo X, Lehtimäki T, Kühnel B, Lopez LM, Polašek O, Boban M, Nelson CP, Morrison AC, Pihur V, Ganesh SK, Hofman A, Kundu S, Mattace-Raso FU, Rivadeneira F, Sijbrands EJ, Uitterlinden AG, Hwang SJ, Vasani RS, Wang TJ, Bergmann S, Vollenweider P, Waeber G,

Laitinen J, Pouta A, Zitting P, McArdle WL, Kroemer HK, Völker U, Völzke H, Glazer NL, Taylor KD, Harris TB, Alavere H, Haller T, Keis A, Tammesoo ML, Aulchenko Y, Barroso I, Khaw KT, Galan P, Hercberg S, Lathrop M, Eyheramendy S, Org E, Söber S, Lu X, Nolte IM, Penninx BW, Corre T, Masciullo C, Sala C, Groop L, Voight BF, Melander O, O'Donnell CJ, Salomaa V, d'Adamo AP, Fabretto A, Faletra F, Ulivi S, Del Greco F, Facheris M, Collins FS, Bergman RN, Beilby JP, Hung J, Musk AW, Mangino M, Shin SY, Soranzo N, Watkins H, Goel A, Hamsten A, Gider P, Loitfelder M, Zeginigg M, Hernandez D, Najjar SS, Navarro P, Wild SH, Corsi AM, Singleton A, de Geus EJ, Willemsen G, Parker AN, Rose LM, Buckley B, Stott D, Orru M, Uda M; LifeLines Cohort Study, van der Klauw MM, Zhang W, Li X, Scott J, Chen YD, Burke GL, Kähönen M, Viikari J, Döring A, Meitinger T, Davies G, Starr JM, Emilsson V, Plump A, Lindeman JH, Hoen PA, König IR; EchoGen consortium, Felix JF, Clarke R, Hopewell JC, Ongen H, Breteler M, Dobbie S, Destefano AL, Fornage M; AortaGen Consortium, Mitchell GF; CHARGE Consortium Heart Failure Working Group, Smith NL; KidneyGen consortium, Holm H, Stefansson K, Thorleifsson G, Thorsteinsdottir U; CKDGen consortium; Cardiogenics consortium; CardioGram, Samani NJ, Preuss M, Rudan I, Hayward C, Deary IJ, Wichmann HE, Raitakari OT, Palmas W, Kooner JS, Stolk RP, Jukema JW, Wright AF, Boomsma DI, Bandinelli S, Gyllenstein UB, Wilson JF, Ferrucci L, Schmidt R, Farrall M, Spector TD, Palmer LJ, Tuomilehto J, Pfeufer A, Gasparini P, Siscovick D, Altshuler D, Loos RJ, Toniolo D, Snieder H, Gieger C, Meneton P, Wareham NJ, Oostra BA, Metspalu A, Launer L, Rettig R, Strachan DP, Beckmann JS, Witteman JC, Erdmann J, van Dijk KW, Boerwinkle E, Boehnke M, Ridker PM, Jarvelin MR, Chakravarti A, Abecasis GR, Gudnason V, **Newton-Cheh C**, Levy D, Munroe PB, Psaty BM, Caulfield MJ, Rao DC, Tobin MD, Elliott P, van Duijn CM. Genome-wide association study identifies six new loci influencing pulse pressure and mean arterial pressure. *Nat Genet.* 2011 Sep 11; 43(10): 1005-11. doi: 10.1038/ng.922. PMID: PMC3445021.

82. Khan AM, Cheng S, Magnusson M, Larson MG, **Newton-Cheh C**, McCabe EL, Coviello AD, Florez JC, Fox CS, Levy D, Robins SJ, Arora P, Bhasin S, Lam CS, Vasani RS, Melander O, Wang TJ. Cardiac natriuretic peptides, obesity, and insulin resistance: evidence from two community-based studies. *J Clin Endocrinol Metab.* 2011 Oct; 96(10):3242-9. PMID: PMC3200240.
83. Schnabel RB, Kerr KF, Lubitz SA, Alkylbekova EL, Marcus GM, Sinner MF, Magnani JW, Wolf PA, Deo R, Lloyd-Jones DM, Lunetta KL, Mehra R, Levy D, Fox ER, Arking DE, Mosley TH, Müller-Nurasyid M, Young TR, Wichmann HE, Seshadri S, Farlow DN, Rotter JI, Soliman EZ, Glazer NL, Wilson JG, Breteler MM, Sotoodehnia N, **Newton-Cheh C**, Kääh S, Ellinor PT, Alonso A, Benjamin EJ, Heckbert SR; for the Candidate Gene Association Resource (CARE) Atrial Fibrillation/Electrocardiography Working Group. Large-Scale Candidate Gene Analysis in Whites and African Americans Identifies IL6R Polymorphism in Relation to Atrial Fibrillation: The National Heart, Lung, and Blood Institute's Candidate Gene Association Resource (CARE) Project. *Circ Cardiovasc Genet.* 2011 Oct 1; 4(5):557-564. Epub 2011 Aug 16. PMID: PMC3224824.
84. Cheng S, Fox CS, Larson MG, Massaro JM, McCabe EL, Khan AM, Levy D, Hoffmann U, O'Donnell CJ, Miller KK, **Newton-Cheh C**, Coviello AD, Bhasin S, Vasani RS, Wang TJ. Relation of visceral adiposity to circulating natriuretic peptides in ambulatory individuals. *Am J Cardiol.* 2011 Oct 1; 108(7):979-84. Epub 2011 Aug 1. PMID: PMC3175269.
85. Oikonen M, Tikkanen E, Juhola J, Tuovinen T, Seppälä I, Juonala M, Taittonen L, Mikkilä V, Kähönen M, Ripatti S, Viikari J, Lehtimäki T, Havulinna AS, Kee F, **Newton-Cheh C**, Peltonen L, Schork NJ, Murray SS, Berenson GS, Chen W, Srinivasan SR, Salomaa V, Raitakari OT. Genetic

variants and blood pressure in a population-based cohort: the cardiovascular risk in young Finns study. *Hypertension*. 2011 Dec; 58(6):1079-85. PMID: PMC3247907.

86. Johnson T, Gaunt TR, Newhouse SJ, Padmanabhan S, Tomaszewski M, Kumari M, Morris RW, Tzoulaki I, O'Brien ET, Poulter NR, Sever P, Shields DC, Thom S, Wannamethee SG, Whincup PH, Brown MJ, Connell JM, Dobson RJ, Howard PJ, Mein CA, Onipinla A, Shaw-Hawkins S, Zhang Y, Smith GD, Day IN, Lawlor DA, Goodall AH; The Cardiogenics Consortium, Fowkes FG, Abecasis GR, Elliott P, Gateva V; The Global BPgen Consortium, Braund PS, Burton PR, Nelson CP, Tobin MD, van der Harst P, Glorioso N, Neuvirth H, Salvi E, Staessen JA, Stucchi A, Devos N, Jeunemaitre X, Plouin PF, Tichet J, Juhanson P, Org E, Putku M, Söber S, Veldre G, Viigimaa M, Levinsson A, Rosengren A, Thelle DS, Hastie CE, Hedner T, Lee WK, Melander O, Wahlstrand B, Hardy R, Wong A, Cooper JA, Palmén J, Chen L, Stewart AF, Wells GA, Westra HJ, Wolfs MG, Clarke R, Franzosi MG, Goel A, Hamsten A, Lathrop M, Peden JF, Seedorf U, Watkins H, Ouwehand WH, Sambrook J, Stephens J, Casas JP, Drenos F, Holmes MV, Kivimaki M, Shah S, Shah T, Talmud PJ, Whittaker J, Wallace C, Delles C, Laan M, Kuh D, Humphries SE, Nyberg F, Cusi D, Roberts R, **Newton-Cheh C**, Franke L, Stanton AV, Dominiczak AF, Farrall M, Hingorani AD, Samani NJ, Caulfield MJ, Munroe PB. Blood Pressure Loci Identified with a Gene-Centric Array. *Am J Hum Genet*. 2011 Dec 9; 89(6):688-700. Epub 2011 Nov 17. PMID: PMC3234370.
87. Gaál EI, Salo P, Kristiansson K, Rehnström K, Kettunen J, Sarin AP, Niemelä M, Jula A, Raitakari OT, Lehtimäki T, Eriksson JG, Widen E, Günel M, Kurki M, von Und Zu Fraunberg M, Jääskeläinen JE, Hernesniemi J, Järvelin MR, Pouta A; The International Consortium for Blood Pressure Genome-Wide Association Studies (ICBP-GWAS), **Newton-Cheh C**, Salomaa V, Palotie A, Perola M. Intracranial Aneurysm Risk Locus 5q23.2 Is Associated with Elevated Systolic Blood Pressure. *PLoS Genet*. 2012 Mar; 8(3):e1002563. Epub 2012 Mar 15. PMID: PMC3305343.
88. Lahtinen AM, Noseworthy PA, Havulinna AS, Jula A, Karhunen PJ, Kettunen J, Perola M, Kontula K, **Newton-Cheh C**, Salomaa V. Common Genetic Variants Associated with Sudden Cardiac Death: The FinSCDgen Study. *PLoS One*. 2012; 7(7):e41675. Epub 2012 Jul 23. PMID: PMC3402479.
89. Voight BF, Kang HM, Ding J, Palmer CD, Sidore C, Chines PS, Burt NP, Fuchsberger C, Li Y, Erdmann J, Frayling TM, Heid IM, Jackson AU, Johnson T, Kilpeläinen TO, Lindgren CM, Morris AP, Prokopenko I, Randall JC, Saxena R, Soranzo N, Speliotes EK, Teslovich TM, Wheeler E, Maguire J, Parkin M, Potter S, Rayner NW, Robertson N, Stirrups K, Winckler W, Sanna S, Mulas A, Nagaraja R, Cucca F, Barroso I, Deloukas P, Loos RJ, Kathiresan S, Munroe PB, **Newton-Cheh C**, Pfeufer A, Samani NJ, Schunkert H, Hirschhorn JN, Altshuler D, McCarthy MI, Abecasis GR, Boehnke M. The metabochip, a custom genotyping array for genetic studies of metabolic, cardiovascular, and anthropometric traits. *PLoS Genet*. 2012;8(8): e1002793. doi: 10.1371/journal.pgen.1002793. Epub 2012 Aug 2. Erratum in: *PLoS Genet*. 2013 Apr; 9(4). doi: 10.1371/annotation/0b4e9c8b-35c5-4dbd-b95b-0640250fbc87. PMID: PMC3410907.
90. Magnusson M, Jujic A, Hedblad B, Engström G, Persson M, Struck J, Morgenthaler NG, Nilsson P, **Newton-Cheh C**, Wang TJ, Melander O. Low plasma level of atrial natriuretic peptide predicts development of diabetes: the prospective Malmo Diet and Cancer study. *J Clin Endocrinol Metab*. 2012 Feb; 97(2):638-45. Epub 2011 Nov 23. PMID: PMC3275360.
91. Lin E, McCabe E, **Newton-Cheh C**, Bloch K, Buys E, Wang T, Miller KK. Effects of transdermal testosterone on natriuretic peptide levels in women: a randomized placebo-controlled pilot study.

