

# Pharmacogenetics



Martin Wilkins

Imperial College London

Hammersmith Hospital

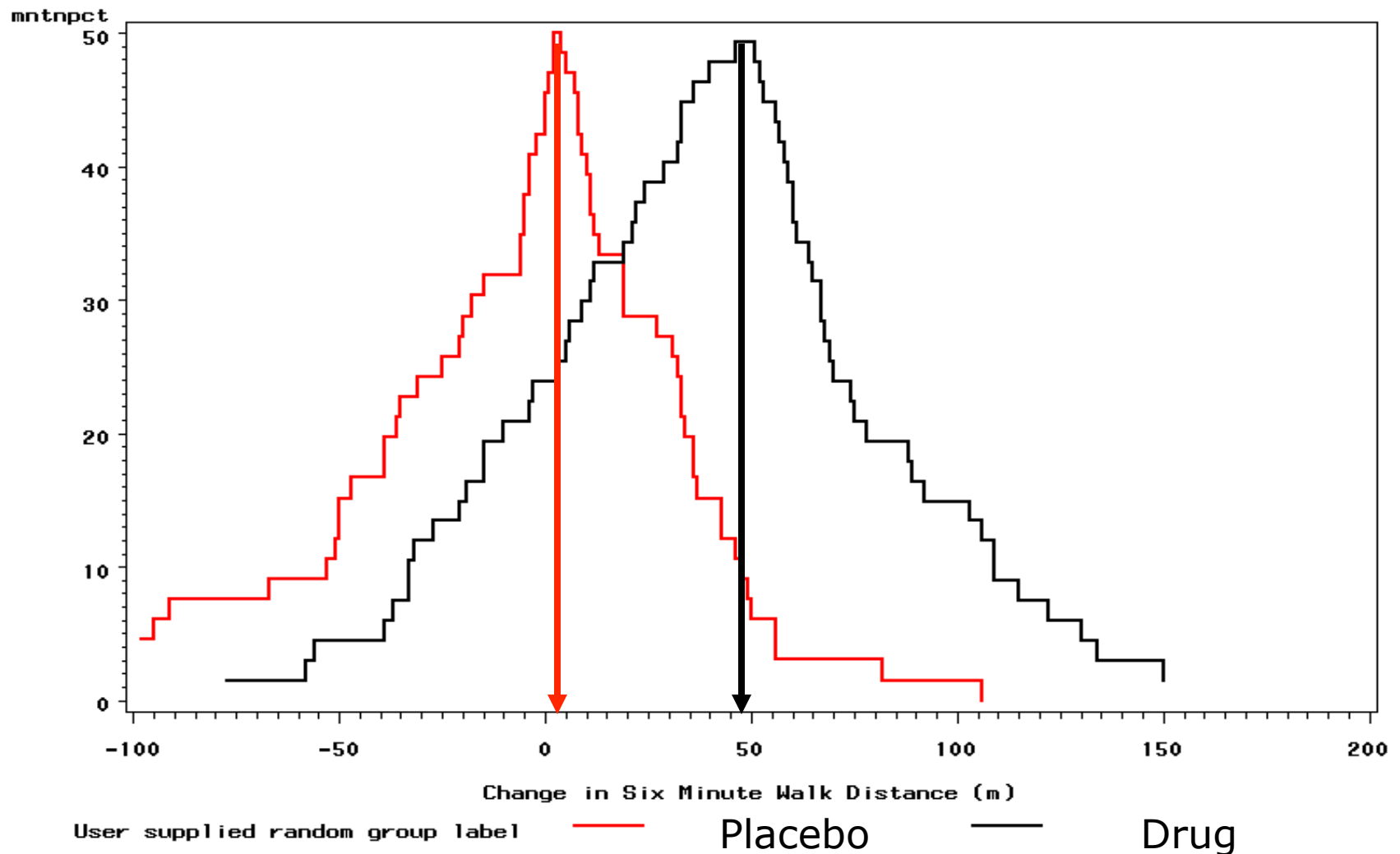
# Nomenclature

---

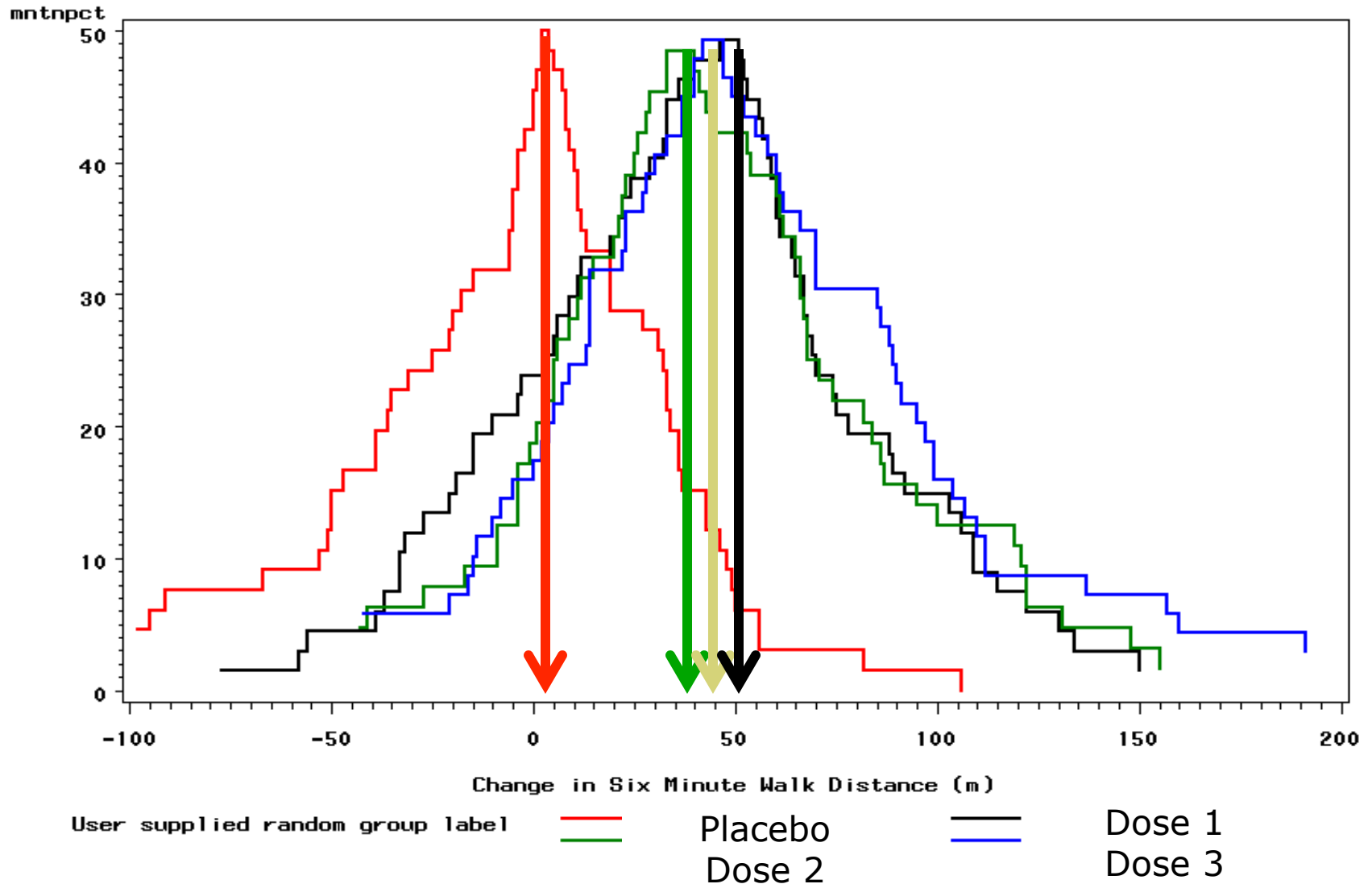
Pharmacogenomics is  
Pharmacogenetics

..with two single nucleotide polymorphisms

# Variability in response to a PAH drug



# Variability in response to a PAH drug





# Response rate to drugs for different diseases

---

Anti-depressant drugs

38%



Asthma drugs

40%



Diabetes drugs

43%



Arthritis drugs

50%



Alzheimer's drugs

73%



Cancer drugs

75%

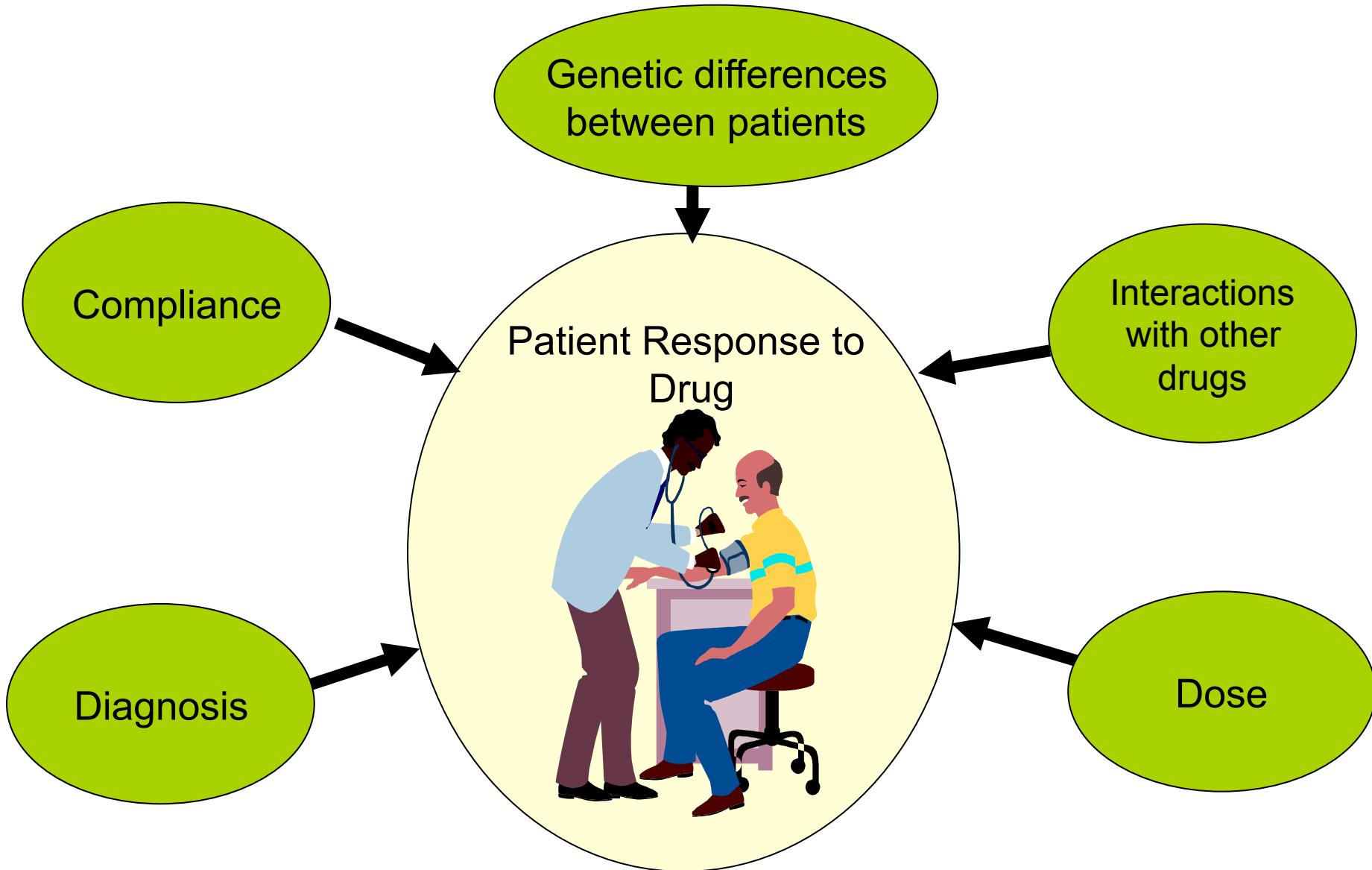


**Glaxo Chief: Our Drugs Do Not  
Work on Most Patients  
by Steve Connor**

