

A new paradigm for pharmacogenomic discoveries: Capturing drug response during surgery



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Motivation

- Highly penetrant variants for a number of drugs have been identified using pharmacogenomics.
- Findings are primarily limited to drugs used to treat cardiovascular diseases, infectious diseases, and cancer therapies.
- Existing pharmacogenomic studies are confounded by environmental factors, drug compliance, and pre-existing co-morbidities.
- They usually focus on drugs taken for a large amount of time, adverse events or dosing, not on acute response.
- Phenotypic heterogeneity and differences in underlying genetic ancestry are also confounders to current approaches.

Aims

- Measure real-time drug response using patient data collected during surgery where bias of drug type and dose and pre-existing conditions are known, environmental impact is minimized, and real-time physiological response is collected.
- Systematically describe differences between patients.
- Uncover potential genetic variants underlying real-time drug response variability.

Phenylephrine drug response

- α_1 -adrenergic receptor agonist \rightarrow known drug target.
- Phenylephrine is responsible for vasoconstriction of blood vessels thereby increasing blood pressure and is commonly administered during surgery.
- Evidence of inter-individual drug response.
- Genomics of phenylephrine drug response is unknown.

Materials and Methods

- Patients enrolled in *BioMe* (Mount Sinai biobank)
 - Who have undergone surgery
 - With available genotype
- Phenotype: difference in systolic blood pressure due to phenylephrine
- Significant confounders were either used as exclusion criteria or adjusted for:



Extreme systolic/diastolic, mean arterial pressures, heart rate

- Emergency procedures, endo-, colonoscopies, intubations, ...
- Patients w/ blood transfusions or who received > 500 ml fluids
- Patients who received propofol and phenylephrine within a 10 min interval
- - African Americans (n = 1217)
 - Hispanic/Latinos (n = 1707)
 - European Americans (n = 1386)
- Health System Diverse population: Biobank

drug

bolus drug drug drug bolus bolus bolus Perioperative Procedure time Period Procedure SBP ΔSBP before after Surgery patient Patient related Genomic data Ancestry Mount Sinai Age/Weight/Height/Sex Biomarkers **ASA** status Charlson score EHR Biobank Comorbidities: Diabetes Hypertension Peripheral Artery Disease Congestive heart failures

Procedure related

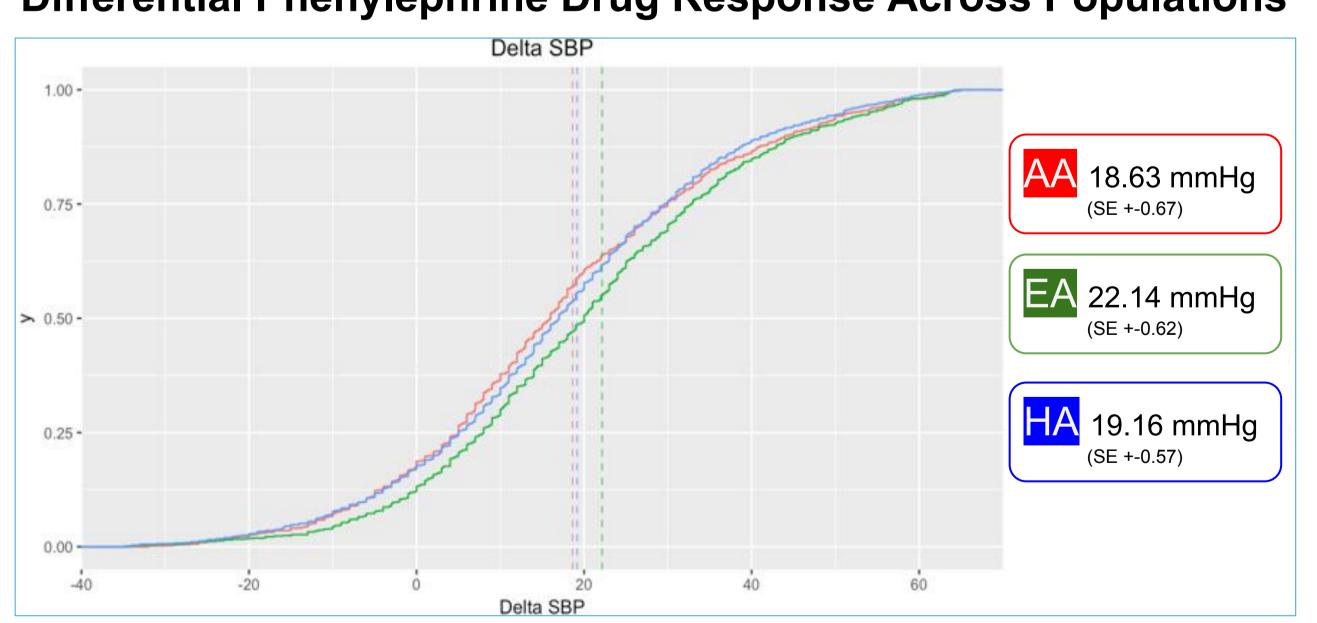
- anesthesia time
- anesthesia type
- procedure time number of boluses
- amount of bolus given
- doses of propofol, remifentanil,
- phenylephrine
- type of surgery (ICD9/class)
- fluids administered/lost inpatient/ambulatory
- systolic/diastolic BP
- heart rate
- etc...