Fertil Steril. 2012 Feb; 97(2):489-93. Epub 2011 Dec 2. PMID: PMC3269509.

92. Musunuru K, Roden DM, Boineau R, Bristow MR, McCaffrey TA, **Newton-Cheh C**, Paltoo DN, Rosenberg Y, Wohlgemuth JG, Zineh I, Hasan AA. Cardiovascular pharmacogenomics: current status and future directions-report of a national heart, lung, and blood institute working group. *J Am Heart Assoc.* 2012 Apr; 1(2): e000554. doi: 10.1161/JAHA.111.000554. Epub 2012 Apr 24. PMID: PMC3487365.
93. Tanguturi VK, Noseworthy PA, **Newton-Cheh C**, Baggish AL. The electrocardiographic early repolarization pattern in athletes: normal variant or sudden death risk factor? *Sports Med.* 2012 May 1; 42(5):359-66. doi: 10.2165/11630790-000000000-00000.
94. Smith JG, **Newton-Cheh C**, Almgren P, Melander O, Platonov PG. Genetic polymorphisms for estimating risk of atrial fibrillation in the general population: a prospective study. *Arch Intern Med.* 2012 May 14; 172(9):742-4. PMID: PMC3763742.
95. Marjamaa A, Oikarinen L, Porthan K, Ripatti S, Peloso G, Noseworthy PA, Viitasalo M, Nieminen MS, Toivonen L, Kontula K, Peltonen L, Havulinna AS, Jula A, O'Donnell CJ, **Newton-Cheh C**, Perola M, Salomaa V. A common variant near the KCNJ2 gene is associated with T-peak to T-end interval. *Heart Rhythm.* 2012 Jul; 9(7): 1099-103. doi: 10.1016/j.hrthm.2012.02.019. Epub 2012 Feb 15. PMID: PMC3690340.
96. Voight BF, Peloso GM, Orho-Melander M, Frikke-Schmidt R, Barbalic M, Jensen MK, Hindy G, Hólm H, Ding EL, Johnson T, Schunkert H, Samani NJ, Clarke R, Hopewell JC, Thompson JF, Li M, Thorleifsson G, **Newton-Cheh C**, Musunuru K, Pirruccello JP, Saleheen D, Chen L, Stewart A, Schillert A, Thorsteinsdottir U, Thorgeirsson G, Anand S, Engert JC, Morgan T, Spertus J, Stoll M, Berger K, Martinelli N, Girelli D, McKeown PP, Patterson CC, Epstein SE, Devaney J, Burnett MS, Mooser V, Ripatti S, Surakka I, Nieminen MS, Sinisalo J, Lokki ML, Perola M, Havulinna A, de Faire U, Gigante B, Ingelsson E, Zeller T, Wild P, de Bakker PI, Klungel OH, Maitland-van der Zee AH, Peters BJ, de Boer A, Grobbee DE, Kamphuisen PW, Deneer VH, Elbers CC, Onland-Moret NC, Hofker MH, Wijmenga C, Verschuren WM, Boer JM, van der Schouw YT, Rasheed A, Frossard P, Demissie S, Willer C, Do R, Ordovas JM, Abecasis GR, Boehnke M, Mohlke KL, Daly MJ, Guiducci C, Burt NP, Surti A, Gonzalez E, Purcell S, Gabriel S, Marrugat J, Peden J, Erdmann J, Diemert P, Willenborg C, König IR, Fischer M, Hengstenberg C, Ziegler A, Buyschaert I, Lambrechts D, Van de Werf F, Fox KA, El Mokhtari NE, Rubin D, Schrezenmeier J, Schreiber S, Schäfer A, Danesh J, Blankenberg S, Roberts R, McPherson R, Watkins H, Hall AS, Overvad K, Rimm E, Boerwinkle E, Tybjaerg-Hansen A, Cupples LA, Reilly MP, Melander O, Mannucci PM, Ardissino D, Siscovick D, Elosua R, Stefansson K, O'Donnell CJ, Salomaa V, Rader DJ, Peltonen L, Schwartz SM, Altshuler D, Kathiresan S. Plasma HDL cholesterol and risk of myocardial infarction: a mendelian randomisation study. *Lancet.* 2012 Aug 11; 380(9841):572-80. Epub 2012 May 17. Erratum in: *Lancet.* 2012 Aug 11; 380(9841):564. PMID: PMC3419820.
97. Fedorowski A, Franceschini N, Brody J, Liu C, Verwoert GC, Boerwinkle E, Couper D, Rice KM, Rotter JI, Mattace-Raso F, Uitterlinden A, Hofman A, Almgren P, Sjögren M, Hedblad B, Larson MG, **Newton-Cheh C**, Wang TJ, Rose KM, Psaty BM, Levy D, Witteman J, Melander O. Orthostatic hypotension and novel blood pressure-associated gene variants: Genetics of Postural Hemodynamics (GPH) Consortium. *Eur Heart J.* 2012 Sep; 33(18):2331-41. Epub 2012 Apr 14. PMID: PMC3442958.

