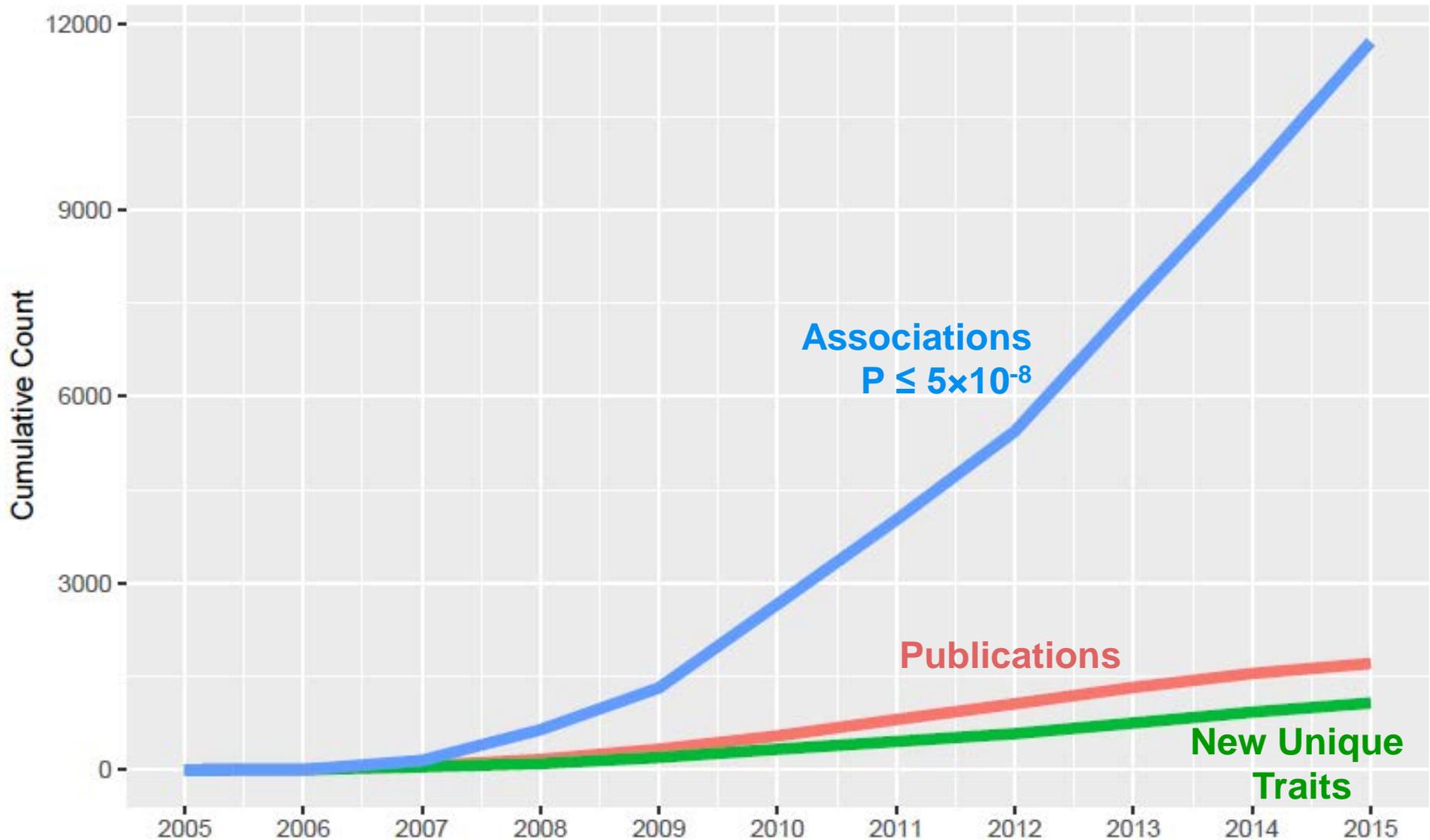


Potential Advantages and Pitfalls to Using Genetics in Drug Development for Complex Diseases

Matthew R. Nelson
March 8, 2017

Genome-wide association studies have identified thousands of variants that influence complex human traits



Early reasons to be optimistic about PGx in PM Abacavir hypersensitivity reaction (HSR) and *HLA-B*57:01*



In 2001, an analysis of a retrospective case/control study identified *HLA-B*57:01* to be strongly associated with HSR risk

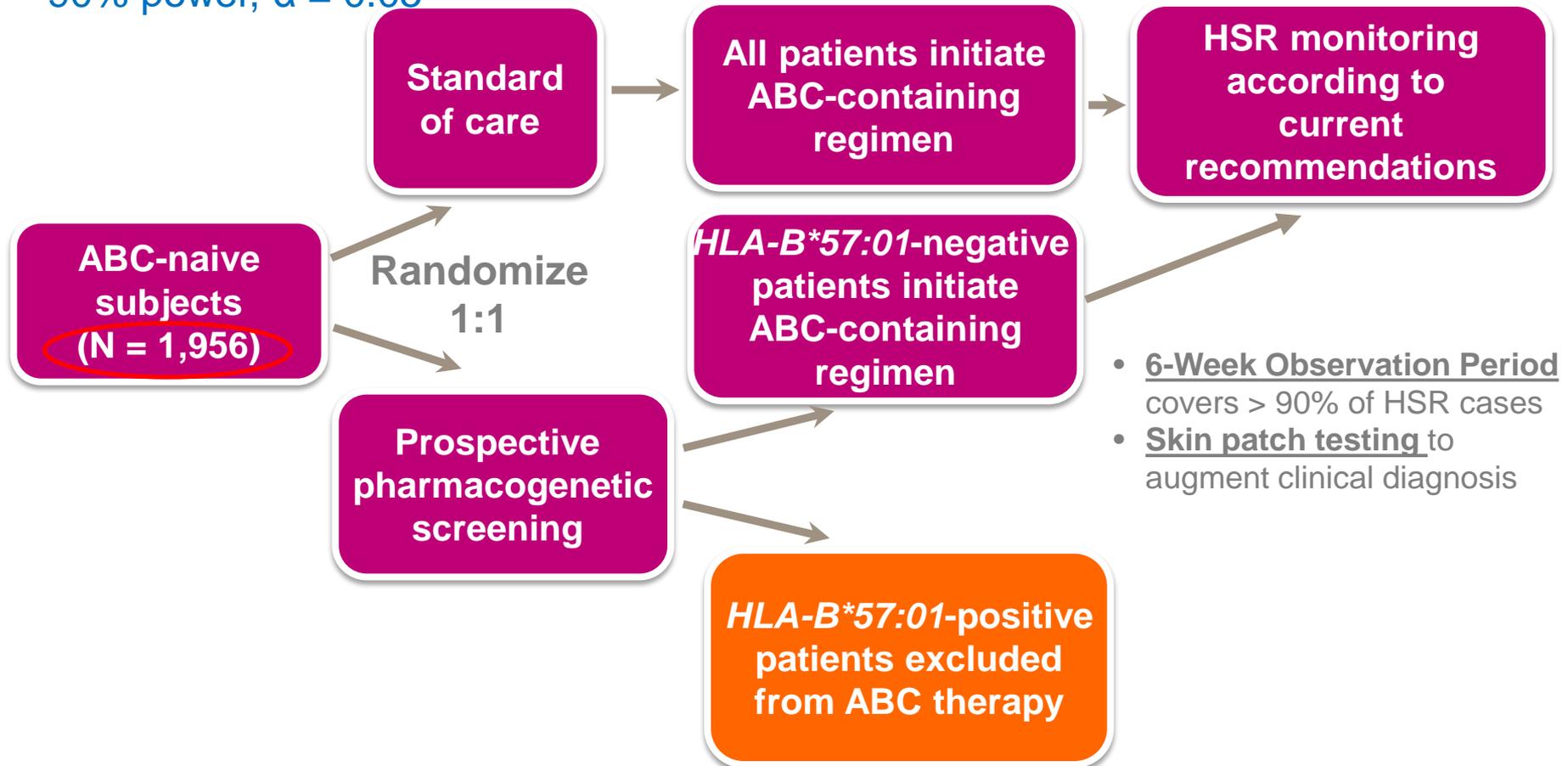
	Cases (n=84)	Controls (n=113)	p-value
<i>HLA-B57 Present</i>	39 (46%)	4 (4%)	<0.0001
<i>HLA-B57 Absent</i>	45 (54%)	109 (96%)	