A senior executive with Britain's biggest drugs company has admitted that most prescription medicines do not work on most people who take them.

*Published on Monday, December 8,  
2003 by the Independent/UK*

# Factors That Influence Medicine Response



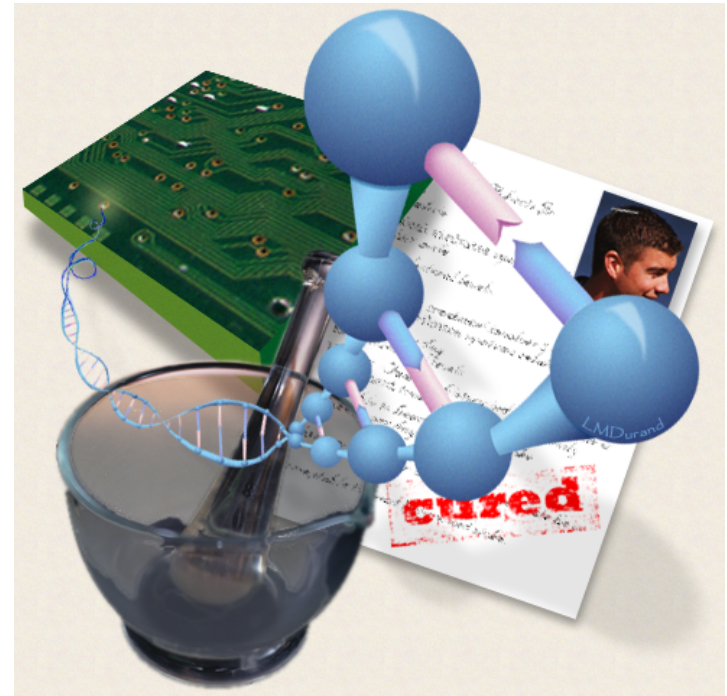
# Pharmacogenomics

## “Drugs by Design?”

---

“In the very near future, primary care physicians will routinely perform genetic tests before writing a prescription because (they will) want to identify the poor responders.”

F. Collins  
(AAFP Annual Meeting,  
1998)



# Objectives

---

1. To understand the principles underlying the use of genetic information to inform prescribing
2. To appreciate the present state-of-the-art and be aware of the hurdles to implementing pharmacogenetics/genomics in clinical practice