98. Bick AG, Flannick J, Ito K, Cheng S, Vasani RS, Parfenov MG, Herman DS, DePalma SR, Gupta N, Gabriel SB, Funke BH, Rehm HL, Benjamin EJ, Aragam J, Taylor HA Jr, Fox ER, **Newton-Cheh C**, Kathiresan S, O'Donnell CJ, Wilson JG, Altshuler DM, Hirschhorn JN, Seidman JG, Seidman C. Burden of rare sarcomere gene variants in the Framingham and Jackson Heart Study cohorts. *Am J Hum Genet.* 2012 Sep 7; 91(3): 513-9. doi: 10.1016/j.ajhg.2012.07.017. PMID: PMC3511985.
99. Noseworthy PA, Peloso GM, Hwang SJ, Larson MG, Levy D, O'Donnell CJ, **Newton-Cheh C**. QT Interval and Long-Term Mortality Risk in the Framingham Heart Study. *Ann Noninvasive Electrocardiol.* 2012 Oct; 17(4): 340-348. doi: 10.1111/j.1542-474X.2012.00535.x. Epub 2012 Aug 13. PMID: PMC3481183.
100. Sinner MF, Porthan K, Noseworthy PA, Havulinna AS, Tikkanen JT, Müller-Nurasyid M, Peloso G, Ulivi S, Beckmann BM, Brockhaus AC, Cooper RR, Gasparini P, Hengstenberg C, Hwang SJ, Iorio A, Junttila MJ, Klopp N, Kähönen M, Laaksonen MA, Lehtimäki T, Lichtner P, Lyytikäinen LP, Martens E, Meisinger C, Meitinger T, Merchant FM, Nieminen MS, Peters A, Pietilä A, Perz S, Oikarinen L, Raitakari O, Reinhard W, Silander K, Thorand B, Wichmann HE, Sinagra G, Viikari J, O'Donnell CJ, Ellinor PT, Huikuri HV, Kääb S, **Newton-Cheh C**, Salomaa V. A meta-analysis of genome-wide association studies of the electrocardiographic early repolarization pattern. *Heart Rhythm.* 2012 Oct; 9(10): 1627-34. doi: 10.1016/j.hrthm.2012.06.008. Epub 2012 Jun 6. PubMed PMID: PMC3459269.
101. Magnani JW, **Newton-Cheh C**, O'Donnell CJ, Levy D. Development and application of a longitudinal electrocardiogram repository: the Framingham Heart Study. *J Electrocardiol.* 2012 Nov; 45(6): 673-6. doi: 10.1016/j.jelectrocard.2012.06.016. Epub 2012 Jul 24. PMID: PMC3483375.
102. Jena AB, Romley JA, **Newton-Cheh C**, Noseworthy P. Therapeutic hypothermia for cardiac arrest: Real-world utilization trends and hospital mortality. *J Hosp Med.* 2012 Nov; 7(9): 684-9. doi: 10.1002/jhm.1974. Epub 2012 Sep 28. PMID: PMC3515738.
103. Smith JG, Almgren P, Engström G, Hedblad B, Platonov PG, **Newton-Cheh C**, Melander O. Genetic polymorphisms for estimating risk of atrial fibrillation: a literature-based meta-analysis. *J Intern Med.* 2012 Dec; 272(6):573-82. doi: 10.1111/j.1365-2796.2012.02563.x. Epub 2012 Jul 27. PMID: PMC3763745.
104. 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 WC, 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, Redline S, Reiner AP, Rotter JI, Schnabel RB, Schork NJ, Singleton AB, Smith JG, Soliman EZ, Srinivasan SR, Zhang ZM, 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 10 African American cohorts. *Circ Cardiovasc Genet.* 2012 Dec; 5(6): 639-46. doi: 10.1161/CIRCGENETICS.112.963991. Epub 2012 Nov 8. PMID: PMC3560365.
105. 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, Li G, Ellinor PT, Magnani JW, Chen W, Bis JC, Curb JD, Hsueh WC, Rotter JI, Liu Y, Newman AB, Limacher MC, North KE,

- Reiner AP, Quibrera PM, Schork NJ, Singleton AB, Psaty BM, Soliman EZ, Solomon AJ, Srinivasan SR, Alonso A, Wallace R, Redline S, Zhang ZM, 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**; CARE and COGENT consortia. Impact of ancestry and common genetic variants on QT interval in African Americans. *Circ Cardiovasc Genet*. 2012 Dec; 5(6):647-55. doi: 10.1161/CIRCGENETICS.112.962787. PMID: PMC3568265.
106. Huertas-Vazquez A, Nelson CP, Guo X, Reinier K, Uy-Evanado A, Teodorescu C, Ayala J, Jerger K, Chugh H, Wtccc, Braund PS, Deloukas P, Hall AS, Balmforth AJ, Jones M, Taylor KD, Pulit SL, **Newton-Cheh C**, Gunson K, Jui J, Rotter JI, Albert CM, Samani NJ, Chugh SS. Novel Loci associated with increased risk of sudden cardiac death in the context of coronary artery disease. *PLoS One*. 2013 Apr 4; 8(4): e59905. doi: 10.1371/journal.pone.0059905. Print 2013. PMID: PMC3617189.
107. Johnson AD, Hwang SJ, Voorman A, Morrison A, Peloso GM, Hsu YH, Thanassoulis G, **Newton-Cheh C**, Rogers IS, Hoffmann U, Freedman JE, Fox CS, Psaty BM, Boerwinkle E, Cupples LA, O'Donnell CJ. Resequencing and clinical associations of the 9p21.3 region: a comprehensive investigation in the Framingham heart study. *Circulation*. 2013 Feb 19; 127(7): 799-810. doi: 10.1161/CIRCULATIONAHA.112.111559. Epub 2013 Jan 11. PMID: PMC3686634.
108. Deo R, Nalls MA, Avery CL, Smith JG, Evans DS, Keller MF, Butler AM, Buxbaum SG, Li G, Miguel Quibrera P, Smith EN, Tanaka T, Akylbekova EL, Alonso A, Arking DE, Benjamin EJ, Berenson GS, Bis JC, Chen LY, Chen W, Cummings SR, Ellinor PT, Evans MK, Ferrucci L, Fox ER, Heckbert SR, Heiss G, Hsueh WC, Kerr KF, Limacher MC, Liu Y, Lubitz SA, Magnani JW, Mehra R, Marcus GM, Murray SS, Newman AB, Njajou O, North KE, Paltoo DN, Psaty BM, Redline SS, Reiner AP, Robinson JG, Rotter JI, Samdarshi TE, Schnabel RB, Schork NJ, Singleton AB, Siscovick D, Soliman EZ, Sotoodehnia N, Srinivasan SR, Taylor HA, Trevisan M, Zhang Z, Zonderman AB, **Newton-Cheh C**, Whitsel EA. Common genetic variation near the connexin-43 gene is associated with resting heart rate in African Americans: A genome-wide association study of 13,372 participants. *Heart Rhythm*. 2013 Mar; 10(3): 401-8. doi: 10.1016/j.hrthm.2012.11.014. Epub 2012 Nov 24. PMID: PMC3718037.
109. Ganesh SK, Tragante V, Guo W, Guo Y, Lanktree MB, Smith EN, Johnson T, Castillo BA, Barnard J, Baumert J, Chang YP, Elbers CC, Farrall M, Fischer ME, Franceschini N, Gaunt TR, Gho JM, Gieger C, Gong Y, Isaacs A, Kleber ME, Mateo Leach I, McDonough CW, Meijs MF, Mellander O, Molony CM, Nolte IM, Padmanabhan S, Price TS, Rajagopalan R, Shaffer J, Shah S, Shen H, Soranzo N, van der Most PJ, Van Iperen EP, Van Setten JA, Vonk JM, Zhang L, Beitelshes AL, Berenson GS, Bhatt DL, Boer JM, Boerwinkle E, Burkley B, Burt A, Chakravarti A, Chen W, Cooper-Dehoff RM, Curtis SP, Dreisbach A, Duggan D, Ehret GB, Fabsitz RR, Fornage M, Fox E, Furlong CE, Gansevoort RT, Hofker MH, Hovingh GK, Kirkland SA, Kottke-Marchant K, Kutlar A, Lacroix AZ, Langa TY, Li YR, Lin H, Liu K, Maiwald S, Malik R; CARDIOGRAM, METASTROKE, Murugesan G, **Newton-Cheh C**, O'Connell JR, Onland-Moret NC, Ouwehand WH, Palmas W, Penninx BW, Pepine CJ, Pettinger M, Polak JF, Ramachandran VS, Ranchalis J, Redline S, Ridker PM, Rose LM, Scharnag H, Schork NJ, Shimbo D, Shuldiner AR, Srinivasan SR, Stolk RP, Taylor HA, Thorand B, Trip MD, van Duijn CM, Verschuren WM, Wijmenga C, Winkelmann BR, Wyatt S, Young JH, Boehm BO, Caulfield MJ, Chasman DI, Davidson KW, Doevendans PA, Fitzgerald GA, Gums JG, Hakonarson H, Hillege HL, Illig T, Jarvik GP, Johnson JA, Kastelein JJ, Koenig W; LifeLines Cohort Study, März W, Mitchell BD, Murray SS, Oldehinkel AJ, Rader DJ,

Reilly MP, Reiner AP, Schadt EE, Silverstein RL, Snieder H, Stanton AV, Uitterlinden AG, van der Harst P, van der Schouw YT, Samani NJ, Johnson AD, Munroe PB, de Bakker PI, Zhu X, Levy D, Keating BJ, Asselbergs FW. Loci influencing blood pressure identified using a cardiovascular gene-centric array. *Hum Mol Genet.* 2013 Apr 15; 22(8): 1663-78. doi: 10.1093/hmg/dds555. Epub 2013 Jan 8. PMID: PMC3657476.