The presence of HLA-B57 is more common in cases (46%) than controls (4%)

PREDICT-1 study design



Hypothesized HSR rate **7.3%**
Post-screening HSR rate **3.8%**
90% power, $\alpha = 0.05$



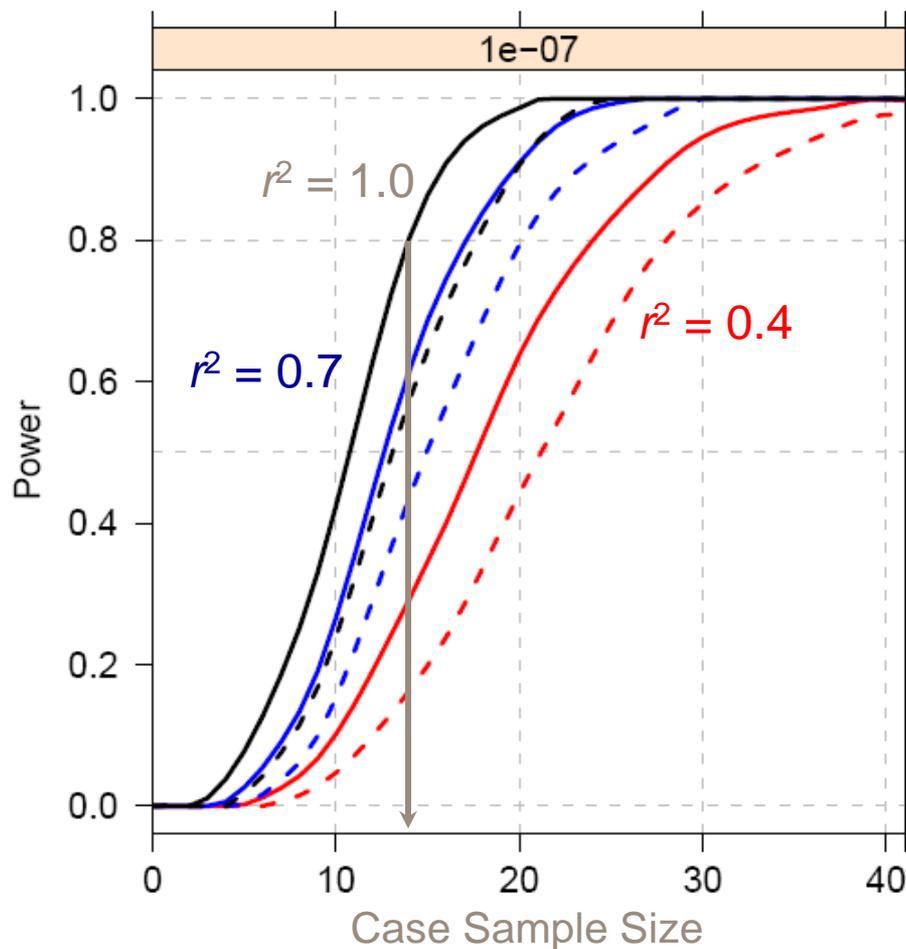
PREDICT-1 trial demonstrated screening utility



Table 4. Performance Characteristics of HLA-B*5701 Screening for Hypersensitivity Reaction to Abacavir in the Control Group.*

Subgroup	Characteristic	Performance Characteristic
	WARNING: HYPERSENSITIVITY REACTIONS, and LACTIC ACIDOSIS AND SEVERE HEPATOMEGALY	
	<i>See full prescribing information for complete boxed warning.</i>	
	<u>Hypersensitivity Reactions</u>	
Immunocompetent	<ul style="list-style-type: none"> • Serious and sometimes fatal hypersensitivity reactions have occurred with ZIAGEN (abacavir). (5.1) • Hypersensitivity to ZIAGEN is a multi-organ clinical syndrome. (5.1) • Patients who carry the HLA-B*5701 allele are at a higher risk of experiencing a hypersensitivity reaction to abacavir. (5.1) • ZIAGEN is contraindicated in patients with a prior hypersensitivity reaction to abacavir and in HLA-B*5701-positive patients. (4) • Discontinue ZIAGEN as soon as a hypersensitivity reaction is suspected. Regardless of HLA-B*5701 status, permanently discontinue ZIAGEN if hypersensitivity cannot be ruled out, even when other diagnoses are possible. (5.1) • Following a hypersensitivity reaction to ZIAGEN, NEVER restart ZIAGEN or any other abacavir-containing product. (5.1) 	
Total patients		100)
Hypersensitivity		100)
No hypersensitivity		-98.0)
White skin		
Hypersensitivity		100)
No hypersensitivity		-97.6)
		PPV: 46.8 (32.1–61.9) NPV: 100 (99.4–100)

Power to Detect Abacavir HSR HLA-B*5701-like Effect



Adjustment for 500,000 hypothesis tests for an experiment-wide $\alpha = 0.05$ yields a test-wise significance level of 10^{-7}

- 200 clinical controls
- - - 200 population controls

Non-iSAEC Findings			iSAEC-related Findings		
Drug	Risk Allele	OR	Drug	Risk Allele	OR
Liver Injury					
Pazopanib	<i>HLA-B*57:01</i>	2	Augmentin	<i>HLA-DRB1*15:01</i>	3
Nevirapine	<i>HLA-DRB1*01</i>	3	Augmentin	<i>HLA-A*02:01</i>	2
Ximelagatran	<i>HLA-DRB1*07:01</i>	4	Augmentin	<i>PTPN22 R620W</i>	2
Lapatinib	<i>HLA-DRB1*07:01</i>	19	Flucloxacillin	<i>HLA-B*57:01</i>	81
Lumiracoxib	<i>HLA-DRB1*15:01</i>	13	Methyldopa	<i>HLA-A*33:01</i>	98
Ticlopidine	<i>HLA-A*33:03</i>	36	Fenofibrate	<i>HLA-A*33:01</i>	51
			Terbinafine	<i>HLA-A*33:01</i>	37
			Sertraline	<i>HLA-A*33:01</i>	22
			Erythromycin	<i>HLA-A*33:01</i>	10
			Cyprofloxacin	<i>HLA-B*50:02</i>	11
			Minocyclin	<i>HLA-B*35:02</i>	28
			Valproic Acid	<i>HLA-DRB1*10:01</i>	14
			Atorvastatin	<i>HLA-DRB1*10:01</i>	11
			Simvastatin	<i>HLA-B*13:02</i>	8
Skin Injury					
Nevirapine	<i>HLA-B*35</i>	2	Carbamazepine	<i>HLA-A*31:01</i>	26
Nevirapine	<i>HLA-C*04</i>	3			
Phenytoin	<i>CYP2C9*3</i>	12			
Phenytoin	<i>HLA-B*15:02</i>	6			
Allopurinol	<i>HLA-B*58:01</i>	678			
Carbamazepine	<i>HLA-B*15:02</i>	>1000			
AHSS/HSR					
Dapsone	<i>HLA-B*13:01</i>	21	Penicillins	<i>HLA-DRB1*10:01</i>	3
Abacavir	<i>HLA-B*57:01</i>	>1000	Augmentin	<i>HLA-DRB1*10:01</i>	3
			Amoxicillin	<i>HLA-DRB1*10:01</i>	3
Other					
Statin myopathy	<i>SLCO1B1*5</i>	5	Thiopurine pancreatitis	<i>HLA-DRB1*07:01</i>	3
Anthracycline cardiotox	<i>SLC28A3 rs7853758</i>	5	5-ASA nephrotoxicity	<i>HLA-DRB1*03:01</i>	2
Anthracycline cardiotox	<i>UGT1A6 rs6759892</i>	3	Clozapine agranulocytosis	<i>HLA-DQB1 126H</i>	5
Pazopanib hyperbilirubin	<i>UGT1A1*28</i>	13	Clozapine agranulocytosis	<i>HLA-B 158T</i>	3
Cisplatin ototoxicity	<i>TPMT rs12201199</i>	17			
Cisplatin ototoxicity	<i>COMT rs9332377</i>	5			
Thiopurine leukopenia	<i>NUDT15 Arg139Cys</i>	35			
Irinotecan neutropenia	<i>UGT1A1*28</i>	28			
Tranilast hyperbilirubin	<i>UGT1A1*28</i>	48			
Clopidogrel CV events	<i>CYP2C19*2/3/4/5</i>	3			