# Historical Background

---

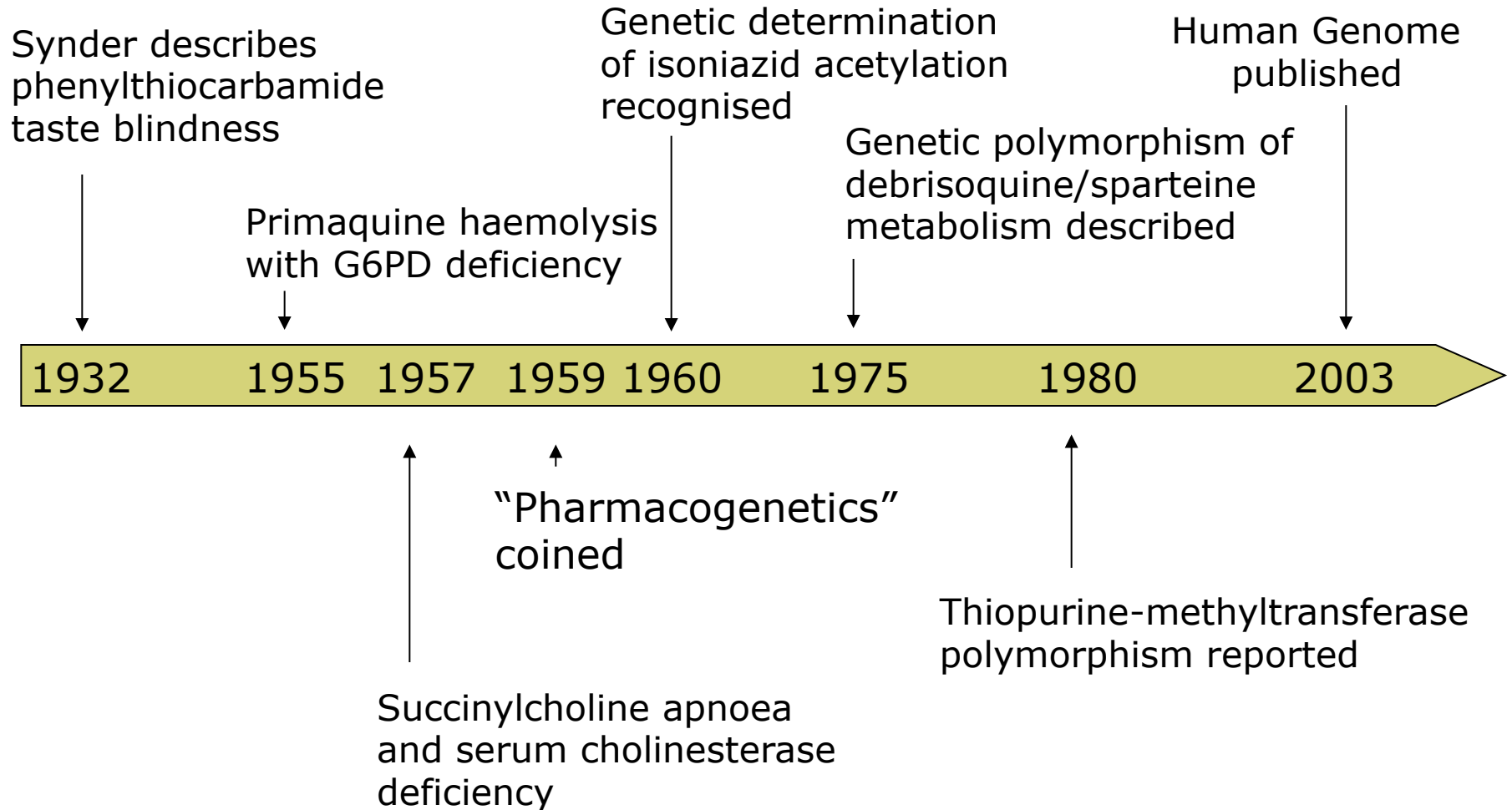


Sir Archibald Garrod 1858-1936

In his 1908 Croonian Lecture he coined the phrase

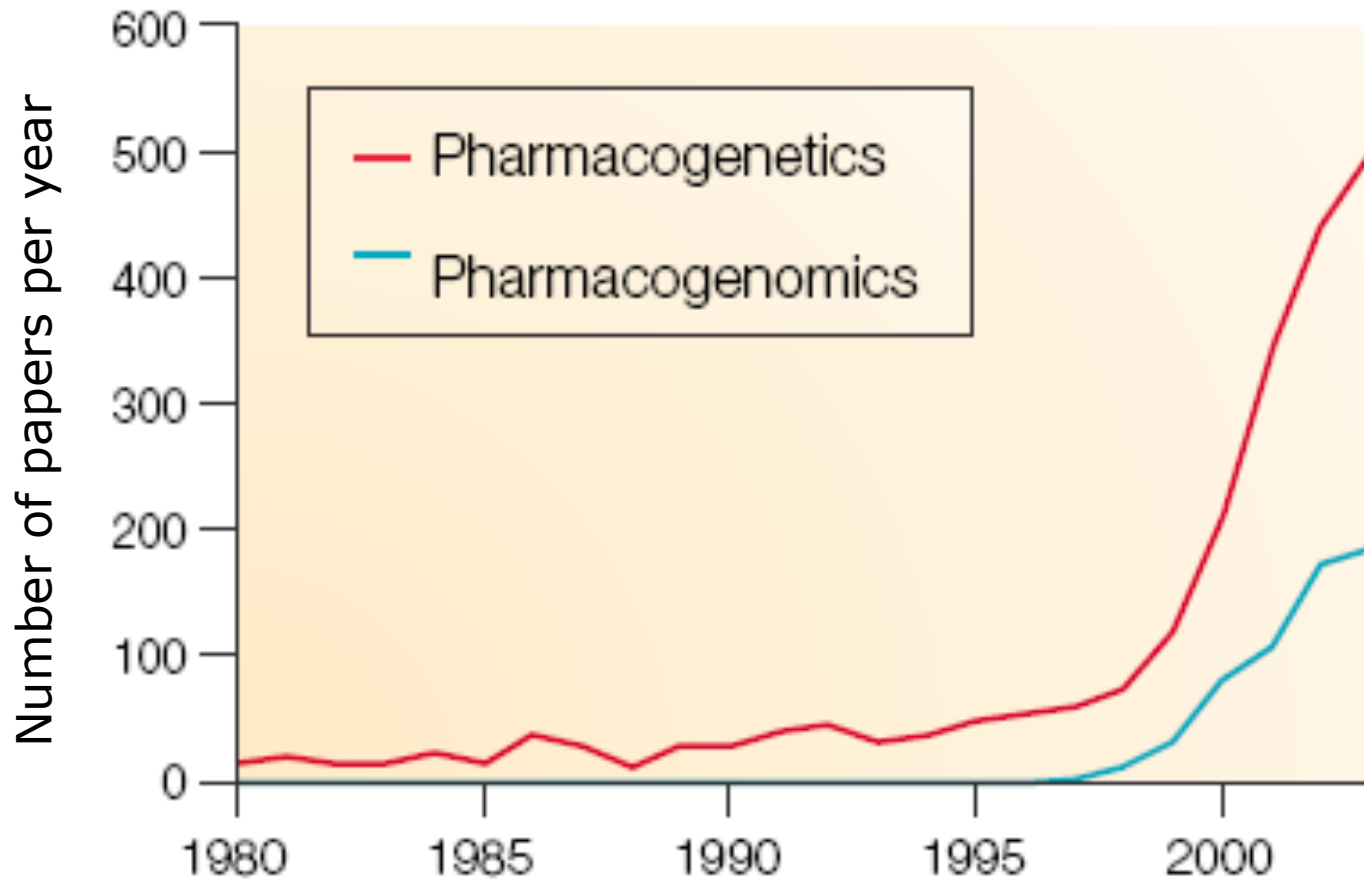
“Chemical  
Individuality”