110. Lieb W, Jansen H, Loley C, Pencina MJ, Nelson CP, **Newton-Cheh C**, Kathiresan S, Reilly MP, Assimes TL, Boerwinkle E, Hall AS, Hengstenberg C, Laaksonen R, McPherson R, Thorsteinsdottir U, Ziegler A, Peters A, Thompson JR, König IR, Erdmann J, Samani NJ, Vasani RS, Schunkert H; CARDIoGRAM, Assimes TL, Deloukas P, Erdmann J, Holm H, Kathiresan S, König IR, McPherson R, Reilly MP, Roberts R, Samani NJ, Schunkert H, Stewart AF. Genetic predisposition to higher blood pressure increases coronary artery disease risk. *Hypertension.* 2013 May; 61(5): 995-1001. doi: 10.1161/HYPERTENSIONAHA.111.00275. Epub 2013 Mar 11. PMID: PMC3855241.
111. Havulinna AS, Kettunen J, Ukkola O, Osmond C, Eriksson JG, Kesäniemi YA, Jula A, Peltonen L, Kontula K, Salomaa V, **Newton-Cheh C**. A blood pressure genetic risk score is a significant predictor of incident cardiovascular events in 32,669 individuals. *Hypertension.* 2013 May; 61(5): 987-94. doi: 10.1161/HYPERTENSIONAHA.111.00649. Epub 2013 Mar 18. PMID: PMC3648219.
112. den Hoed M, Eijgelsheim M, Esko T, Brundel BJ, Peal DS, Evans DM, Nolte IM, Segrè AV, Holm H, Handsaker RE, Westra HJ, Johnson T, Isaacs A, Yang J, Lundby A, Zhao JH, Kim YJ, Go MJ, Almgren P, Bochud M, Boucher G, Cornelis MC, Gudbjartsson D, Hadley D, van der Harst P, Hayward C, den Heijer M, Igl W, Jackson AU, Kutalik Z, Luan J, Kemp JP, Kristiansson K, Ladenvall C, Lorentzon M, Montasser ME, Njajou OT, O'Reilly PF, Padmanabhan S, St Pourcain B, Rankinen T, Salo P, Tanaka T, Timpson NJ, Vitart V, Waite L, Wheeler W, Zhang W, Draisma HH, Feitosa MF, Kerr KF, Lind PA, Mihailov E, Onland-Moret NC, Song C, Weedon MN, Xie W, Yengo L, Absher D, Albert CM, Alonso A, Arking DE, de Bakker PI, Balkau B, Barlassina C, Benaglio P, Bis JC, Bouatia-Naji N, Brage S, Chanock SJ, Chines PS, Chung M, Darbar D, Dina C, Dörr M, Elliott P, Felix SB, Fischer K, Fuchsberger C, de Geus EJ, Goyette P, Gudnason V, Harris TB, Hartikainen AL, Havulinna AS, Heckbert SR, Hicks AA, Hofman A, Holewijn S, Hoogstra-Berends F, Hottenga JJ, Jensen MK, Johansson A, Junttila J, Kääb S, Kanon B, Ketkar S, Khaw KT, Knowles JW, Kooner AS, Kors JA, Kumari M, Milani L, Laiho P, Lakatta EG, Langenberg C, Leusink M, Liu Y, Luben RN, Lunetta KL, Lynch SN, Markus MR, Marques-Vidal P, Mateo Leach I, McArdle WL, McCarroll SA, Medland SE, Miller KA, Montgomery GW, Morrison AC, Müller-Nurasyid M, Navarro P, Nelis M, O'Connell JR, O'Donnell CJ, Ong KK, Newman AB, Peters A, Polasek O, Pouta A, Pramstaller PP, Psaty BM, Rao DC, Ring SM, Rossin EJ, Rudan D, Sanna S, Scott RA, Sehmi JS, Sharp S, Shin JT, Singleton AB, Smith AV, Soranzo N, Spector TD, Stewart C, Stringham HM, Tarasov KV, Uitterlinden AG, Vandenput L, Hwang SJ, Whitfield JB, Wijmenga C, Wild SH, Willemsen G, Wilson JF, Witteman JC, Wong A, Wong Q, Jamshidi Y, Zitting P, Boer JM, Boomsma DI, Borecki IB, van Duijn CM, Ekelund U, Forouhi NG, Froguel P, Hingorani A, Ingelsson E, Kivimäki M, Kronmal RA, Kuh D, Lind L, Martin NG, Oostra BA, Pedersen NL, Quertermous T, Rotter JI, van der Schouw YT, Verschuren WM, Walker M, Albanes D, Arnar DO, Assimes TL, Bandinelli S, Boehnke M, de Boer RA, Bouchard C, Caulfield WL, Chambers JC, Curhan G, Cusi D, Eriksson J, Ferrucci L, van Gilst WH, Glorioso N, de Graaf J, Groop L, Gyllenstein U, Hsueh WC, Hu FB, Huikuri HV, Hunter DJ, Iribarren C, Isomaa B, Jarvelin MR, Jula A, Kähönen M, Kiemeny LA, van der Klauw MM, Kooner JS, Kraft P, Iacoviello L, Lehtimäki T, Lokki ML, Mitchell BD, Navis G, Nieminen MS, Ohlsson C, Poulter NR, Qi L, Raitakari OT, Rimm EB, Rioux JD, Rizzi F, Rudan I, Salomaa V, Sever PS, Shields DC, Shuldiner AR, Sinisalo J, Stanton AV, Stolk RP, Strachan DP,

- Tardif JC, Thorsteinsdottir U, Tuomilehto J, van Veldhuisen DJ, Virtamo J, Viikari J, Vollenweider P, Waeber G, Widen E, Cho YS, Olsen JV, Visscher PM, Willer C, Franke L; Global BPgen Consortium; CARDIoGRAM Consortium, Erdmann J, Thompson JR; PR GWAS Consortium, Pfeufer A; QRS GWAS Consortium, Sotoodehnia N; QT-IGC Consortium, **Newton-Cheh C**; CHARGE-AF Consortium, Ellinor PT, Stricker BH, Metspalu A, Perola M, Beckmann JS, Smith GD, Stefansson K, Wareham NJ, Munroe PB, Sibon OC, Milan DJ, Snieder H, Samani NJ, Loos RJ. Identification of heart rate-associated loci and their effects on cardiac conduction and rhythm disorders. *Nat Genet.* 2013 Jun; 45(6): 621-31. doi: 10.1038/ng.2610. Epub 2013 Apr 14. PMID: PMC3696959.
113. Lahtinen AM, Havulinna AS, Noseworthy PA, Jula A, Karhunen PJ, Perola M, **Newton-Cheh C**, Salomaa V, Kontula K. Prevalence of arrhythmia-associated gene mutations and risk of sudden cardiac death in the Finnish population. *Ann Med.* 2013 Jun; 45(4): 328-35. doi: 10.3109/07853890.2013.783995. Epub 2013 May 8. PMID: PMC3778376.
114. Huertas-Vazquez A, Teodorescu C, Reinier K, Uy-Evanado A, Chugh H, Jerger K, Ayala J, Gunson K, Jui J, **Newton-Cheh C**, Albert CM, Chugh SS. A common missense variant in the neuregulin 1 gene is associated with both schizophrenia and sudden cardiac death. *Heart Rhythm.* 2013 Jul; 10(7): 994-8. doi: 10.1016/j.hrthm.2013.03.020. Epub 2013 Mar 21. PMID: PMC3692570.
115. Arora P, Wu C, Khan AM, Bloch DB, Davis-Dusenbery BN, Ghorbani A, Spagnolli E, Martinez A, Ryan A, Tainsh LT, Kim S, Rong J, Huan T, Freedman JE, Levy D, Miller KK, Hata A, Del Monte F, Vandenwijngaert S, Swinnen M, Janssens S, Holmes TM, Buys ES, Bloch KD, **Newton-Cheh C**, Wang TJ. Atrial natriuretic peptide is negatively regulated by microRNA-425. *J Clin Invest.* 2013 Aug 1; 123(8): 3378-82. doi: 10.1172/JCI67383. Epub 2013 Jul 15. PMID: PMC3726159.
116. Porthan K, Viitasalo M, Toivonen L, Havulinna AS, Jula A, Tikkanen JT, Väänänen H, Nieminen MS, Huikuri HV, **Newton-Cheh C**, Salomaa V, Oikarinen L. Predictive value of electrocardiographic T-wave morphology parameters and T-wave peak to T-wave end interval for sudden cardiac death in the general population. *Circ Arrhythm Electrophysiol.* 2013 Aug; 6(4): 690-6. doi: 10.1161/CIRCEP.113.000356. Epub 2013 Jul 23.
117. Li A, Ahsen OO, Liu JJ, Du C, McKee ML, Yang Y, Wasco W, **Newton-Cheh CH**, O'Donnell CJ, Fujimoto JG, Zhou C, Tanzi RE. Silencing of the Drosophila ortholog of SOX5 in heart leads to cardiac dysfunction as detected by optical coherence tomography. *Hum Mol Genet.* 2013 Sep 15; 22(18): 3798-806. doi: 10.1093/hmg/ddt230. Epub 2013 May 21. PMID: PMC3749865.
118. Ho JE, Chen WY, Chen MH, Larson MG, McCabe EL, Cheng S, Ghorbani A, Coglianese E, Emilsson V, Johnson AD, Walter S, Franceschini N, O'Donnell CJ, Dehghan A, Lu C, Levy D, **Newton-Cheh C**, Lin H, Felix JF, Schreiter ER, Vasani RS, Januzzi JL, Lee RT, Wang TJ. Common genetic variation at the IL1RL1 locus regulates IL-33/ST2 signaling. *J Clin Invest.* 2013 Oct 1; 123(10): 4208-18. doi: 10.1172/JCI67119. Epub 2013 Sep 3. PMID: PMC3784527.
119. Mak GS, Sawaya H, Khan AM, Arora P, Martinez A, Ryan A, Ernande L, **Newton-Cheh C**, Wang TJ, Scherrer-Crosbie M. Effects of subacute dietary salt intake and acute volume expansion on diastolic function in young normotensive individuals. *Eur Heart J Cardiovasc Imaging.* 2013 Nov; 14(11): 1092-8. doi: 10.1093/ehjci/jet036. Epub 2013 Mar 20. PMID: PMC3806580.
120. Flannick J, Beer NL, Bick AG, Agarwala V, Molnes J, Gupta N, Burtt NP, Florez JC, Meigs JB,

