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# The need for genetics/genomics literacy in cardiovascular/stroke care



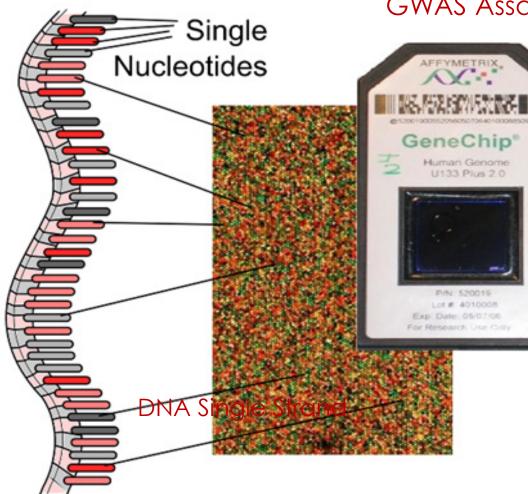
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#### State-of-the-science



Tremendous progress in understanding of the genetic basis of cardiovascular and stroke disorders via GWAS studies



#### GWAS Assay

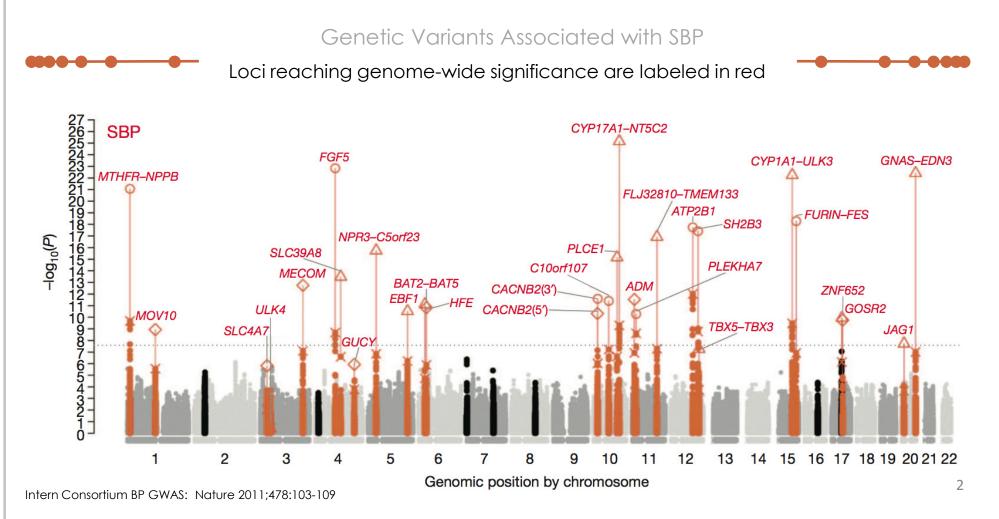
#### **GWAS** studies

selectively assay a subset of nucleotides ("singlenucleotide polymorphisms," SNPs) on all chromosomes in the genome. These polymorphisms are relatively common in populations.

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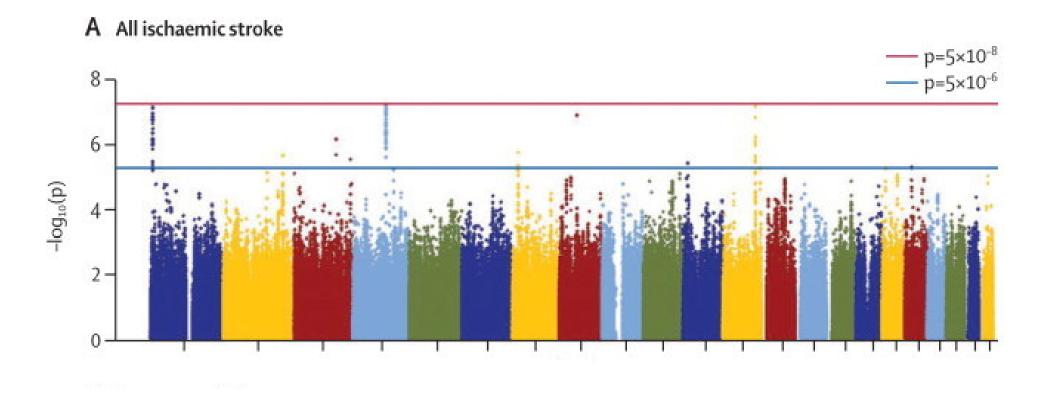
#### GWAS of 200,000 Individuals of European Descent