# Historical Background



# Increasing interest in genetics and medicines

---





# Genetic variation can influence patient response at two levels

---

**Drug**



**Absorption**

**Distribution**

Pharmacokinetics

**Drug metabolism**



**Blood levels achieved**

Pharmacodynamics

**Drug target**

**Intermediate  
pathways**



**Response**

# Genetic variation can influence patient response at two levels

---

**Drug**



**Absorption**

**Distribution**

**Drug metabolism**



**Blood levels achieved**

Pharmacokinetics

Pharmacodynamics

Drug target

Intermediate  
pathways

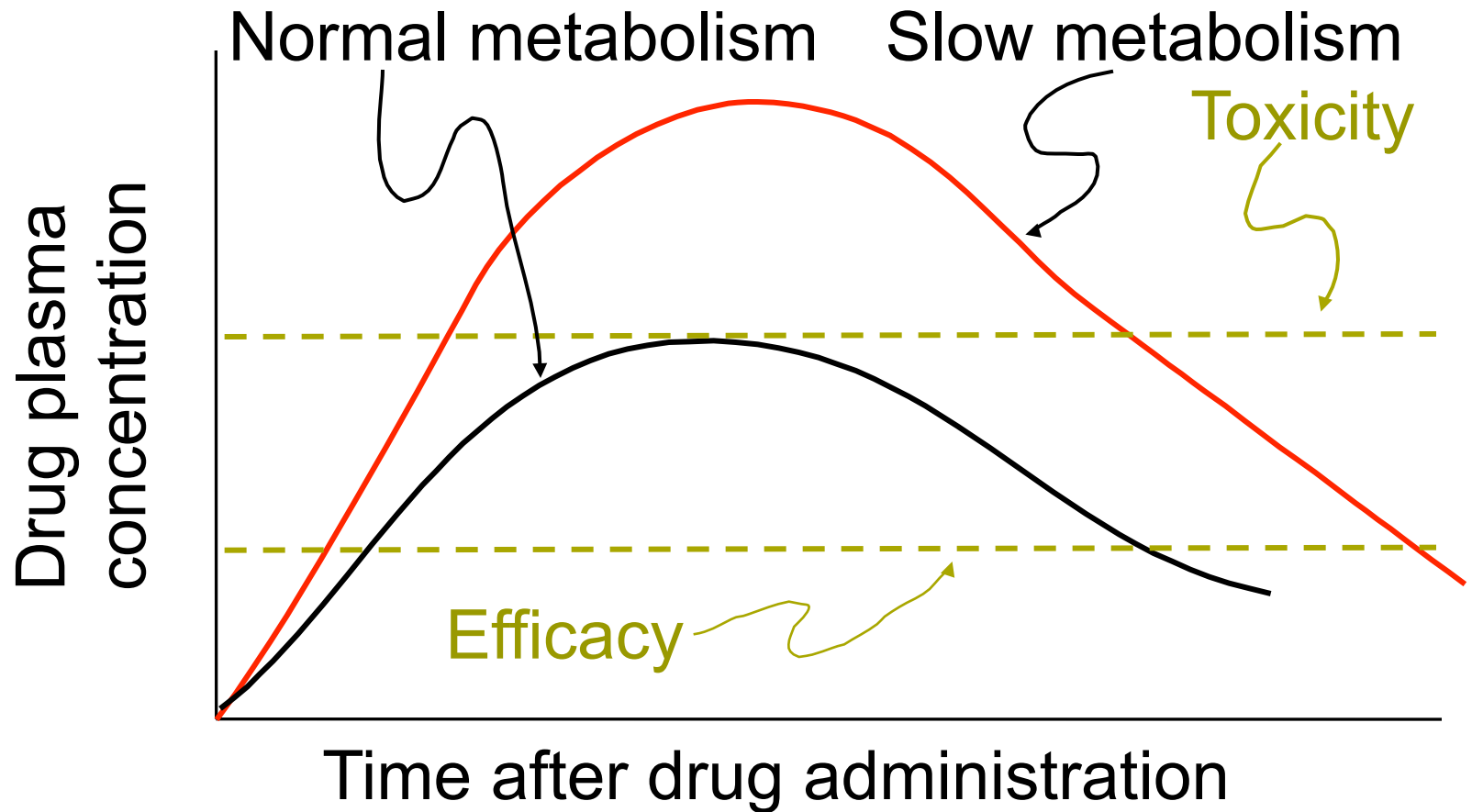


Response

Genetic variation can influence  
drug metabolism

# Poor drug metabolisers can be exposed to increased blood levels

---

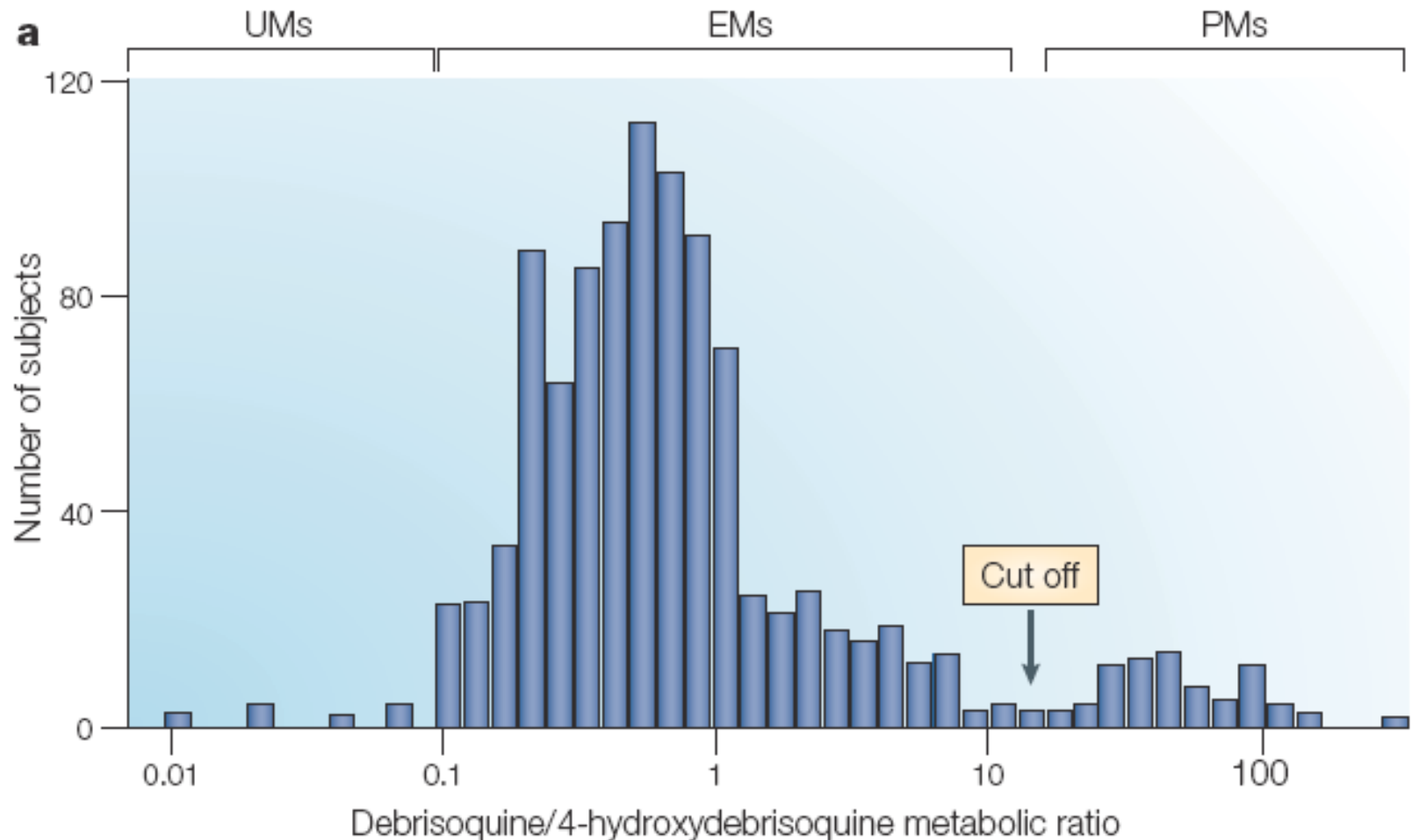


# When is genetic variation in drug metabolism important?

---

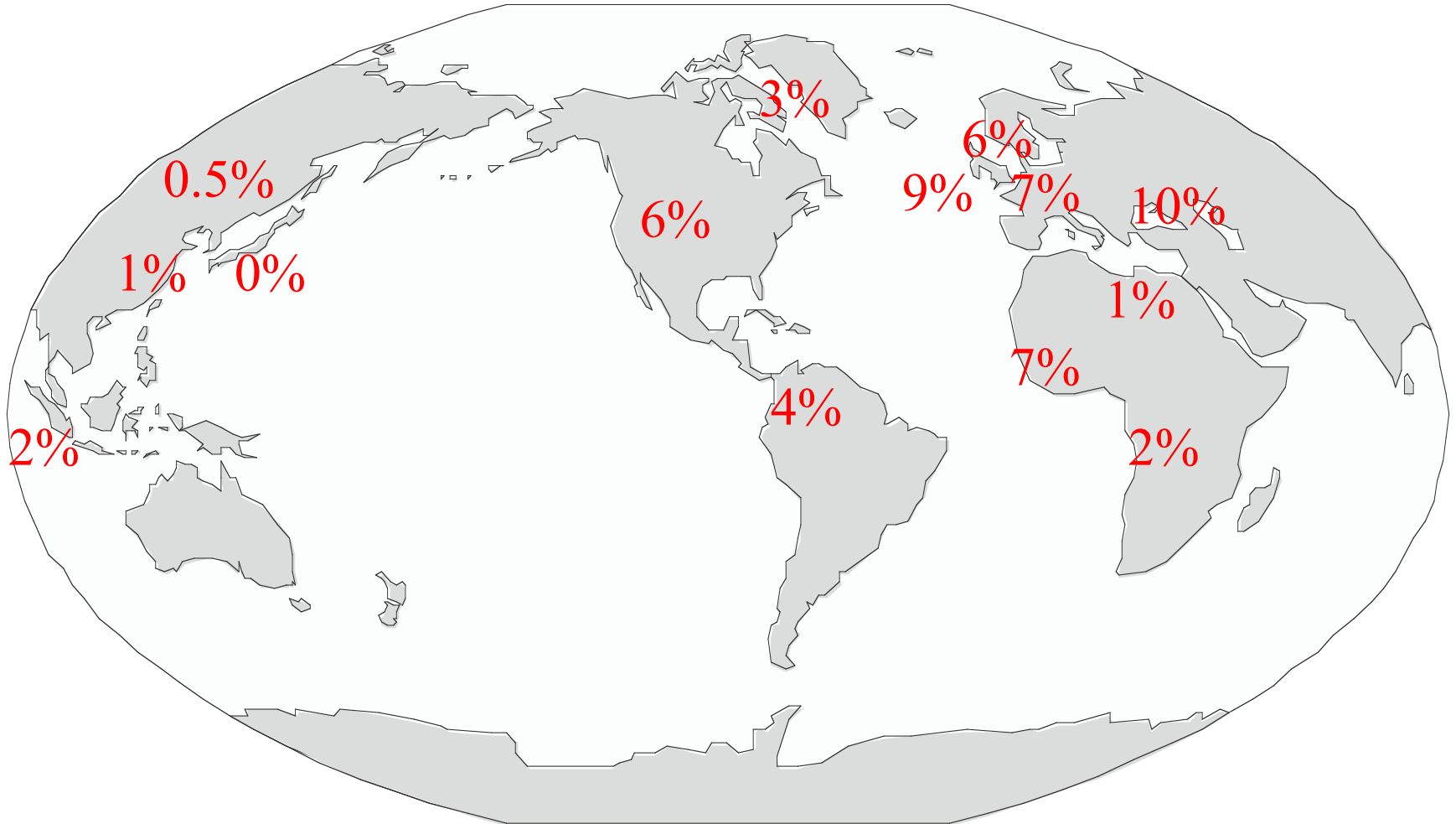
- When the drug has a steep dose-response curve
- When the therapeutic window is narrow
- When the drug is metabolised mainly by one enzyme