- Taylor H, Lyssenko V, Irgens H, Fox E, Burslem F, Johansson S, Brosnan MJ, Trimmer JK, **Newton-Cheh C**, Tuomi T, Molven A, Wilson JG, O'Donnell CJ, Kathiresan S, Hirschhorn JN, Njølstad PR, Rolph T, Seidman JG, Gabriel S, Cox DR, Seidman CE, Groop L, Altshuler D. Assessing the phenotypic effects in the general population of rare variants in genes for a dominant Mendelian form of diabetes. *Nat Genet.* 2013 Nov; 45(11): 1380-5. doi: 10.1038/ng.2794. Epub 2013 Oct 6. PMID: PMC4051627.
121. Akylbekova EL, Payne JP, **Newton-Cheh C**, May WL, Fox ER, Wilson JG, Sarpong DF, Taylor HA, Maher JF. Gene-environment interaction between SCN5A-1103Y and hypokalemia influences QT interval prolongation in African Americans: the Jackson Heart Study. *Am Heart J.* 2014 Jan; 167(1): 116-122.e1. doi: 10.1016/j.ahj.2013.10.009. Epub 2013 Oct 22. PMID: PMC3884587.
122. Avery CL, Sitlani CM, Arking DE, Arnett DK, Bis JC, Boerwinkle E, Buckley BM, Ida Chen YD, de Craen AJ, Eijgelsheim M, Enquobahrie D, Evans DS, Ford I, Garcia ME, Gudnason V, Harris TB, Heckbert SR, Hochner H, Hofman A, Hsueh WC, Isaacs A, Jukema JW, Knekt P, Kors JA, Krijthe BP, Kristiansson K, Laaksonen M, Liu Y, Li X, Macfarlane PW, **Newton-Cheh C**, Nieminen MS, Oostra BA, Peloso GM, Porthan K, Rice K, Rivadeneira FF, Rotter JI, Salomaa V, Sattar N, Siscovick DS, Slagboom PE, Smith AV, Sotoodehnia N, Stott DJ, Stricker BH, Stürmer T, Trompet S, Uitterlinden AG, van Duijn C, Westendorp RG, Witteman JC, Whitsel EA, Psaty BM. Drug-gene interactions and the search for missing heritability: a cross-sectional pharmacogenomics study of the QT interval. *Pharmacogenomics J.* 2014 Feb; 14(1): 6-13. doi: 10.1038/tpj.2013.4. Epub 2013 Mar 5. PMID: PMC3766418.
123. Ito K, Bick AG, Flannick J, Friedman DJ, Genovese G, Parfenov MG, Depalma SR, Gupta N, Gabriel SB, Taylor HA Jr, Fox ER, **Newton-Cheh C**, Kathiresan S, Hirschhorn JN, Altshuler DM, Pollak MR, Wilson JG, Seidman JG, Seidman C. Increased burden of cardiovascular disease in carriers of APOL1 genetic variants. *Circ Res.* 2014 Feb 28; 114(5):845-50. doi: 10.1161/CIRCRESAHA.114.302347. Epub 2013 Dec 30. PMID: PMC3982584. Ito K, Bick AG, Flannick J, Friedman DJ, Genovese G, Parfenov MG, Depalma SR, Gupta N, Gabriel SB, Taylor HA Jr, Fox ER, **Newton-Cheh C**, Kathiresan S, Hirschhorn JN, Altshuler DM, Pollak MR, Wilson JG, Seidman JG, Seidman C. Increased Burden of Cardiovascular Disease in Carriers of APOL1 Genetic Variants. *Circ Res.* 2014 Feb 28; 114(5): 845-50. doi: 10.1161/CIRCRESAHA.114.302347. Epub 2013 Dec 30. PMID: PMC3982584.
124. Lemaitre RN, Johnson CO, Hesselton S, Sotoodehnia N, McKnight B, Sitlani CM, Rea TD, King IB, Kwok PY, Mak A, Li G, Brody J, Larson E, Mozaffarian D, Psaty BM, Huertas-Vazquez A, Tardif JC, Albert CM, Lyytikäinen LP, Arking DE, Kääb S, Huikuri HV, Krijthe BP, Eijgelsheim M, Wang YA, Reinier K, Lehtimäki T, Pulit SL, Brugada R, Müller-Nurasyid M, **Newton-Cheh CH**, Karhunen PJ, Stricker BH, Goyette P, Rotter JI, Chugh SS, Chakravarti A, Jouven X, Siscovick DS. Common variation in fatty acid metabolic genes and risk of incident sudden cardiac arrest. *Heart Rhythm.* 2014 Mar; 11(3): 471-7. doi: 10.1016/j.hrthm.2014.01.008. Epub 2014 Jan 10. PMID: PMC3966996.
125. Tragante V, Barnes MR, Ganesh SK, Lanktree MB, Guo W, Franceschini N, Smith EN, Johnson T, Holmes MV, Padmanabhan S, Karczewski KJ, Almoguera B, Barnard J, Baumert J, Chang YP, Elbers CC, Farrall M, Fischer ME, Gaunt TR, Gho JM, Gieger C, Goel A, Gong Y, Isaacs A, Kleber ME, Mateo Leach I, McDonough CW, Meijis MF, Melander O, Nelson CP, Nolte IM, Pankratz N, Price TS, Shaffer J, Shah S, Tomaszewski M, van der Most PJ, Van Iperen EP, Vonk JM, Witkowska

K, Wong CO, Zhang L, Beitelshes AL, Berenson GS, Bhatt DL, Brown M, Burt A, Cooper-Dehoff RM, Connell JM, Cruickshanks KJ, Curtis SP, Davey-Smith G, Delles C, Gansevoort RT, Guo X, Haiqing S, Hastie CE, Hofker MH, Hovingh GK, Kim DS, Kirkland SA, Klein BE, Klein R, Li YR, Maiwald S, **Newton-Cheh C**, O'Brien ET, Onland-Moret NC, Palmas W, Parsa A, Penninx BW, Pettinger M, Vasan RS, Ranchalis JE, M Ridker P, Rose LM, Sever P, Shimbo D, Steele L, Stolk RP, Thorand B, Trip MD, van Duijn CM, Verschuren WM, Wijmenga C, Wyatt S, Young JH, Zwinderman AH, Bezzina CR, Boerwinkle E, Casas JP, Caulfield MJ, Chakravarti A, Chasman DI, Davidson KW, Doevendans PA, Dominiczak AF, Fitzgerald GA, Gums JG, Fornage M, Hakonarson H, Halder I, Hillege HL, Illig T, Jarvik GP, Johnson JA, Kastelein JJ, Koenig W, Kumari M, März W, Murray SS, O'Connell JR, Oldehinkel AJ, Pankow JS, Rader DJ, Redline S, Reilly MP, Schadt EE, Kottke-Marchant K, Snieder H, Snyder M, Stanton AV, Tobin MD, Uitterlinden AG, van der Harst P, van der Schouw YT, Samani NJ, Watkins H, Johnson AD, Reiner AP, Zhu X, de Bakker PI, Levy D, Asselbergs FW, Munroe PB, Keating BJ. Gene-centric Meta-analysis in 87,736 Individuals of European Ancestry Identifies Multiple Blood-Pressure-Related Loci. *Am J Hum Genet.* 2014 Mar 6; 94(3):349-60. doi: 10.1016/j.ajhg.2013.12.016. Epub 2014 Feb 20. PMID: PMC3951943.

126. Magnani JW, Brody JA, Prins BP, Arking DE, Lin H, Yin X, Liu CT, Morrison AC, Zhang F, Spector TD, Alonso A, Bis JC, Heckbert SR, Lumley T, Sitlani CM, Cupples LA, Lubitz SA, Soliman EZ, Pulit SL, **Newton-Cheh C**, O'Donnell CJ, Ellinor PT, Benjamin EJ, Muzny DM, Gibbs RA, Santibanez J, Taylor HA, Rotter JI, Lange LA, Psaty BM, Jackson R, Rich SS, Boerwinkle E, Jamshidi Y, Sotoodehnia N; CHARGE Consortium; NHLBI Exome Sequencing Project (ESP); UK10K. Sequencing of SCN5A identifies rare and common variants associated with cardiac conduction: Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium. *Circ Cardiovasc Genet.* 2014 Jun; 7(3):365-73. doi: 10.1161/CIRCGENETICS.113.000098.