#### Manhattan plots for All Ischemic Stroke –log10(p) by genomic position

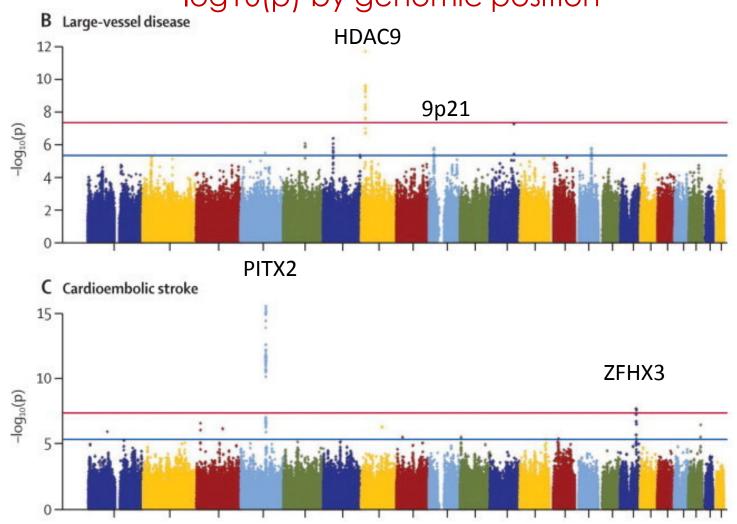


Lancet Neurol. 2012 November; 11(11): 951-962

American Heart Stroke Association Association. Association Stroke

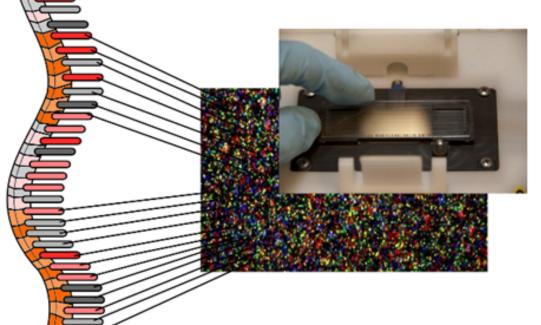


#### Manhattan plots for Large-Vessel Disease and Cardioembolic Stroke –log10(p) by genomic position





#### Exome and Whole Genome Sequencing



Protein

coding region

**DNA Single Strand** 

Whole **exome/ genome** sequencing studies selectively assay the protein coding regions ("exons") on all chromosomes in the genome.

Sequence data finds polymorphisms that are common and rare in populations.



#### State-of-the-science

- Exome sequencing has discovered new genes involved in cholesterol metabolism and blood pressure control
- Functional characterization of novel cardiovascular/stroke loci and genes is underway – among the most advanced of any discipline



#### From genotype to phenotype

# ARTICLES

# Biological, clinical and population relevance of 95 loci for blood lipids

A list of authors and their affiliations appears at the end of the paper.

Plasma concentrations of total cholesterol, low-density lipoprotein cholesterol, high-density lipoprotein cholesterol and triglycerides are among the most important risk factors for coronary artery disease (CAD) and are targets for therapeutic

#### ARTICLES

# From noncoding variant to phenotype via SORT1 at the 1p13 cholesterol locus

Kiran Musunuru<sup>1,2,3</sup>\*, Alanna Strong<sup>4</sup>\*, Maria Frank-Kamenetsky<sup>5</sup>, Noemi E. Lee<sup>1</sup>, Tim Ahfeldt<sup>1,6</sup>, Katherine V. Sachs<sup>4</sup>, Xiaoyu Li<sup>4</sup>, Hui Li<sup>4</sup>, Nicolas Kuperwasser<sup>1</sup>, Vera M. Ruda<sup>1</sup>, James P. Pirruccello<sup>1,2</sup>, Brian Muchmore<sup>7</sup>, Ludmila Prokunina-Olsson<sup>7</sup>, Jennifer L. Hall<sup>2,8</sup>, Eric E. Schadt<sup>9</sup>, Carlos R. Morales<sup>10</sup>,