# Population distribution of metaboliser status



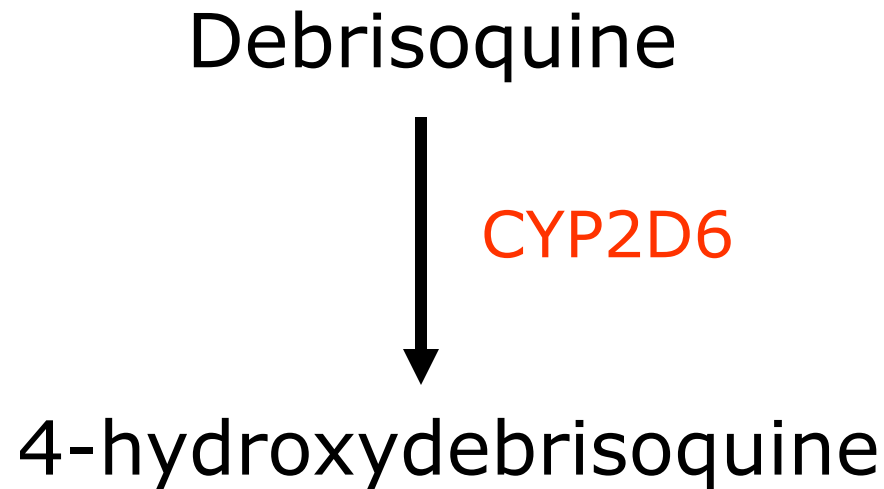
# Worldwide distribution of poor drug metabolisers (CYP2D6)

---



# Debrisoquine is metabolised largely by one enzyme

---





# Some drugs cleared by CYP2D6

---

$\beta$ -Adrenergic antagonists

e.g. metoprolol, bufuralol, propranolol

Neuroleptics

e.g. haloperidol, perphenazine

Anti-depressants

e.g. imipramine, amitriptyline, nortriptyline, paroxetine

Anti-arrhythmics

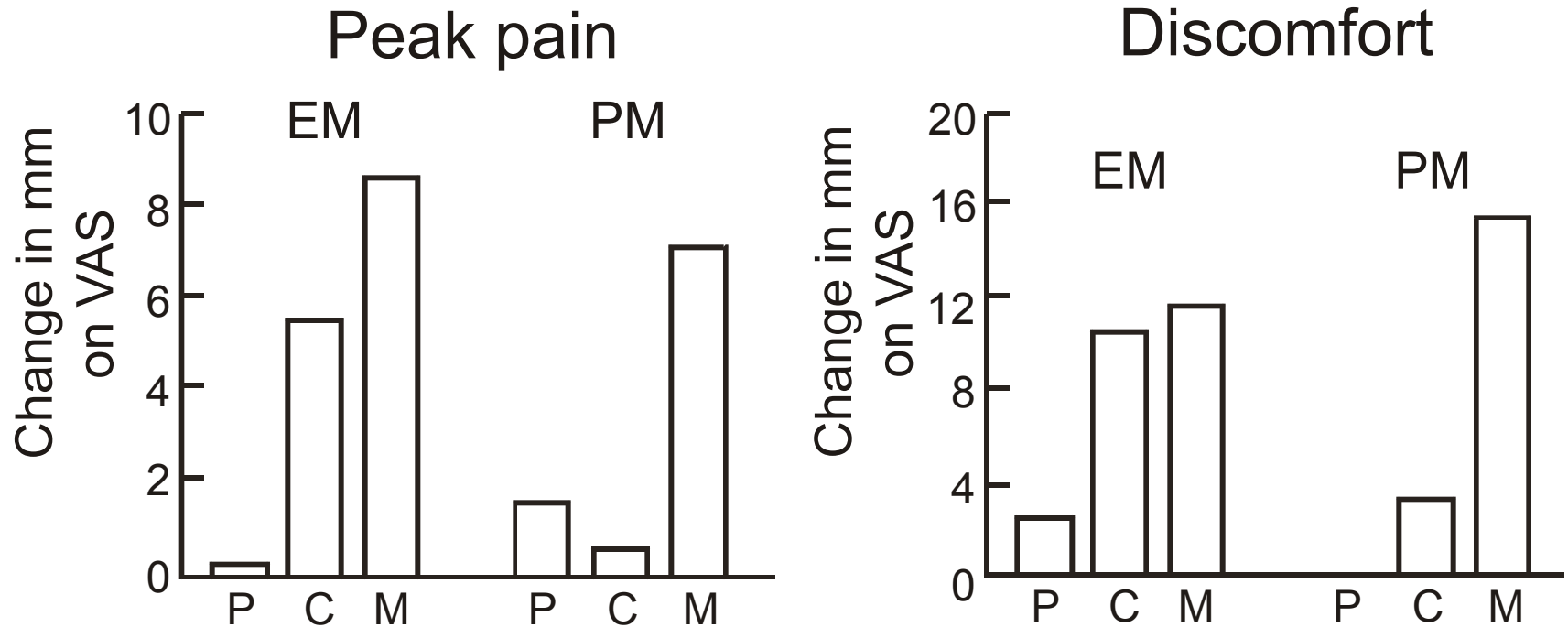
e.g. flecainide, encainide

Miscellaneous drugs

e.g. dextromethorphan, debrisoquine, sparteine,  
codeine

# CYP2D6 phenotype reduces effectiveness of codeine analgesia

From Poulsen *et al.* (1996)