Kapoor A, Sekar RB, Hansen NF, Fox-Talbot K, Morley M, Pihur V, Chatterjee S, Brandimarto J, Moravec CS, Pulit SL; QT Interval-International GWAS Consortium, Pfeufer A, Mullikin J, Ross M, Green ED, Bentley D, **Newton-Cheh C**, Boerwinkle E, Tomaselli GF, Cappola TP, Arking DE, Halushka MK, Chakravarti A. An enhancer polymorphism at the cardiomyocyte intercalated disc protein NOS1AP locus is a major regulator of the QT interval. *Am J Hum Genet.* 2014 Jun 5; 94(6):854-69. doi: 10.1016/j.ajhg.2014.05.001. Epub 2014 May 22. PMID: PMC4121472.

127. Ganesh SK, Chasman DI, Larson MG, Guo X, Verwoert G, Bis JC, Gu X, Smith AV, Yang ML, Zhang Y, Ehret G, Rose LM, Hwang SJ, Papanicolau GJ, Sijbrands EJ, Rice K, Eiriksdottir G, Pihur V, Ridker PM, Vasan RS, **Newton-Cheh C**; Global Blood Pressure Genetics Consortium, Raffel LJ, Amin N, Rotter JI, Liu K, Launer LJ, Xu M, Caulfield M, Morrison AC, Johnson AD, Vaidya D, Dehghan A, Li G, Bouchard C, Harris TB, Zhang H, Boerwinkle E, Siscovick DS, Gao W, Uitterlinden AG, Rivadeneira F, Hofman A, Willer CJ, Franco OH, Huo Y, Witteman JC, Munroe PB, Gudnason V, Palmas W, van Duijn C, Fornage M, Levy D, Psaty BM, Chakravarti A. Effects of long-term averaging of quantitative blood pressure traits on the detection of genetic associations. *Am J Hum Genet.* 2014 Jul 3; 95(1):49-65. doi: 10.1016/j.ajhg.2014.06.002. Epub 2014 Jun 26. PMID: PMC4085637.

128. Rosenberg MA, Kaplan RC, Siscovick DS, Psaty BM, Heckbert SR, **Newton-Cheh C**, Mukamal KJ. Genetic variants related to height and risk of atrial fibrillation: the cardiovascular health study. *Am J Epidemiol.* 2014 Jul 15; 180(2):215-22. doi: 10.1093/aje/kwu126. Epub 2014 Jun 18. PMID: PMC4082343.

129. Lundby A, Rossin EJ, Steffensen AB, Acha MR, **Newton-Cheh C**, Pfeufer A, Lynch SN; QT Interval International GWAS Consortium (QT-IGC), Olesen SP, Brunak S, Ellinor PT, Jukema JW, Trompet S, Ford I, Macfarlane PW, Krijthe BP, Hofman A, Uitterlinden AG, Stricker BH, Nathoe HM, Spiering W, Daly MJ, Asselbergs FW, van der Harst P, Milan DJ, de Bakker PI, Lage K, Olsen JV. Annotation of loci from genome-wide association studies using tissue-specific quantitative interaction proteomics. *Nat Methods*. 2014 Aug; 11(8):868-74. doi: 10.1038/nmeth.2997. Epub 2014 Jun 22. PMID: PMC4117722.

Arking DE, Pulit SL, Crotti L, van der Harst P, Munroe PB, Koopmann TT, Sotoodehnia N, Rossin EJ, Morley M, Wang X, Johnson AD, Lundby A, Gudbjartsson DF, Noseworthy PA, Eijgelsheim M, Bradford Y, Tarasov KV, Dörr M, Müller-Nurasyid M, Lahtinen AM, Nolte IM, Smith AV, Bis JC, Isaacs A, Newhouse SJ, Evans DS, Post WS, Waggott D, Lytikäinen LP, Hicks AA, Eisele L, Ellinghaus D, Hayward C, Navarro P, Ulivi S, Tanaka T, Tester DJ, Chatel S, Gustafsson S, Kumari M, Morris RW, Naluai AT, Padmanabhan S, Kluttig A, Strohmer B, Panayiotou AG, Torres M, Knoflach M, Hubacek JA, Slowikowski K, Raychaudhuri S, Kumar RD, Harris TB, Launer LJ, Shuldiner AR, Alonso A, Bader JS, Ehret G, Huang H, Kao WH, Strait JB, Macfarlane PW, Brown M, Caulfield MJ, Samani NJ, Kronenberg F, Willeit J; CARE Consortium; COGENT Consortium, Smith JG, Greiser KH, Meyer Zu Schwabedissen H, Werdan K, Carella M, Zelante L, Heckbert SR, Psaty BM, Rotter JI, Kolcic I, Polašek O, Wright AF, Griffin M, Daly MJ; DCCT/EDIC, Arnar DO, Hólm H, Thorsteinsdóttir U; eMERGE Consortium, Denny JC, Roden DM, Zuvich RL, Emilsson V, Plump AS, Larson MG, O'Donnell CJ, Yin X, Bobbo M, D'Adamo AP, Iorio A, Sinagra G, Carracedo A, Cummings SR, Nalls MA, Jula A, Kontula KK, Marjamaa A, Oikarinen L, Perola M, Porthan K, Erbel R, Hoffmann P, Jöckel KH, Kälisch H, Nöthen MM; HRGEN Consortium, den Hoed M, Loos RJ, Thelle DS, Gieger C, Meitinger T, Perz S, Peters A, Prucha H, Sinner MF, Waldenberger M, de Boer RA, Franke L, van der Vleuten PA, Beckmann BM, Martens E, Bardai A, Hofman N, Wilde AA, Behr ER, Dalageorgou C, Giudicessi JR, Medeiros-Domingo A, Barc J, Kyndt F, Probst V, Ghidoni A, Insolia R, Hamilton RM, Scherer SW, Brandimarto J, Margulies K, Moravec CE, Greco M FD, Fuchsberger C, O'Connell JR, Lee WK, Watt GC, Campbell H, Wild SH, El Mokhtari NE, Frey N, Asselbergs FW, Mateo Leach I, Navis G, van den Berg MP, van Veldhuisen DJ, Kellis M, Krijthe BP, Franco OH, Hofman A, Kors JA, Uitterlinden AG, Witteman JC, Kedenko L, Lamina C, Oostra BA, Abecasis GR, Lakatta EG, Mulas A, Orrú M, Schlessinger D, Uda M, Markus MR, Völker U, Snieder H, Spector TD, Arnlöv J, Lind L, Sundström J, Syvänen AC, Kivimäki M, Kähönen M, Mononen N, Raitakari OT, Viikari JS, Adamkova V, Kiechl S, Brion M, Nicolaidis AN, Paulweber B, Haerting J, Dominiczak AF, Nyberg F, Whincup PH, Hingorani AD, Schott JJ, Bezzina CR, Ingelsson E, Ferrucci L, Gasparini P, Wilson JF, Rudan I, Franke A, Mühleisen TW, Pramstaller PP, Lehtimäki TJ, Paterson AD, Parsa A, Liu Y, van Duijn CM, Siscovick DS, Gudnason V, Jamshidi Y, Salomaa V, Felix SB, Sanna S, Ritchie MD, Stricker BH, Stefansson K, Boyer LA, Cappola TP, Olsen JV, Lage K, Schwartz PJ, Käb S, Chakravarti A, Ackerman MJ, Pfeufer A, de Bakker PI, **Newton-Cheh C**. Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. *Nat Genet*. 2014 Aug; 46(8):826-36. doi: 10.1038/ng.3014. Epub 2014 Jun 22. PMID: PMC4124521.