Gene Categories
HLA
ADME
Immune-related

Pharmacogenetics timeline at GSK

Several notable successes, but where's the impact on efficacy?




Collected and hold blood for later use

~\$10



DNA extracted real time and banked -

~\$50 DNA extraction & banking

DNA extracted as required

3wks to ship and extract

~\$40 per extraction



Custom genotyping

~\$1-5 per SNP (Taqman, Sanger Seq)

~\$500 GWAS (600K), ~\$1000 HLA

~\$400 Roche Amplichip (CYP2D6)

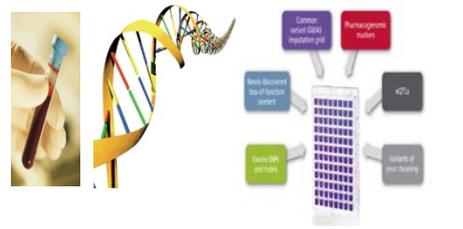
Bespoke GT

~\$30 candidate SNPs

~\$400 GWAS (1M)

~\$350 ADME arrays

~\$700 HLA

Genome	\$1000-3000
Exome	\$600-1,500
NGS Panels	\$200-400
GWAS	\$80-100
HLA	\$400
Imputation	\$0

Many Candidate Gene Studies Published Implicating Genes in Steroid Treatment Response in Asthma Subjects

Association between *WDR21A* polymorphisms and airway responsiveness to inhaled corticosteroids in asthmatic patients

Sung-Hwan Cho^{a,b}, Byung-Lae Park^c, Seung Woo Shin^{a,b}, Jeong-Seok Heo^{a,b}, Jong-Sook Park^{a,b}, Sung Woo Park^{a,b}, An-Soo Jang^{a,b}, Il Yup Chung^e, Hounng-Doo Shin^{c,d} and Choon-Sik Park^{a,b,e}

Objective Genetic polymorphism is partially responsible for the wide variation in the response of moderate-to-severe asthmatic patients to inhaled

of genotype. The predicted *WDR21A* protein structure was similar to the Gji, protein structure (template modeling-score=0.93).

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Genomewide Association between *GLCCI1* and Response to Glucocorticoid Therapy in Asthma

Mechanisms of allergy and clinical immunology

The glucocorticoid receptor heterocomplex gene *STIP1* is associated with improved lung function in asthmatic subjects treated with inhaled corticosteroids

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Human Molecular Genetics, 2004, Vol. 13, No. 13 1353–1359
DOI: 10.1093/hmg/ddh149
Advance Access published on May 5, 2004

Corticosteroid pharmacogenetics: association of sequence variants in *CRHR1* with improved lung function in asthmatics treated with inhaled corticosteroids

Kelan G. Tantisira^{1,2}, Stephen Lake¹, Eric S. Silverman², Lyle J. Palmer³, Ross Lazarus¹, Edwin K. Silverman^{1,2}, Stephen B. Liggett⁴, Erwin W. Gelfand⁵, Lanny J. Rosenwasser⁵, Brent Richter¹, Elliot Israel², Michael Wechsler^{1,2}, Stacey Gabriel^{6,8}, David Altshuler^{6,7,8}, Eric Lander^{6,8}, Jeffrey Drazen² and Scott T. Weiss^{1,*}

TBX21: A functional variant predicts improved response to inhaled corticosteroids in asthma with the use of inhaled corticosteroids

Kelan G. Tantisira^{***}, Eun Sook Hwang⁵, Benjamin A. Raby^{**}, Eric S. Silverman^{†5}, Stephen L. Lake^{*}, Brent G. Richter^{*}, Stanford L. Peng[‡], Jeffrey M. Drazen^{†5}, Laurie H. Glimcher⁵, and Scott T. Weiss^{*}

^{*}Channing Laboratory and [†]Pulmonary Division, Brigham and Women's Hospital and Harvard Medical School, Boston, MA 02115; and [‡]Washington University School of Medicine, St. Louis, MO 63110

Contributed by Laurie H. Glimcher, November 17, 2004

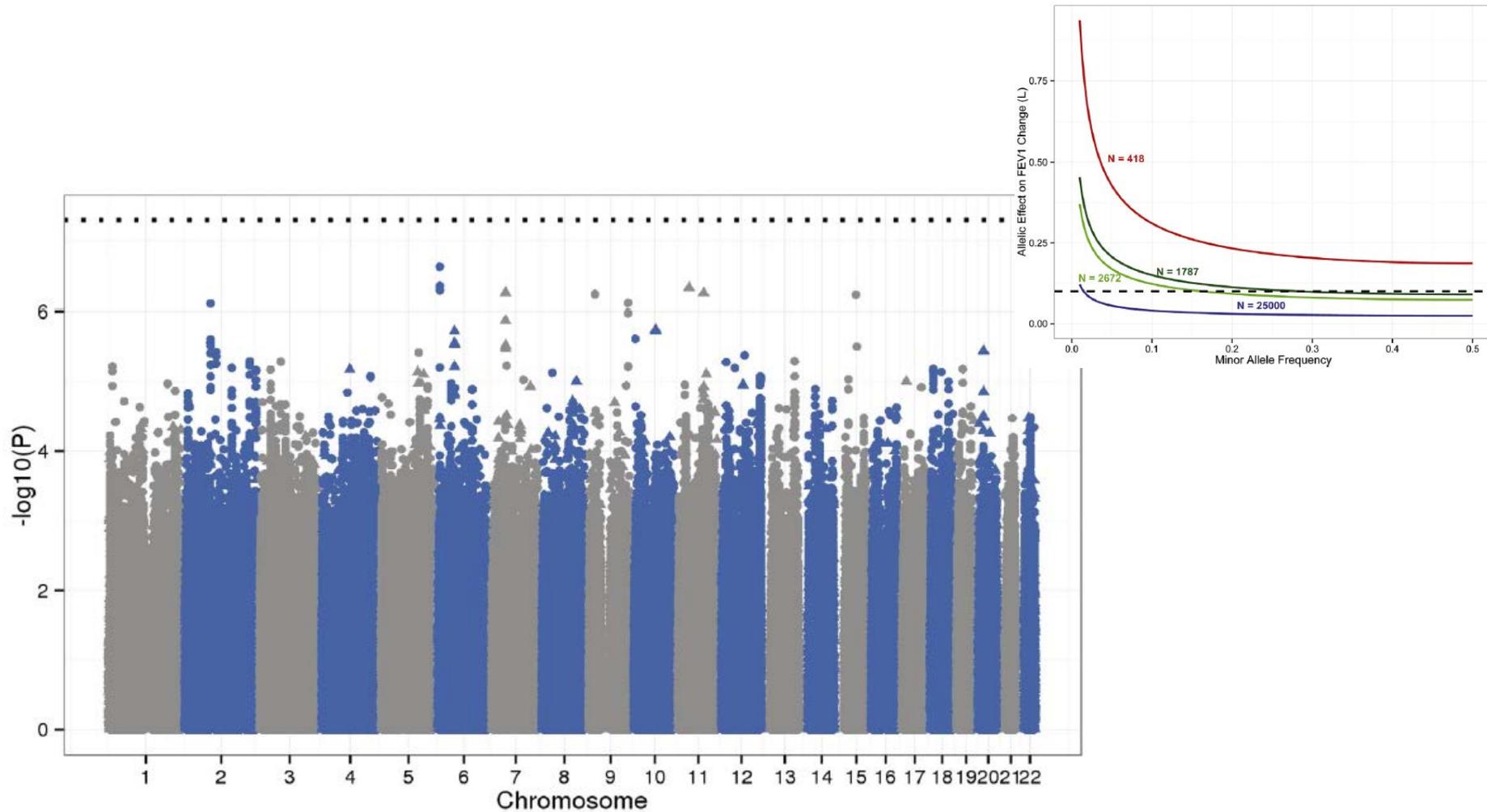
TBX21 encodes for the transcription factor T-bet (T-box expressed in T cells) and is expressed in T cells