Results

Differential Phenylephrine Drug Response Across Populations

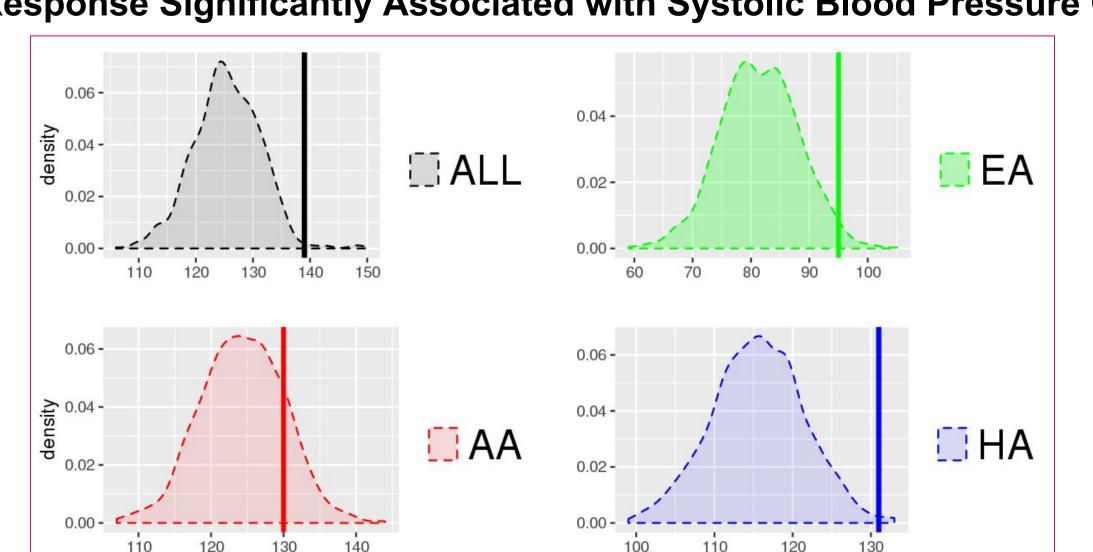
for > 40 million SNPs inputed from 1000G reference panels