P=placebo; C=codeine; M=morphine

# Poor metaboliser status can work both ways

---

## Poor metaboliser

Prodrug eg codeine  
activated by 2D6

Risk lack  
of effect

Active drugs  
inactivated by  
metabolism

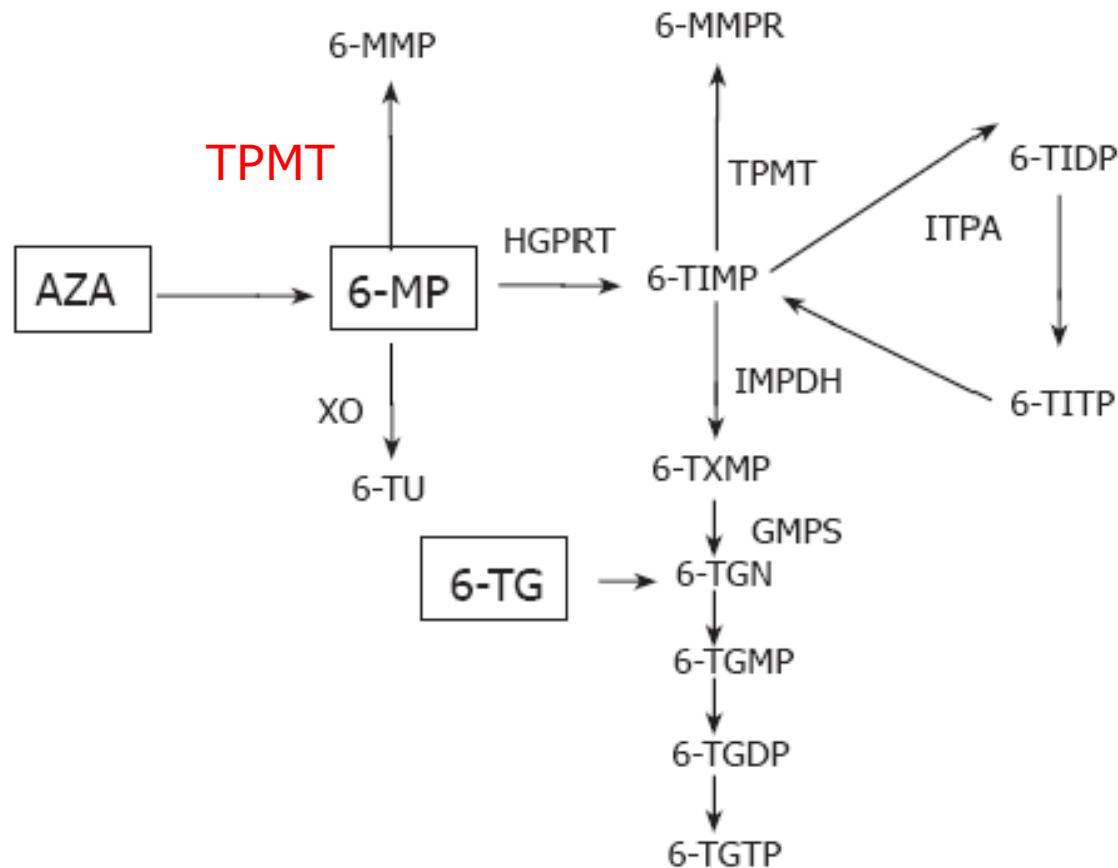
Risk of toxicity

# Genes on a chip

---



# Thiopurine S-Methyltransferase (TPMT)



# TPMT deficiency

---

- 1 in 300 (UK) have TPMT deficiency
- Several polymorphisms: G238C, G460A and A719G
- TPMT deficiency predicts severe neutropaenia following treatment with 6-mercaptopurine or azathioprine (but not other side effects)
- Some cases of severe neutropaenia occur with normal TPMT phenotype

# TPMT deficiency

---

- Azathioprine used in a wide range of inflammatory diseases, prescribed by specialists in respiratory medicine, dermatology, neurology, rheumatology, oncology, gastroenterology
- Product labelling amended to include reference to TPMT status, FDA have approved (2004) test for determination of TPMT status
- British Association of Dermatologists
  - Pretreatment TPMT determination in all patients requiring azathioprine
- British Thoracic Society
  - No requirement for testing TPMT status in interstitial lung disease guidelines before prescribing azathioprine
- British Society of Gastroenterology
  - TPMT status determination 'cannot yet be recommended as a prerequisite to therapy'