Asthma and lower airway disease

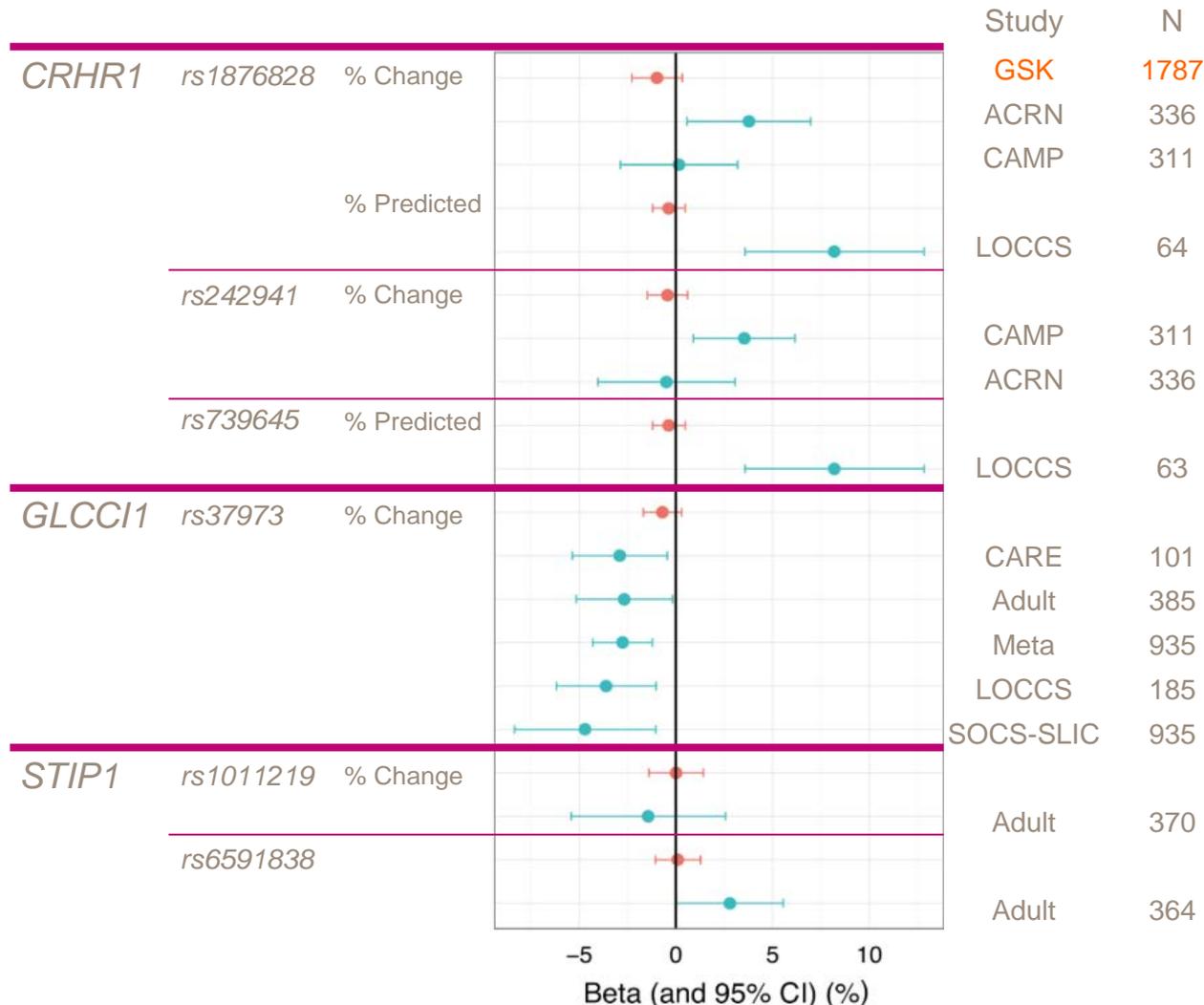
Genetic predictors associated with improvement of asthma symptoms in response to inhaled corticosteroids

Heung-Woo Park, MD, PhD,^{a,b} Amber Dahlin, PhD, MMSc,^a Szeman Tse, MDCM, MPH,^a Qing Ling Duan, PhD,^a Brooke Schuemann, BS,^a Fernando D. Martinez, MD,^a Stephen P. Peters, MD, PhD,^d Stanley J. Szefler, MD,^a John J. Lima, PharmD,^f Michiaki Kubo, MD, PhD,^g Mayumi Tamari, MD, PhD,^g and Kelan G. Tantisira, MD, MPH^{a,b}
Boston, Mass, Seoul, Korea, Tucson, Ariz, Winston-Salem, NC, Denver, Colo, Jacksonville, Fla, and Kanagawa, Japan

Study of Patient Response to Inhaled Corticosteroids across 7 Clinical Trials in 2672 patients (1787 European)



No Previously Reported Associations with Steroid Response were Statistically Significant at $P < 0.05$



Several GWAS for drug efficacy have been conducted and many high confidence associations found



2,223 GWAS reported in NHGRI/EBI GWAS Catalog as of July, 2015

76 (3.4%) GWAS of drug efficacy

- 41 different drugs, drug classes or combinations

13 (17%) GWAS of drug efficacy with high confidence associations

- 6 of 41 (15%) different drugs, drug classes or combinations

13 high confidence efficacy associations for five drug classes identified by GWAS



Drug	Efficacy Trait	Gene (Allele)
Warfarin	Maintenance dose	<i>VKORC1</i> <i>CYP2C9</i> (*3, *2, other) <i>CYP4F2</i>
Clopidogrel	Platelet aggregation, CV events	<i>CYP2C19</i> (*2)
Rosuvastatin	Change in LDL-C	<i>ABCG2</i>
Statins	Change in LDL-C	<i>LPA</i> <i>APOE</i> <i>SORT1</i> <i>SLCO1B1</i>
Metformin	HbA1c below 7%	<i>ATM</i>
Peg-Interferon	Sustained virologic response	<i>IL28B</i>