*denotes equal contribution

Other peer-reviewed publications

1. **Newton-Cheh C**, Hirschhorn JN. Genetic association studies of complex traits: design and analysis issues. *Mutation Research*. 2005; 573(1-2): 54-69. Review

2. **Newton-Cheh C**, Shah R. Genetic determinants of QT interval variation and sudden cardiac death. *Curr Opin Genet Dev.* 2007; 17(3): 213-221. Review
3. Noseworthy PA and **Newton-Cheh C**. Genetic determinants of sudden cardiac death. *Circulation.* 2008 118(18): 1854-63. Review
4. Smith JG, **Newton-Cheh C**. Genome-wide association study in humans. *Methods in Molecular Biology*, 2009; 573: 231-258. Review
5. Arora P, **Newton-Cheh C**. Blood pressure and human genetic variation in the general population. *Curr Opin Cardiol.* 2010; 25: 229-237. Review. PMID: PMC2947583.
6. Rodriguez I, Erdman A, Padhi D, Garnett CE, Zhao H, Targum SL, Balakrishnan S, Strnadova C, Viner N, Geiger MJ, **Newton-Cheh C**, Litwin J, Pugsley MK, Sager PT, Krucoff MW, Finkle JK. Electrocardiographic assessment for therapeutic proteins--scientific discussion. *Am Heart J.* 2010 Oct; 160(4): 627-34. Review.
7. Sauer AJ, **Newton-Cheh C**. Clinical and genetic determinants of torsade de pointes risk. *Circulation.* 2012 Apr 3; 125(13):1684-94. Review. PMID: PMC3347483.
8. Padmanabhan S, **Newton-Cheh C**, Dominiczak AF. Genetic basis of blood pressure and hypertension. *Trends Genet.* 2012 Aug; 28(8):397-408. Epub 2012 May 21. Review

Research publications without named authorship

1. Ikram MK, Xueling S, Jensen RA, Cotch MF, Hewitt AW, Ikram MA, Wang JJ, Klein R, Klein BE, Breteler MM, Cheung N, Liew G, Mitchell P, Uitterlinden AG, Rivadeneira F, Hofman A, de Jong PT, van Duijn CM, Kao L, Cheng CY, Smith AV, Glazer NL, Lumley T, McKnight B, Psaty BM, Jonasson F, Eiriksdottir G, Aspelund T; Global BPgen Consortium, Harris TB, Launer LJ, Taylor KD, Li X, Iyengar SK, Xi Q, Sivakumaran TA, Mackey DA, Macgregor S, Martin NG, Young TL, Bis JC, Wiggins KL, Heckbert SR, Hammond CJ, Andrew T, Fahy S, Attia J, Holliday EG, Scott RJ, Islam FM, Rotter JI, McAuley AK, Boerwinkle E, Tai ES, Gudnason V, Siscovick DS, Vingerling JR, Wong TY. Four novel Loci (19q13, 6q24, 12q24, and 5q14) influence the microcirculation in vivo. *PLoS Genet.* 2010 Oct 28; 6(10): e1001184. PMID: PMC2965750.
2. Tabara Y, Kohara K, Kita Y, Hirawa N, Katsuya T, Ohkubo T, Hiura Y, Tajima A, Morisaki T, Miyata T, Nakayama T, Takashima N, Nakura J, Kawamoto R, Takahashi N, Hata A, Soma M, Imai Y, Kokubo Y, Okamura T, Tomoike H, Iwai N, Ogiwara T, Inoue I, Tokunaga K, Johnson T, Caulfield M, Munroe P; Global Blood Pressure Genetics Consortium, Umemura S, Ueshima H, Miki T. Common variants in the ATP2B1 gene are associated with susceptibility to hypertension: the Japanese Millennium Genome Project. *Hypertension.* 2010 Nov; 56(5): 973-80. Epub 2010 Oct 4.
3. Fox ER, Young JH, Li Y, Dreisbach AW, Keating BJ, Musani SK, Liu K, Morrison AC, Ganesh S, Kutlar A, Ramachandran VS, Polak JF, Fabsitz RR, Dries DL, Farlow DN, Redline S, Adeyemo A, Hirschorn JN, Sun YV, Wyatt SB, Penman AD, Palmas W, Rotter JI, Townsend RR, Doumatey AP, Tayo BO, Mosley TH Jr, Lyon HN, Kang SJ, Rotimi CN, Cooper RS, Franceschini N, Curb JD, Martin LW, Eaton CB, Kardina SL, Taylor HA, Caulfield MJ, Ehret GB, Johnson T; The International Consortium for Blood Pressure Genome-wide Association Studies (ICBP-GWAS); CARDIoGRAM consortium; CKDGen consortium; KidneyGen consortium; EchoGen consortium; CHARGE-HF

consortium, Chakravarti A, Zhu X, Levy D. Association of genetic variation with systolic and diastolic blood pressure among African Americans: the Candidate Gene Association Resource study. *Hum Mol Genet.* 2011 Apr 8. [Epub ahead of print]. PMID: PMC3090190.

4. Sim X, Jensen RA, Ikram MK, Cotch MF, Li X, MacGregor S, Xie J, Smith AV, Boerwinkle E, Mitchell P, Klein R, Klein BE, Glazer NL, Lumley T, McKnight B, Psaty BM, de Jong PT, Hofman A, Rivadeneira F, Uitterlinden AG, van Duijn CM, Aspelund T, Eiriksdottir G, Harris TB, Jonasson F, Launer LJ; Wellcome Trust Case Control Consortium 2, Attia J, Baird PN, Harrap S, Holliday EG, Inouye M, Rohtchina E, Scott RJ, Viswanathan A; Global BPGen Consortium, Li G, Smith NL, Wiggins KL, Kuo JZ, Taylor KD, Hewitt AW, Martin NG, Montgomery GW, Sun C, Young TL, Mackey DA, van Zuydam NR, Doney AS, Palmer CN, Morris AD, Rotter JI, Tai ES, Gudnason V, Vingerling JR, Siscovick DS, Wang JJ, Wong TY. Genetic loci for retinal arteriolar microcirculation. *PLoS One.* 2013 Jun 12;8(6):e65804. doi: 10.1371/journal.pone.0065804. Print 2013. PMID: PMC3680438.
5. O'Connor TD, Kiezun A, Bamshad M, Rich SS, Smith JD, Turner E; NHLBIGO Exome Sequencing Project; ESP Population Genetics, Statistical Analysis Working Group, Leal SM, Akey JM. Fine-scale patterns of population stratification confound rare variant association tests. *PLoS One.* 2013 Jul 4;8(7):e65834. doi: 10.1371/journal.pone.0065834. Print 2013. PMID: PMC3701690.
6. Fu W, O'Connor TD, Jun G, Kang HM, Abecasis G, Leal SM, Gabriel S, Altshuler D, Shendure J, Nickerson DA, Bamshad MJ; NHLBI Exome Sequencing Project, Akey JM. Analysis of 6,515 exomes reveals the recent origin of most human protein-coding variants. *Nature.* 2013 Jan 10; 493(7431):216-20. PMID: PMC3676746
7. Johnsen JM, Auer PL, Morrison AC, Jiao S, Wei P, Haessler J, Fox K, McGee SR, Smith JD, Carlson CS, Smith N, Boerwinkle E, Kooperberg C, Nickerson DA, Rich SS, Green D, Peters U, Cushman M, Reiner AP; NHLBI Exome Sequencing Project. Common and rare von Willebrand factor (VWF) coding variants, VWF levels, and factor VIII levels in African Americans: the NHLBI Exome Sequencing Project. *Blood.* 2013 Jul 25; 122(4):590-7. doi: 10.1182/blood-2013-02-485094. Epub 2013 May 20. PMID: PMC3724194.
8. Chen HH, Anstrom KJ, Givertz MM, Stevenson LW, Semigran MJ, Goldsmith SR, Bart BA, Bull DA, Stehlik J, LeWinter MM, Konstam MA, Huggins GS, Rouleau JL, O'Meara E, Tang WH, Starling RC, Butler J, Deswal A, Felker GM, O'Connor CM, Bonita RE, Margulies KB, Cappola TP, Ofili EO, Mann DL, Dávila-Román VG, McNulty SE, Borlaug BA, Velazquez EJ, Lee KL, Shah MR, Hernandez AF, Braunwald E, Redfield MM; NHLBI Heart Failure Clinical Research Network. Low-dose dopamine or low-dose nesiritide in acute heart failure with renal dysfunction: the ROSE acute heart failure randomized trial. *JAMA.* 2013 Dec 18; 310(23):2533-43. doi: 10.1001/jama.2013.282190.
9. Lange LA, Hu Y, Zhang H, Xue C, Schmidt EM, Tang ZZ, Bizon C, Lange EM, Smith JD, Turner EH, Jun G, Kang HM, Peloso G, Auer P, Li KP, Flannick J, Zhang J, Fuchsberger C, Gaulton K, Lindgren C, Locke A, Manning A, Sim X, Rivas MA, Holmen OL, Gottesman O, Lu Y, Ruderfer D, Stahl EA, Duan Q, Li Y, Durda P, Jiao S, Isaacs A, Hofman A, Bis JC, Correa A, Griswold ME, Jakobsdottir J, Smith AV, Schreiner PJ, Feitosa MF, Zhang Q, Huffman JE, Crosby J, Wassel CL, Do R, Franceschini N, Martin LW, Robinson JG, Assimes TL, Crosslin DR, Rosenthal EA, Tsai M, Rieder MJ, Farlow DN, Folsom AR, Lumley T, Fox ER, Carlson CS, Peters U, Jackson RD, van

Duijn CM, Uitterlinden AG, Levy D, Rotter JI, Taylor HA, Gudnason V Jr, Siscovick DS, Fornage M, Borecki IB, Hayward C, Rudan I, Chen YE, Bottinger EP, Loos RJ, Sætrum P, Hveem K, Boehnke M, Groop L, McCarthy M, Meitinger T, Ballantyne CM, Gabriel SB, O'Donnell CJ, Post WS, North KE, Reiner AP, Boerwinkle E, Psaty BM, Altshuler D, Kathiresan S, Lin DY, Jarvik GP, Cupples LA, Kooperberg C, Wilson JG, Nickerson DA, Abecasis GR, Rich SS, Tracy RP, Willer CJ; NHLBI Grand Opportunity Exome Sequencing Project. Whole-exome sequencing identifies rare and low-frequency coding variants associated with LDL cholesterol. *Am J Hum Genet.* 2014 Feb 6; 94(2):233-45. doi: 10.1016/j.ajhg.2014.01.010. PMID: PMC3928660.