Genome-wide association tests were performed



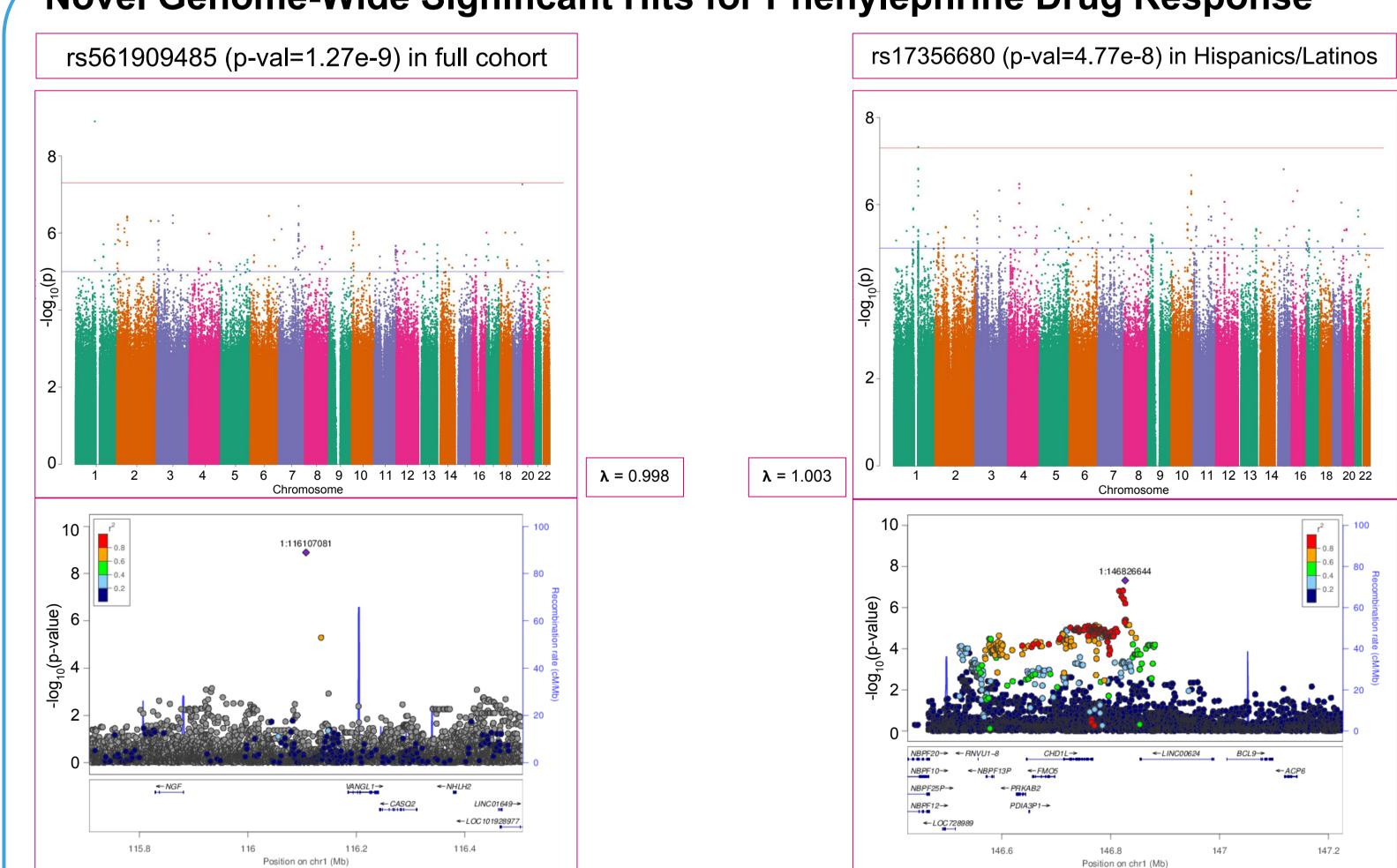
European Americans have a significantly higher increase of African after phenylephrine than Americans pressure and Hispanic/Latinos (p-values < 9e-6 and 10e-5)

Drug Response Significantly Associated with Systolic Blood Pressure Genes



165 genes associated with systolic blood pressure in the UK BioBank cohort are enriched for association with phenylephrine response in the BioMe cohort in European Americans (p-val = 0.016) and Hispanic Latinos (p-val = 0.005). Lower association in African Americans from our cohort is consistent with mostly European cohort from the UK BioBank.

Novel Genome-Wide Significant Hits for Phenylephrine Drug Response



GWAS hit downstream of VANGL1 (VANGL1 (+/-) mice show aberrant subclavian artery)

GWAS hit near CHD1L (highly expressed in arterial tissues); eQTLs w/ rs17356680 and CHD1L in skeletal muscles and arteries.

Conclusions

- We established a robust definition for real-time drug response that can be applied to successfully conduct large-scale pharmacogenomic studies.
- We identified inter-population differences of phenylephrine response.
- We discovered putative novel associations near genes CHD1L and VANGL1.
- Systolic blood pressure genes are enriched in association with drug response.