Compelling candidate gene drug efficacy associations



11 additional associations for 7 drugs not identified or confirmed by GWAS

Drug	Efficacy Trait	Gene (Allele)
Codeine	Pain tolerance	<i>CYP2D6</i> (PM)
Tamoxifen	Disease-free survival (inv)	<i>CYP2D6</i> (PM)
Eculizumab	Hemolytic activity	<i>C5</i>
EGFR TKIs	NSCLC progression	<i>BIM</i>
Platinum-based chemotherapy	Ovarian cancer progression and survival	<i>BRCA1</i> <i>BRCA2</i>
Olaparib	Ovarian cancer complete response	<i>BRCA1</i> <i>BRCA2</i>
Sulfonylureas	HbA1c/glucose control	<i>KCNJ11</i> <i>ABCC8</i> <i>HNF1A</i>



Efficacy associations fall into three functional mechanisms

Though with some ambiguity based on current understanding

ADME <200 genes

Warfarin–*CYP2C9*
Warfarin–*CYP4F2*
Clopidogrel–*CYP2C19*
Rosuvastatin–*ABCG2*
Statin–*SLCO1B1*
Codeine–*CYP2D6*
Tamoxifen–*CYP2D6*

Drug Target ~350 genes

Warfarin–*VKORC1*
Eculizumab–*C5*
Sulfonylureas–*KCNJ11*
Sulfonylureas–*ABCC8*

Disease Mechanism >>2000 genes

Statin–*LPA*
Statin–*APOE*
Statin–*SORT1*
Metformin–*ATM*
Peg Ifn–*IL28B*
Platinums–*BRCA1/2*
Olaparib–*BRCA1/2*
EGFR TKIs–*BIM*
Sulfonylureas–*HNF1A*

Significant excess of ADME gene associations (OR = 5.5) and underrepresentation of disease gene associations (OR = 0.24; $p = 0.001$)

What have we learned about the role of genetics in drug safety and efficacy?



- **Drug Safety**

- Mechanisms primarily through HLA and ADME genes
- Most HLA genetic risk factors are drug and SAE specific
- Common genetic risk factors (*alone*) do not accurately predict patient risk for rare SAEs
- <50% of drug SAEs appear to have *major* genetic risks

- **Drug Efficacy**

- Mechanisms primarily through ADME, drug targets, and disease genes
 - We expect ~10% of drugs to have genetic predictors of drug efficacy that could influence clinical decision making
-

- Probability of success can be improved with tailored analysis strategies
 - Where possible, take advantage of sensitive, quantitative biomarkers of response
 - Balance hypothesis-driven and hypothesis-free approaches
 - Combine data across trials of same or similar drugs
 - Don't shy away from investigating very effective drugs; this is where the best opportunities are
- With proper consent or secondary use, anonymized clinical trial data could advance understanding of disease genetics

Advantages

- Cost effective strategies to consent and screen routinely
- Discover early, *hopefully* validate pre-submission
- Opportunity to develop companion diagnostics
- Quickly respond to regulatory and academic questions
- Use unique clinical resources to advance discovery and development

Pitfalls

- Limitations in statistical power and strong temptation to over interpret “suggestive” or just significant findings
 - Tendency to focus on trials demonstrating poor efficacy
 - Lowest probability of success
 - Low “success” rates require a long view
 - Cost
-

iSAEC Acknowledgments



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- Scott Patterson (Amgen)
- Martin Armstrong & Fredrik Nyberg (Astra-Zeneca)
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- Michael Dunn (Wellcome Trust)

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- EUDRAGENE -- Mariam Molokhia (Kings College, UK)
- ITCH -- Munir Pirmohamed (Liverpool, UK)
- TdP -- Elijah Behr (SGUL) & Dan Roden (Vanderbilt)
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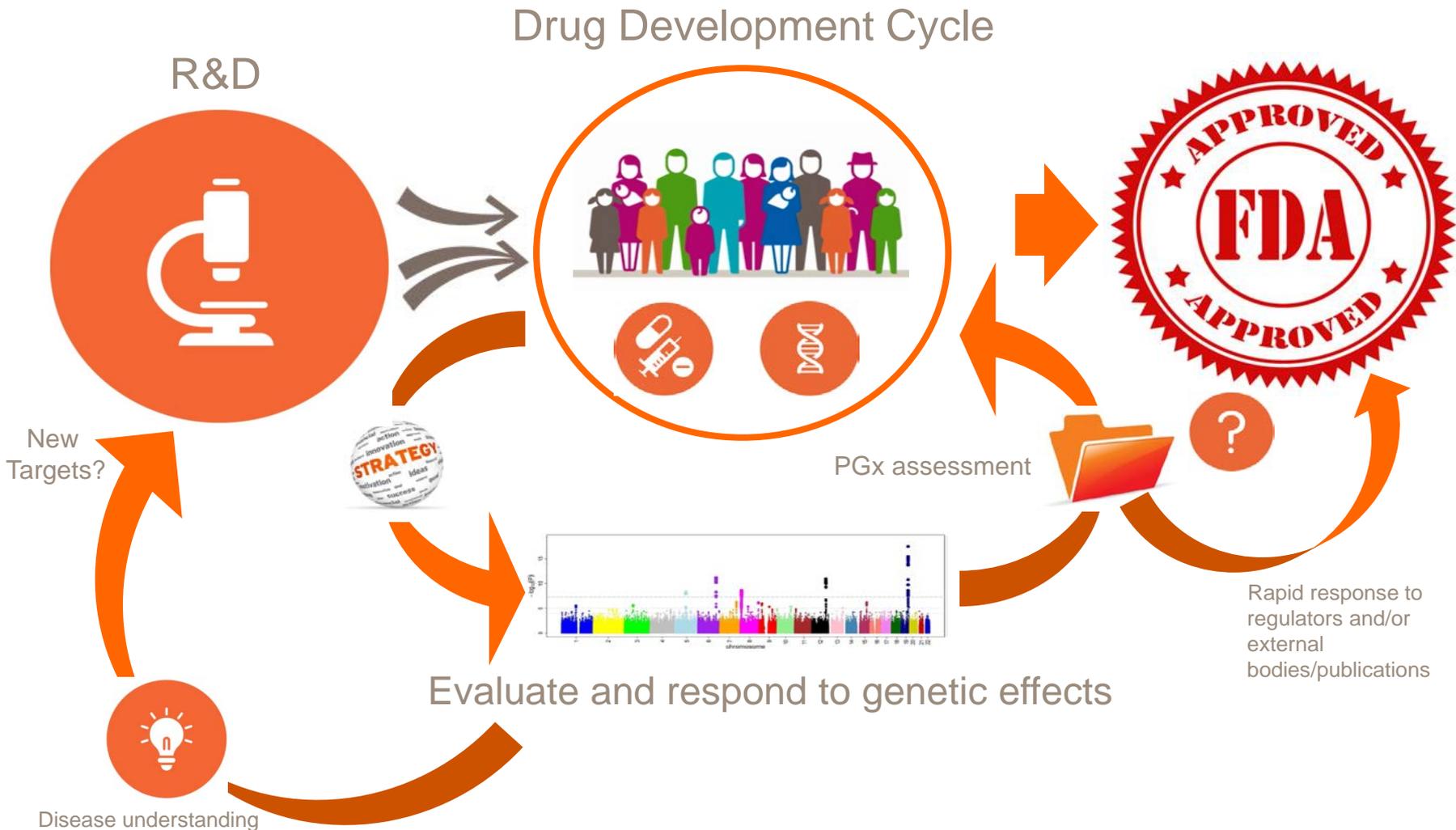
GSK