# **Clinical applications**

- Despite the wealth of new knowledge...
- ...cardiovascular and stroke clinical care has not yet been significantly impacted
- Translation into the clinic will ensue and accelerate over the next 5-10 years



#### Cardiovascular risk prediction

- Common DNA variants
- Rare DNA variants



### **Common DNA variants**

- In most cases, have been discovered by GWAS
- Typically have small effects no one variant will singlehandedly determine if a patient will develop disease
- Can aggregate the effects of many variants by combining them into genetic risk scores



#### Genetic risk scores

- Genetic risk scores have been tested for predictive power for several cardiovascular/stroke disorders
- Modest predictive power for incident coronary heart disease – high genetic risk score confers only a ~70% increase in risk
- Limited clinical usefulness (except perhaps for intermediate-risk, "on the fence" patients)



## Rare DNA variants

- Rare DNA variants have potential to have large effects, may increase (or decrease) risk several-fold
- However, each variant only found in one or a few people
- Genetic risk scores with common DNA variants do not capture – and may be trumped by – rare variants



#### Needle in a haystack

- Whole-genome sequencing will identify hundreds of rare variants in each person – most of which are irrelevant to disease
- How to know which rare variants are clinically important? Huge challenge for the field
- May fall in genes already implicated in disease, e.g., hypertrophic cardiomyopathy (HCM)



#### What does it mean to have a mutation?

- Even if in known gene, two problems
- First: the particular mutation may or may not affect gene function difficult to predict *a priori*
- Second: even if the mutation changes gene function, penetrance may be affected by genetic background
- Even within a family, some members with HCM mutation will develop disease, some will not



#### Dangers of genetic information

- DTC testing (e.g., 23andMe) informs patients of genetic risk scores but (for now) not rare variants – possibility of false reassurance (or false worry)
- Problem will get worse with DTC genome sequencing
- Providers need to be aware of the limitations of this information to appropriately counsel patients
- Providers' lack of knowledge about genetics will serve patients poorly – education is critical



#### Pharmacogenomics

- Getting the right dose of the right medication to the right patient
- Though integrated into clinical practice in some disciplines (e.g., oncology), not yet adopted in cardiovascular/stroke care
- Several emerging applications: warfarin, clopidogrel, beta-blockers, lipid-modifying medications



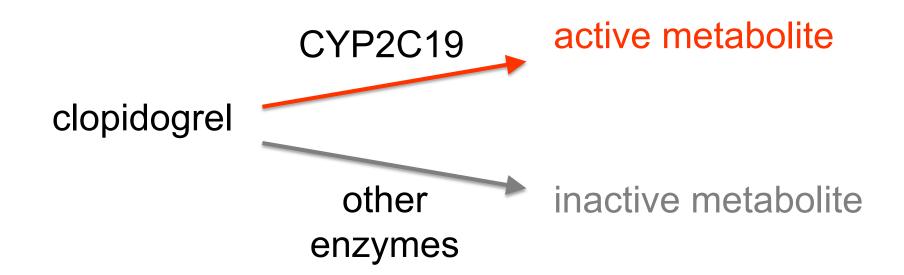
# Clopidogrel and CYP2C19

- Patients presenting with myocardial infarction
- Routine practice is to give anti-platelet agent clopidogrel (Plavix), a type of thienopyridine
- Clopidogrel reduces risk of future cardiovascular event as well as the risk of in-stent thrombosis