# Genetic variation can influence patient response at two levels

---

**Drug**



Absorption

Distribution

Drug metabolism



Blood levels achieved

Pharmacokinetics

Pharmacodynamics

**Drug target**

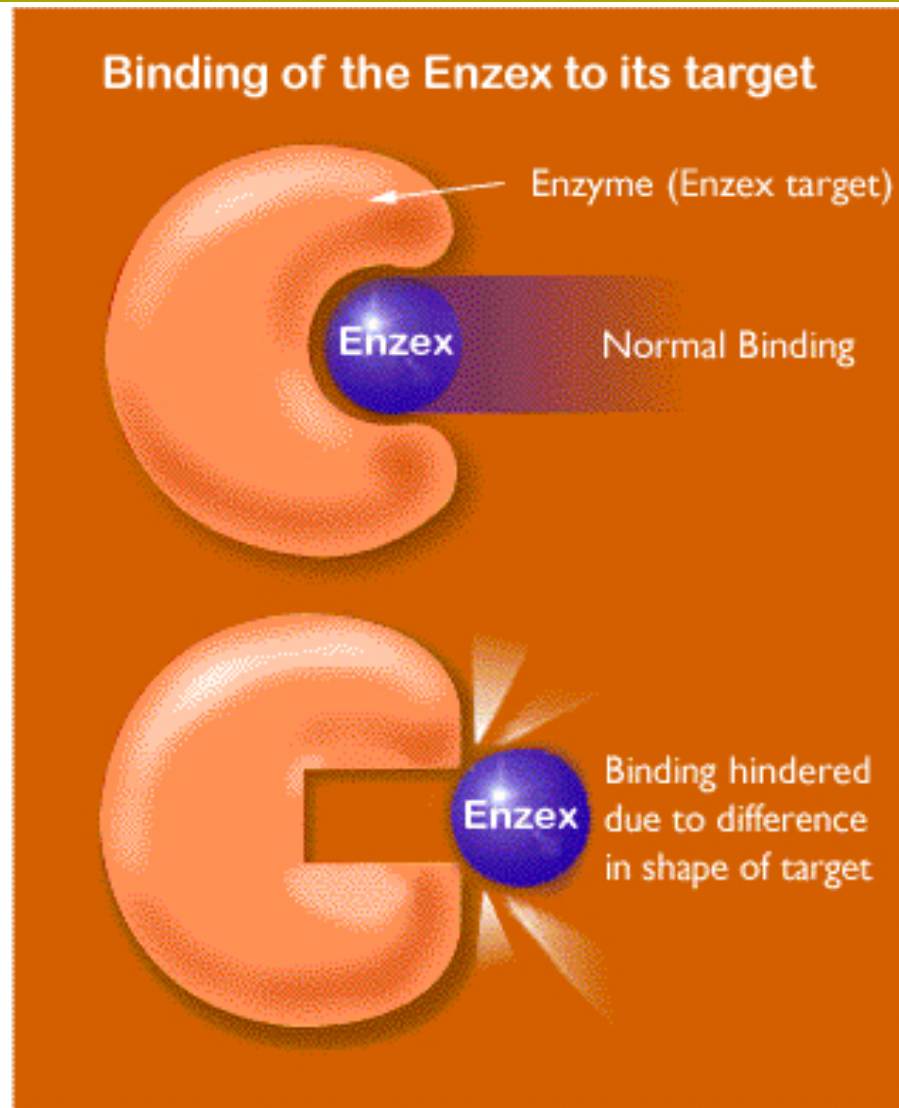
**Intermediate  
pathways**



**Response**



# Genetic variation in drug target

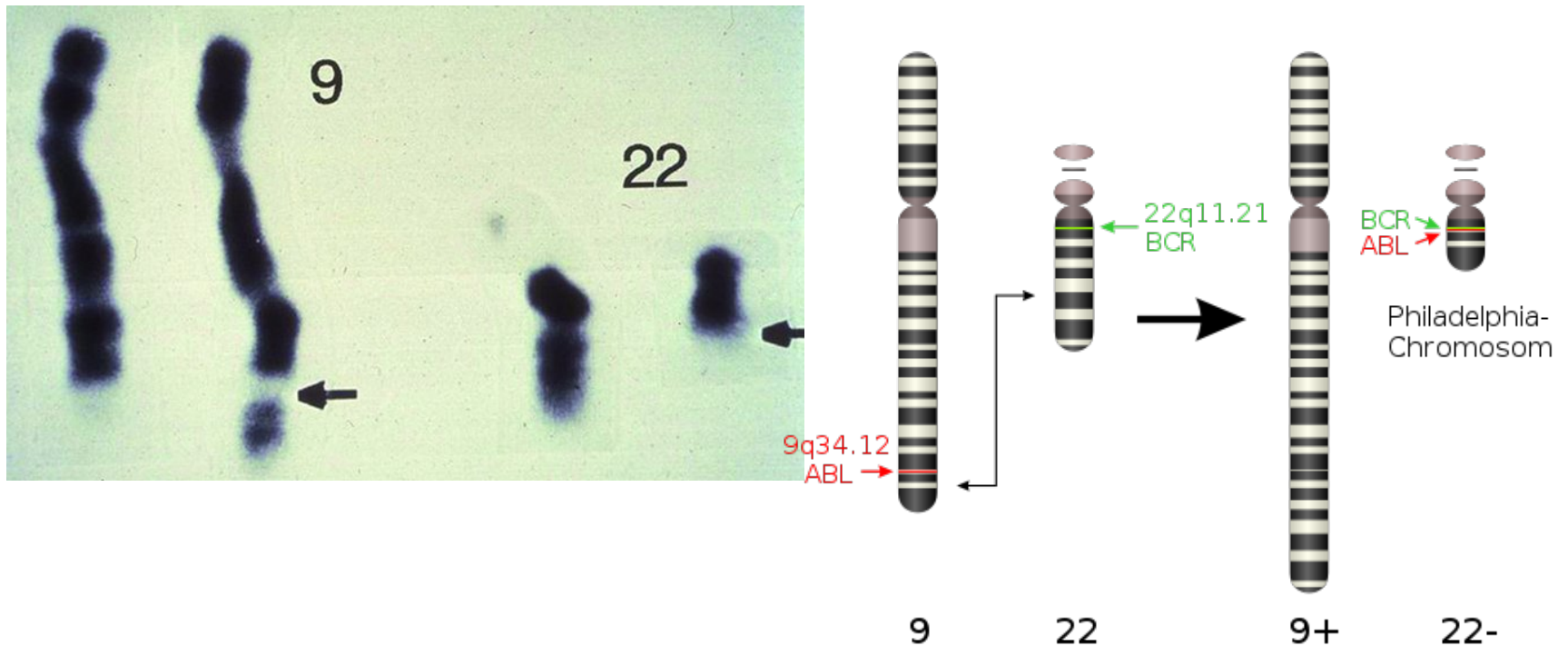


# Genomic variation affecting drug targets

---

- Malignancy
  - Gleevec
  - Herceptin
- Infectious diseases

# Imatinib (Gleevec): designer drug



# Genomics can be used identify good responder subgroups in cancer

---

## **Breast cancer – Herceptin**

Patients who respond best are those with tumours that overexpress the *ERBB2* (also known as *HER2/neu*) gene

# A polymorphism within a conserved $\beta_1$ -adrenergic receptor motif alters cardiac function and $\beta$ -blocker response in human heart failure

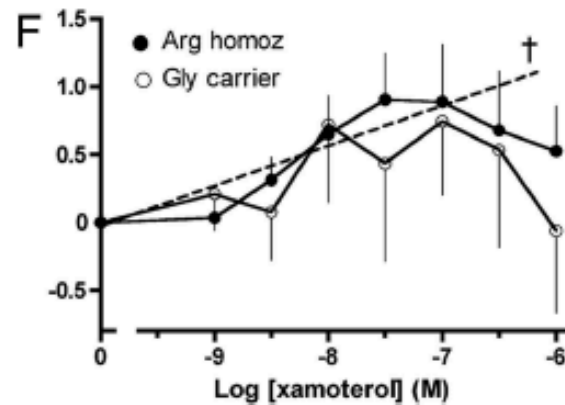
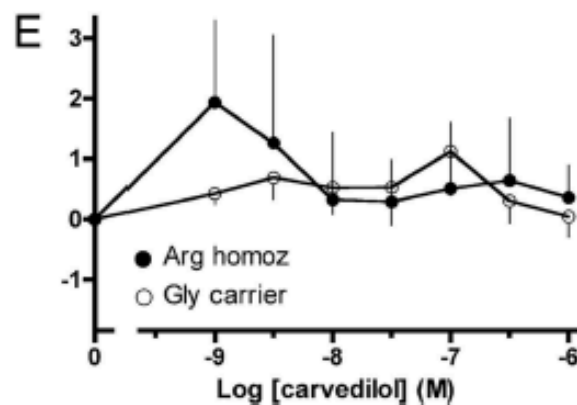
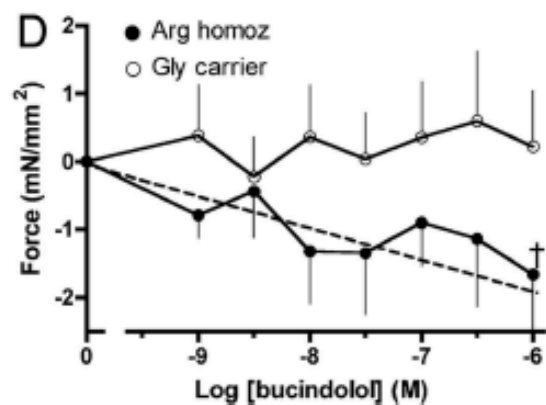
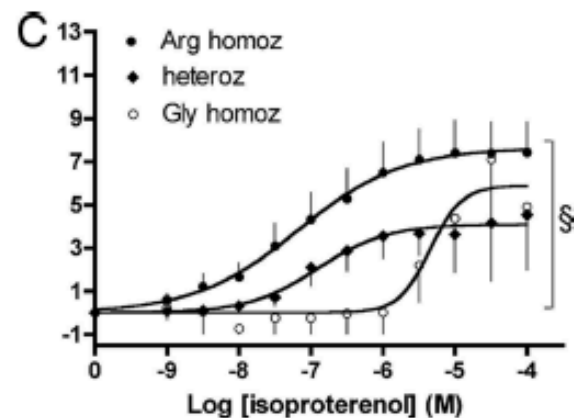
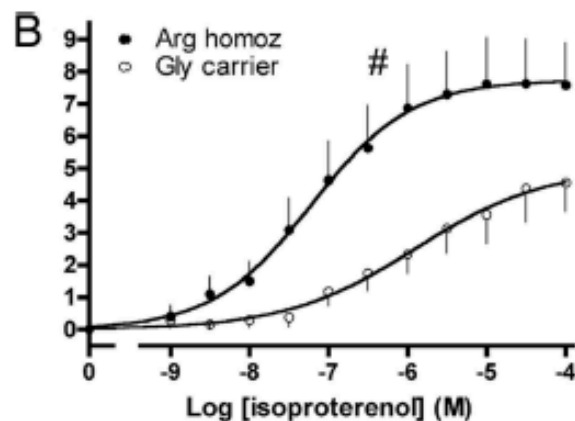
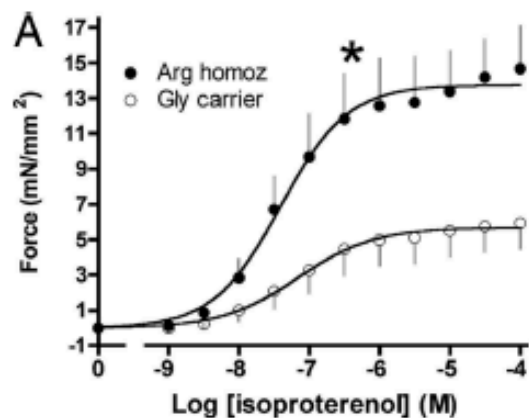
Stephen B. Liggett<sup>\*,†</sup>, Jeanne Mialet-Perez<sup>‡</sup>, Surai Thaneemit-Chen<sup>§</sup>, Stewart A. Weber<sup>¶</sup>, Scott M. Greene<sup>¶</sup>, Danielle Hodne<sup>¶</sup>, Bradley Nelson<sup>¶</sup>, Jennifer Morrison<sup>¶</sup>, Michael J. Domanski<sup>¶</sup>, Lynne E. Wagoner<sup>‡</sup>, William T. Abraham<sup>\*\*</sup>, Jeffrey L. Anderson<sup>††</sup>, John F. Carlquist<sup>††</sup>, Heidi J. Krause-Steinrauf<sup>§</sup>, Laura C. Lazzeroni<sup>§</sup>, J. David Port<sup>¶</sup>, Philip W. Lavori<sup>§</sup>, and Michael R. Bristow<sup>¶</sup>