- **Toby Johnson**
- Liling Warren (currently independent)
- Arlene R. Hughes (currently at PAREXEL)
- Charlie Cox
- Stephanie L. Chissoe
- Chun-Fang Xu
- **Dawn M. Waterworth**

Our PGx vision: To identify genetic predictors of response and safety in time to inform medicine development decision making



Driven by asset specific strategies, targeting key impact points



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^{*}Channing Laboratory and [†]Pulmonary Division, Brigham and Women's Hospital and Harvard Medical School, Boston, MA 02115; and [‡]Washington University School of Medicine, St. Louis, MO 63110

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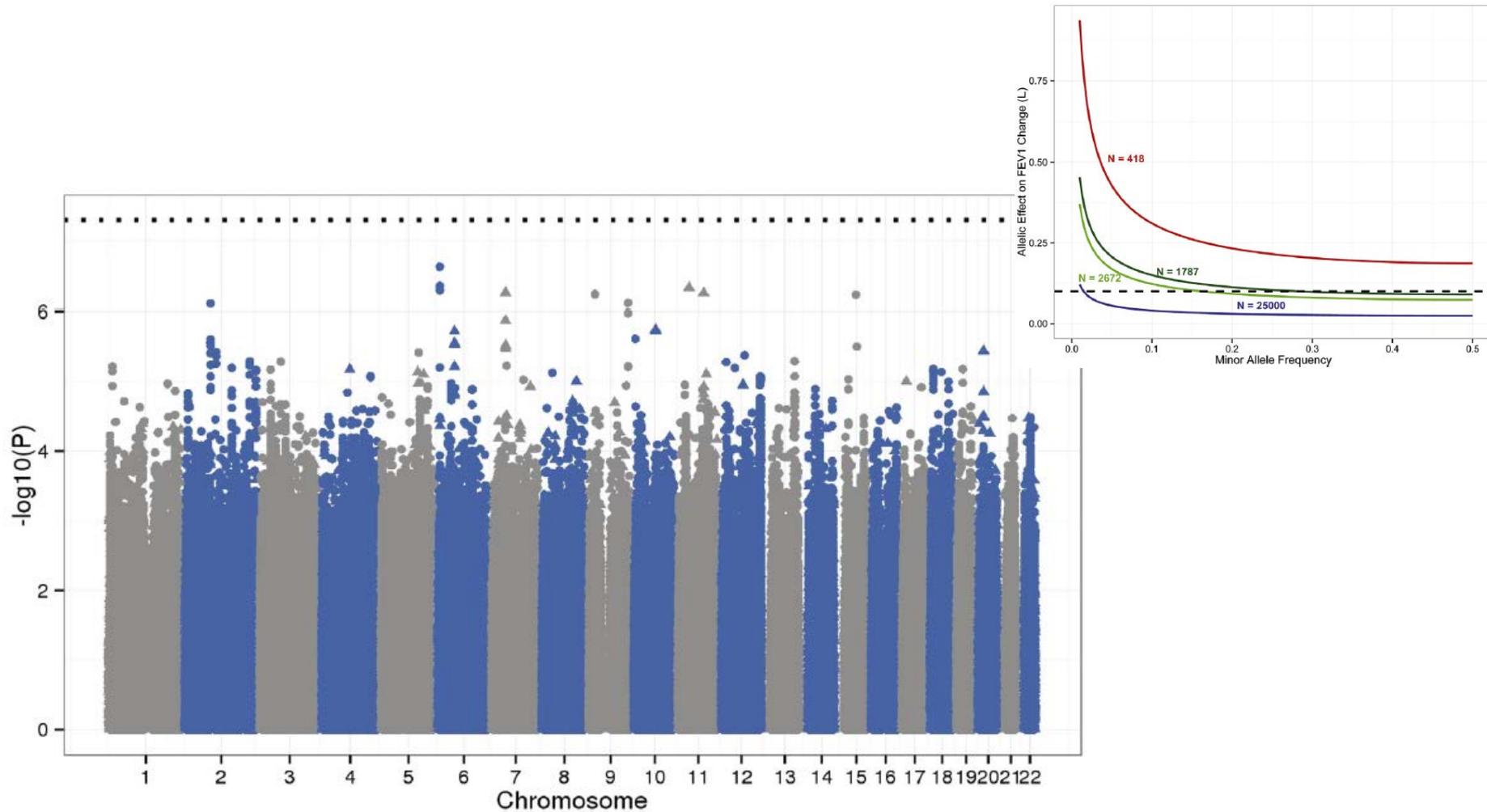
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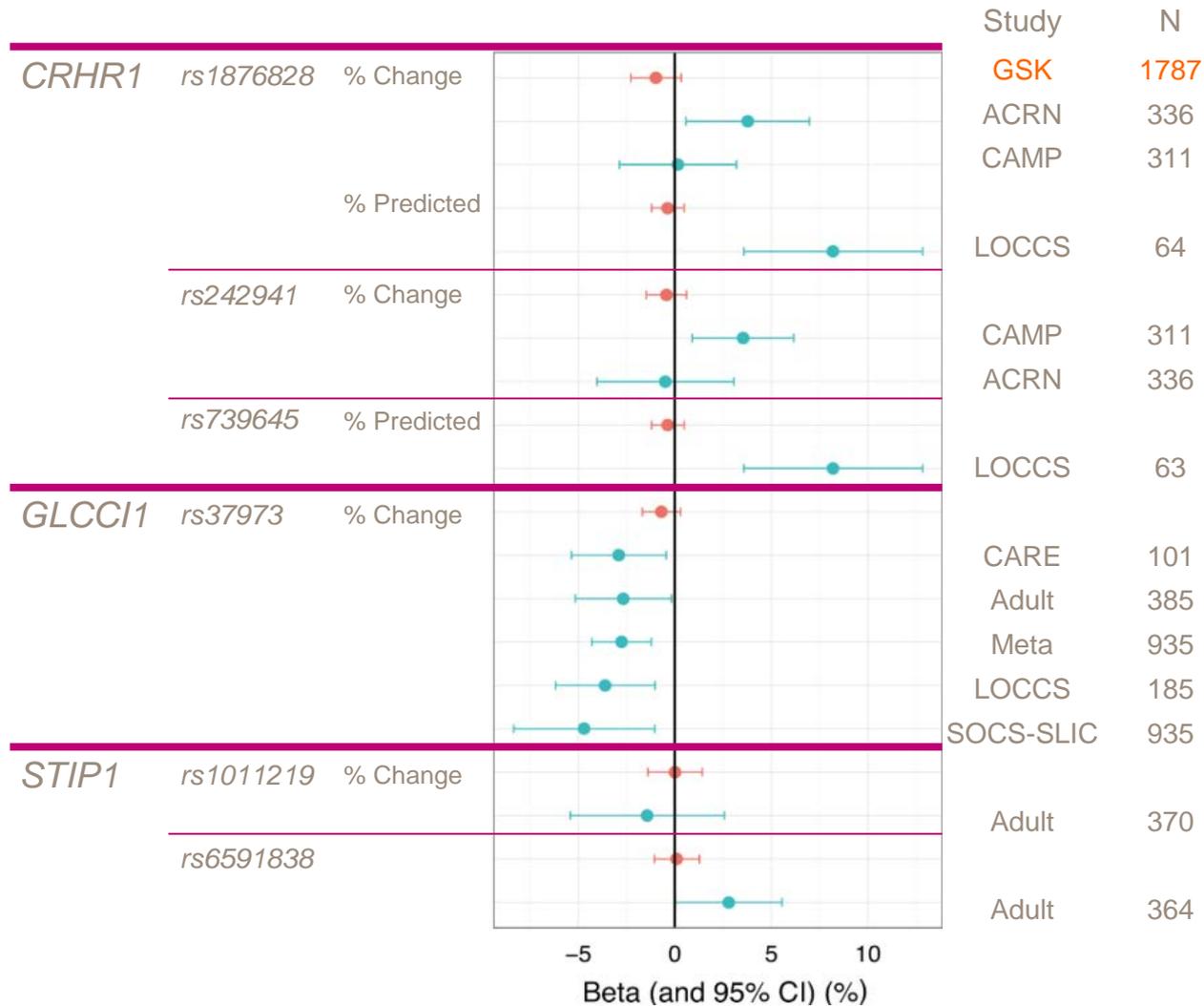
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PNAS

Study of Patient Response to Inhaled Corticosteroids across 7 Clinical Trials in 2672 patients (1787 European)



No Previously Reported Associations with Steroid Response were Statistically Significant at $P < 0.05$



How often do we expect drugs to have efficacy genetic effects with potential to inform patient treatment?