# Clopidogrel and CYP2C19

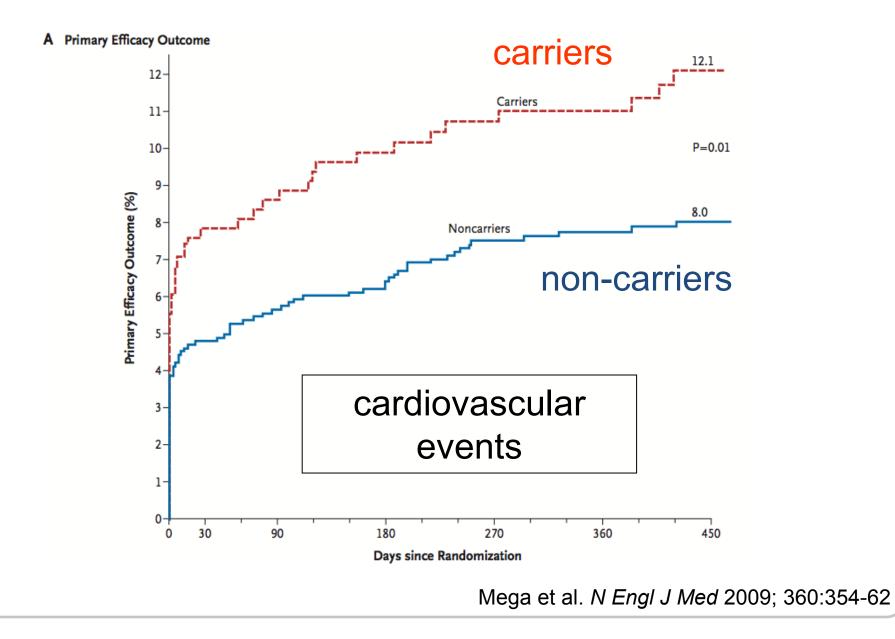
• CYP2C19 encodes a key enzyme in making active metabolite of clopidogrel



 Reduced-function CYP2C19 alleles result in decreased clopidogrel efficacy (but not other thienopyridines) – has lead to FDA black box warning



#### Effect of CYP2C19 reduced-function alleles





#### Pharmacogenomic strategy using CYP2C19

patient presents to hospital with myocardial infarction



give clopidogrel, cath lab, etc.

normal alleles

reduced-function alleles

give a different thienopyridine (e.g., ticagrelor) or 2X dose of clopidogrel, cath lab, etc.



### Clopidogrel and CYP2C19

- Point-of-care CYP2C19 genotyping presently being piloted by a number of hospitals
- Remains to be validated by clinical studies unclear which patients, if any, would benefit
- Must be validated before can be recommended for routine clinical use



#### Adoption of pharmacogenomics

- With several cardiovascular pharmacogenomic applications under evaluation, likely that at least one will be validated within 5 years
- Expect rapid adoption by academic centers and large healthcare networks
- Expect slower adoption by individual practitioners, small group practices due to "knowledge gap" – education is critical



#### **Provider education**

- Although lack of provider education on genetics/genomics issues not currently jeopardizing clinical care – anticipate that negative effects of a "knowledge gap" will emerge in 5-10 years
- AHA/ASA is moving to address the anticipated need for provider genetics/genomics literacy in two ways



#### Scientific Statements

- To date, AHA/ASA has published several scientific statements related to genetics – e.g., "Genetics and Genomics for the Prevention and Treatment of Cardiovascular Disease" – but these are oriented to state-of-the-science and policy, not provider education
- Currently has a working group formulating a statement on "Use of Genetics and Genomics in Cardiovascular and Stroke Patient Care" – exclusively focused on provider education



# **Outline of Statement**

- Primer on genetics and genomics
- Monogenic cardiovascular and stroke disorders
- Polygenic cardiovascular and stroke disorders
- Cardiovascular and stroke risk prediction
- Pharmacogenomics
- Cardiovascular and stroke risk prediction
- Social and ethical implications
- Educational resources



#### Massively open online course (MOOC)

- AHA/ASA monitoring latest trends in education
- Formulating an online course in Genetics/Genomics that will cover all of the topics outlined in the "Use of Genetics and Genomics in Cardiovascular and Stroke Patient Care" statement
- Plan to make course freely available to the biomedical community – pitched at an undergraduate level, with target audience of physicians, RNs, pharmacists, etc.



#### Massively open online course (MOOC)

- Will eventually be supplemented with modules on specific cardiovascular genetic disorders – e.g., hypertrophic cardiomyopathy, arrhythmogenic right ventricular dysplasia, long QT syndrome, Brugada syndrome
- Initial versions in English, to potentially be followed by translations into other languages



# **Special Thanks To:**

# Dr. Kiran Musunuru MD PhD MPH Assistant Professor Harvard University