Non-peer reviewed scientific or medical publications/materials in print or other media

1. Kathiresan S*, **Newton-Cheh C***, Gerszten RE. On the interpretation of genetic association studies. *Eur Heart J.* 2004 Aug; 25(16):1378-81. Editorial.
2. **Newton-Cheh C**, O'Donnell CJ. Sex differences and genetic associations with myocardial infarction. *JAMA.* 2004 Jun 23; 291(24):3008-10. Editorial.
3. **Newton-Cheh C**, Larson M, Kathiresan S, O'Donnell C. On the significance of linkage studies of complex traits. *Am J Hum Genet.* 2004 Jul; 75(1):151-2. Editorial.
4. **Newton-Cheh C**, Semigran MJ. Selection and Management of Cardiac Transplantation Candidates. Treatment of Advanced Heart Disease. Eds Baughman KL, Baumgartner WA. New York: Taylor & Francis, 2006; 491-510. Chapter.
5. Smith JG, **Newton-Cheh C**. Candidate gene and genome-wide association studies. Cardiovascular Genetics and Genomics. Ed. Roden D. Wiley-Blackwell, 2009; 3-19. Chapter.
6. **Newton-Cheh C**, Smith JG. What can human genetics teach us about the causes of cardiovascular disease? *Journal of the American College of Cardiology.* 2010 Jun 22; 55(25): 2843-5. Editorial.
9. **Newton-Cheh C**. What can genetic studies of left ventricular mass tell us? *Circ Cardiovasc Genet.* 2011 Dec 1; 4(6):581-4. Editorial. PMID: PMC3247759.

Narrative Report

My laboratory in the Massachusetts General Hospital Center for Human Genetic Research and Cardiovascular Research Center is focused on identification of the genetic factors that underlie variation in hypertension and arrhythmia risk by large population-based studies of blood pressure and QT interval duration, determination of the role of these factors in human physiology, elucidation of novel mechanisms of disease that highlight potential therapeutic targets, with the goal of improving human health at the population level. I spend 83% of my time on research, 10% on patient care, 5% on teaching and 2% on administration. Areas of excellence in the Newton-Cheh laboratory include the genetics of blood pressure and of myocardial repolarization. Elevated blood pressure is an important modifiable risk factor for stroke, heart failure, myocardial infarction and end-stage renal disease. Electrocardiographic QT interval prolongation at rest is a risk factor for sudden cardiac death and occurs as a potentially fatal cardiotoxic side effect of many medications, leading to loss of valuable compounds from the drug development pipeline.

I established that a common genetic variant in the 3' untranslated region of *NPPA* is associated with higher plasma atrial natriuretic peptide (ANP) concentration, and as a consequence lower blood pressure

and hypertension risk (Newton-Cheh, *Nature Genetics* 2009). This was the first *in vivo* demonstration of the role of inter-individual variability in the natriuretic peptide system in blood pressure regulation and was the first common variant for blood pressure reported. This work served as the basis of my R01, on which I am PI, to conduct translational studies using dietary sodium challenge and acute intravenous saline administration in individuals selected on the basis of natriuretic peptide genotype to define the impact of the natriuretic peptide system on acute and chronic responses to salt exposure. We have just published in the *Journal of Clinical Investigation* the results of this work in which we demonstrate that the genetic effect on ANP is as strong as the impact of a 20-fold reduction in sodium intake and identify that the variant acts by interrupting a previously unrecognized negative regulator of ANP, micro-RNA miR-425. We have filed a patent on the use of antagonists of miR-425 to treat hypertension.

I have led international consortia (Steering Committee co-chair) that have completed genome-wide association studies identifying 60 novel genetic loci that regulate blood pressure in humans, (Newton-Cheh, *Nature Genetics* 2009; Ehret, *Nature* 2011, manuscript in preparation). The first paper was recognized by the American Heart Association as one of the top ten discoveries of 2009. These unbiased surveys of the entire genome have focused attention on that natriuretic peptide/guanylate cyclase receptor A and nitric oxide/soluble guanylate cyclase (sGC) systems, both of which cause vasodilation and natriuresis via production of the intracellular second messenger cyclic guanosine monophosphate (cGMP). This work served as the basis for an R01 on which I am principal investigator, which has just been funded. Through this grant, we are recruiting individuals on the basis of their genotype at a SNP in sGC and administering inhaled nitric oxide and performing platelet assays to determine the role of the SNP in sGC-mediated cGMP response, as well as attempting to elucidate the molecular mechanism underlying the blood pressure association.

I have also led the QTGEN and QT-IGC consortia (Steering Committee chair) that have identified in total 68 independent genetic variants at 35 loci that influence QT interval variation (Newton-Cheh, *Nature Genetics* 2009; manuscript under review). While several loci harboring common QT variants are also known to harbor rare variants that cause congenital Long QT Syndrome, most of the loci are novel. Collectively, these studies highlight the role of myocardial calcium signaling as a mediator of repolarization. My laboratory is currently testing the hypothesis that these variants in aggregate mark individuals at risk of cardiotoxic side effects of marketed pharmaceutical agents through physiologic experiments in the GCRC setting.

In the area of teaching, I served as chief medical resident and administrative cardiology fellow at the Massachusetts General Hospital in part for my skills as a teacher. I have attended on the general medical wards at MGH (2004, 05, 06), in the coronary care unit (2006, 07, 08, 09, 10, 11, 12), on the transplant service (2004, 05, 06, 07, 08, 09, 10, 11, 12, 13) and cardiology consultation service (2004, 05, 06). I tutor in the HMS first year genetics course (2006, 07, 08, 09, 10, 12) and teach in (2008, 09, 10, 11, 12, 13) and co-direct since 2009 the Genetics 228 course in the Biological and Biomedical Sciences Program at Harvard Medical School, and direct the Primer on Medical and Population Genetics course at the Broad Institute of Harvard and MIT, now in its 8th year.

My clinical contributions include serving as a staff cardiologist in the Heart Failure and Transplantation Section, directing the care of inpatients and outpatients requiring advanced heart failure therapies including ventricular assist device therapy and transplantation. I carry a panel of 300 outpatients, with clinic every two weeks.

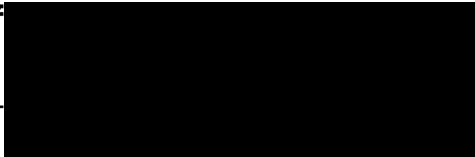
In summary, my laboratory is generating key insights into the genetic basis of blood pressure regulation and myocardial repolarization, which have implications for our understanding of the physiology of hypertension and cardiotoxic drug response. We are expanding our research efforts to translate these genetic findings with the use of high-resolution phenotyping of carriers of known genetic factors that operate at the population level, with the promise that we may improve both the efficacy and reduce the toxicity of therapies for the prevention of human disease.

**Committee on Energy and Commerce
U.S. House of Representatives**

Witness Disclosure Requirement - "Truth in Testimony"
Required by House Rule XI, Clause 2(g)

1. Your Name: Christopher Newton-Cheh, MD, MPH		
2. Are you testifying on behalf of the Federal, or a State or local government entity?	Yes	No X
3. Are you testifying on behalf of an entity that is not a government entity?	Yes X	No
4. Other than yourself, please list which entity or entities you are representing: American Heart Association		
5. Please list any Federal grants or contracts (including subgrants or subcontracts) that you or the entity you represent have received on or after October 1, 2011: HL088456, HL098283, HL113933		
6. If your answer to the question in item 3 in this form is "yes," please describe your position or representational capacity with the entity or entities you are representing: I am a volunteer for the American Heart Association, a non-profit organization focused on the prevention and treatment of cardiovascular disease and stroke		
7. If your answer to the question in item 3 is "yes," do any of the entities disclosed in item 4 have parent organizations, subsidiaries, or partnerships that you are not representing in your testimony?	Yes	No X
8. If the answer to the question in item 3 is "yes," please list any Federal grants or contracts (including subgrants or subcontracts) that were received by the entities listed under the question in item 4 on or after October 1, 2011, that exceed 10 percent of the revenue of the entities in the year received, including the source and amount of each grant or contract to be listed: NA		
9. Please attach your curriculum vitae to your completed disclosure form. see attached		

Signature: _____



Date: 9/8/14 _____