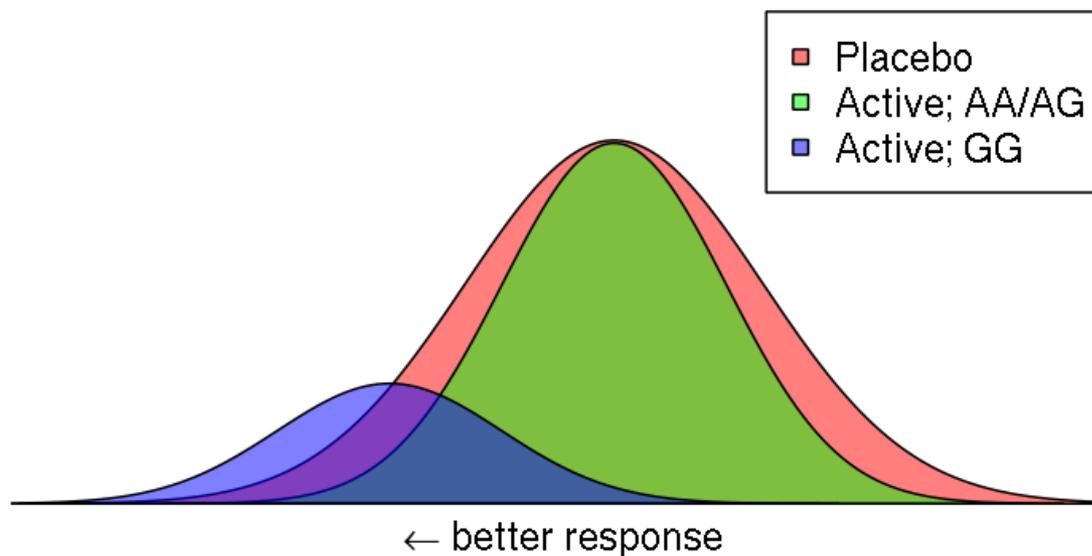
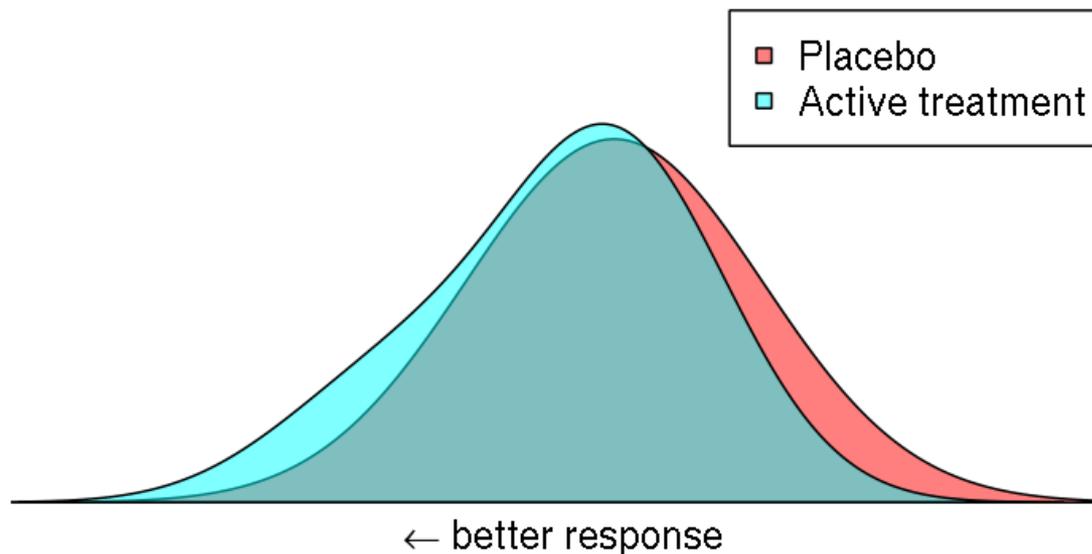
$$15\% \quad \times \quad 64\% \quad = \quad 8.1\% \quad (2.0\text{--}17.1\%)$$

6/41 drugs/classes
with effect found by
GWAS

5/9 drugs/classes
with common
genetic effects with
potential clinical
utility

Expected drugs having
genetic effects with
potential clinical utility

Efficacy effects, and the power to identify them, are constrained by drug effect sizes observed in trials

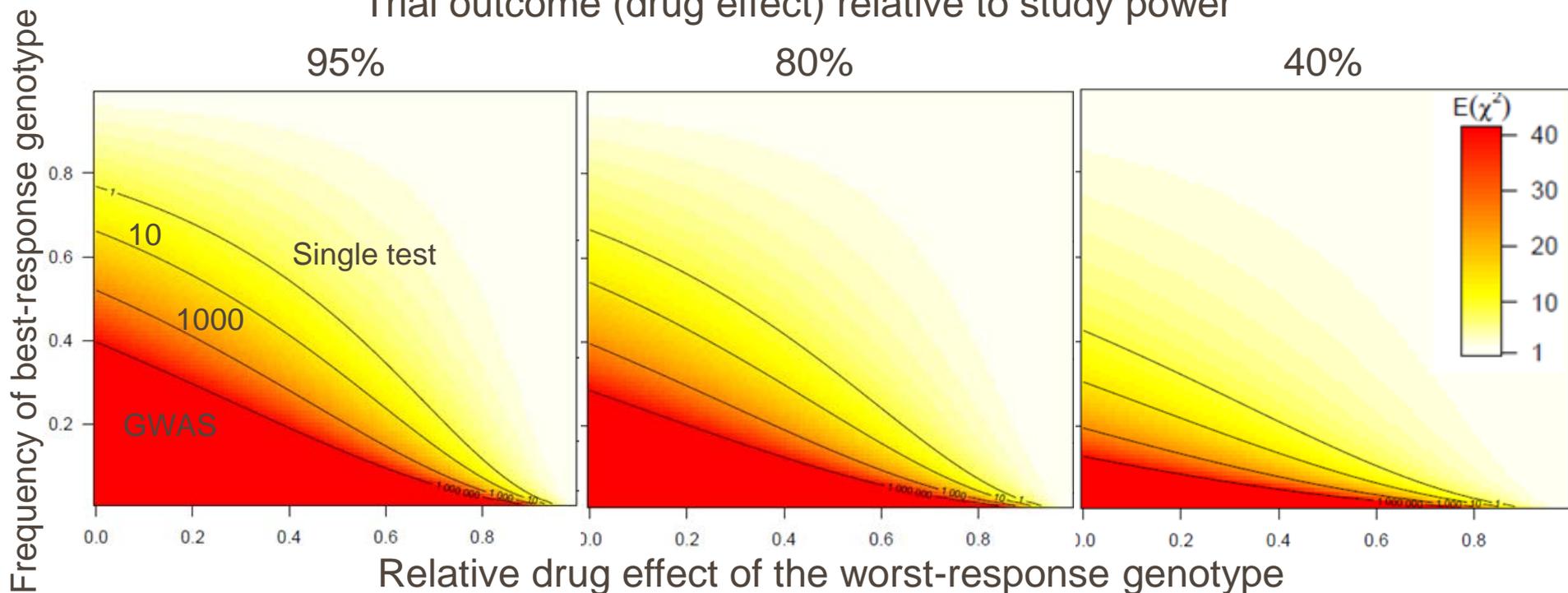


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$$\text{NCP}_{\text{PGx}} \leq \text{NCP}_{\text{treatment}} \times 2 \frac{1 - p_{\text{carrier}}}{p_{\text{carrier}}}$$

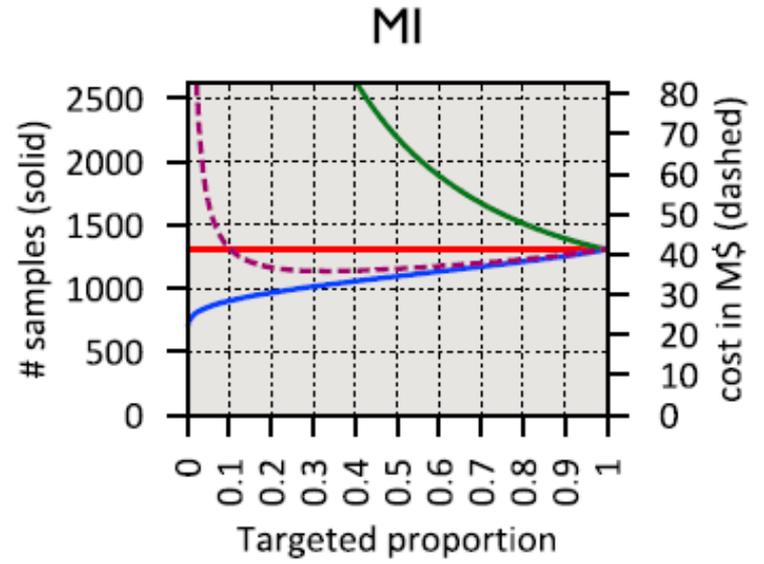
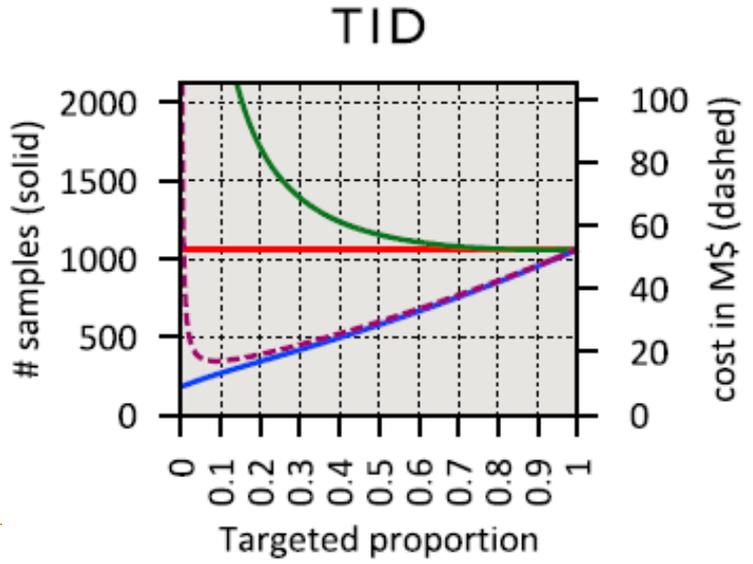
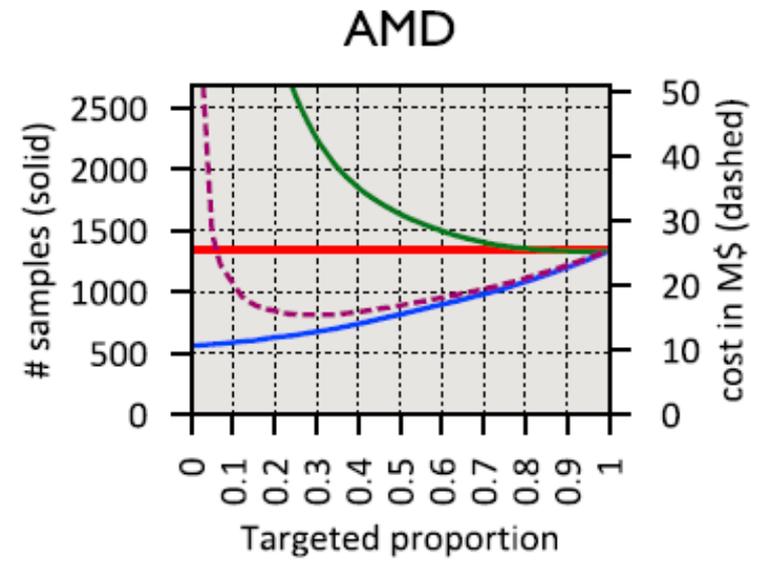
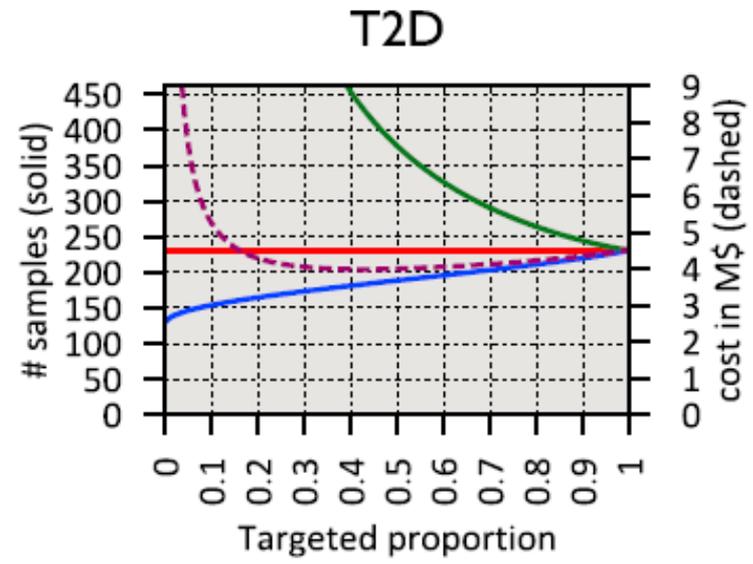
Trial outcome (drug effect) relative to study power



1. We should be mindful how we pay our multiple test penalties
2. Very few situations where PGx will rescue a failed trial
3. Trials with larger drug effects have more power to find efficacy PGx
4. Combine data across trials to increase PGx power

Enrichment by genetic risk can result in more efficient preventive clinical trials

randomized - conventional trial (red solid line)
 # randomized - enrichment trial (blue solid line)
 # screened - enrichment trial (green solid line)
 Total conventional trial cost (red dashed line)
 Total enrichment trial cost (purple dashed line)



Drug Safety

- Important role for genetics in immune-mediated and exposure-related adverse events
- 5 key discoveries within GSK
- External discoveries for 2 GSK drugs

Drug Exposure

- Expectations from regulators that PGx studies will be conducted to understand PK variability
- Avoidance of key polymorphic enzymes have reduced risk

Drug Efficacy*

- 5-20% of drugs estimated to have clinically meaningful PGx effects
- Known mechanisms focus on ADME, drug targets, and disease genes
- Larger effects can be

Be patient focused



*"The genetics of drug efficacy: opportunities and challenges": Nelson, M. et al., Nat Rev Genet (2016) 17: 197