species

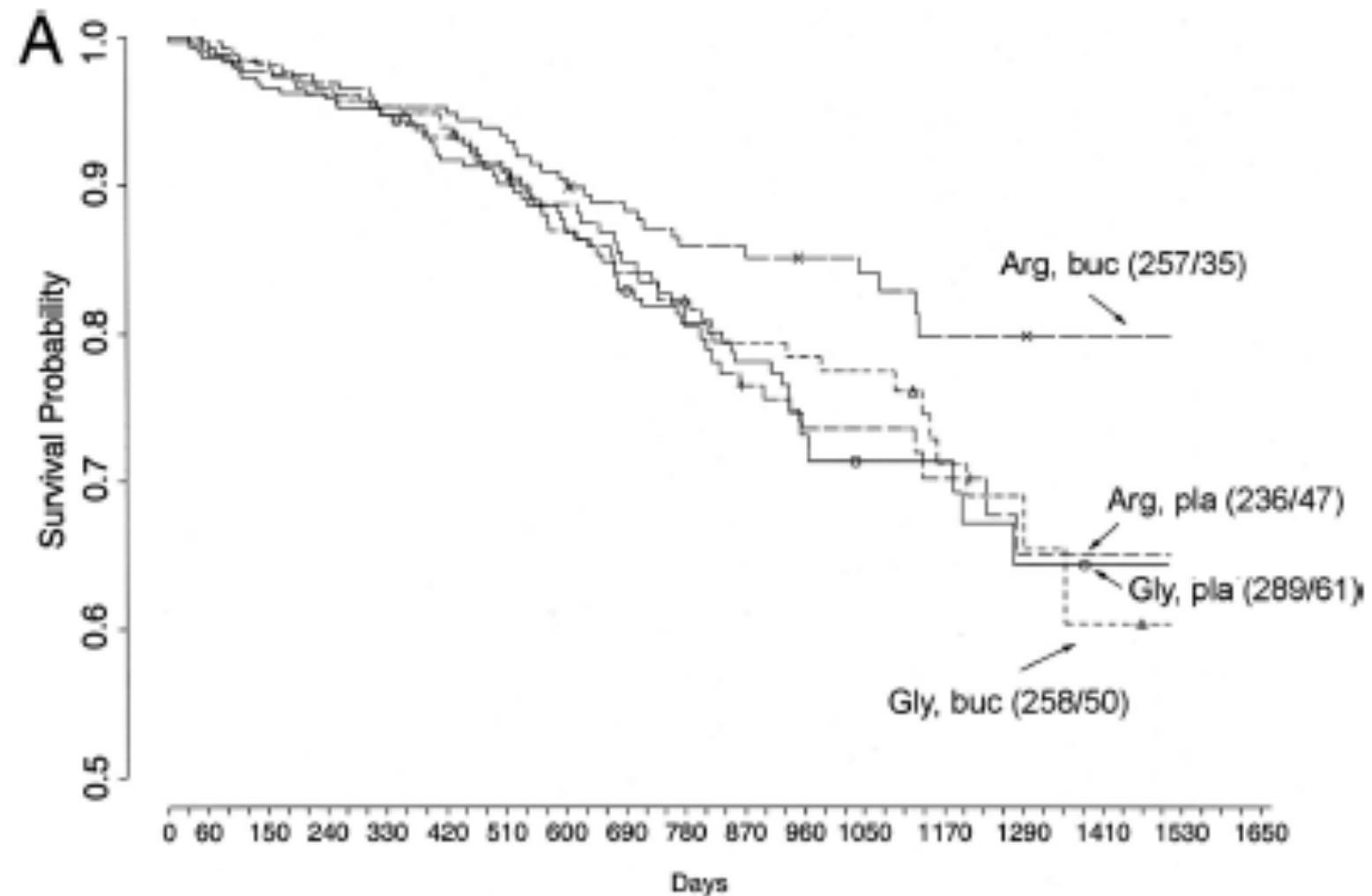
% identity

species	R	S	P	D	F	R	K	A	F	Q	R/G	L	L	C	C	A	R	R	A	% identity
Homo sapien	R	S	P	D	F	R	K	A	F	Q	R/G	L	L	C	C	A	R	R	A	100
Ovis aries	R	S	P	D	F	R	K	A	F	Q	R	L	L	C	C	A	R	R	A	100
Bos taurus	R	S	P	D	F	R	K	A	F	Q	R	L	L	C	C	A	R	R	A	100
Rattus norvegicus	R	S	P	D	F	R	K	A	F	Q	R	L	L	C	C	A	R	R	A	100
Mus musculus	R	S	P	D	F	R	K	A	F	Q	R	L	L	C	C	A	R	R	A	100
Pan troglodytes	R	S	P	D	F	R	K	A	F	Q	R	L	L	C	C	V	R	R	A	94
Rhesus macaque	R	S	P	D	F	R	N	A	F	Q	R	L	L	C	C	A	R	R	A	94
Canis familiaris	R	S	P	D	F	R	R	A	F	Q	R	L	L	C	C	A	R	R	A	94
Felis catus	R	S	P	D	F	R	K	A	F	Q	R	L	L	C	F	A	R	R	A	94
Sus scrofa	R	C	P	D	F	R	K	A	F	Q	R	L	L	C	C	A	R	R	V	94
Tetraodon nigroviridis	R	S	P	D	F	R	K	A	F	K	R	L	L	C	C	A	R	Q	A	84
Xenopus laevis	R	S	P	D	F	R	K	A	F	K	R	L	L	C	C	P	K	K	A	78
Meleagris gallopavo	R	S	P	D	F	R	S	A	F	K	R	L	L	C	F	P	R	K	A	78

# A polymorphism within a conserved $\beta_1$ -adrenergic receptor motif alters cardiac function and $\beta$ -blocker response in human heart failure

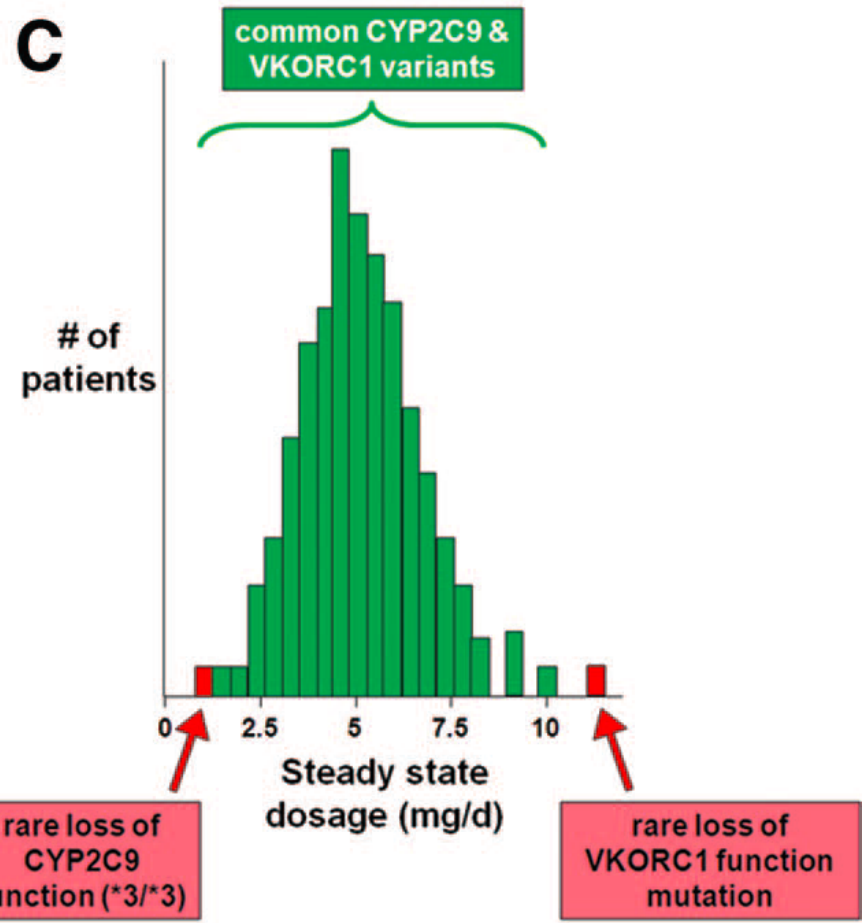
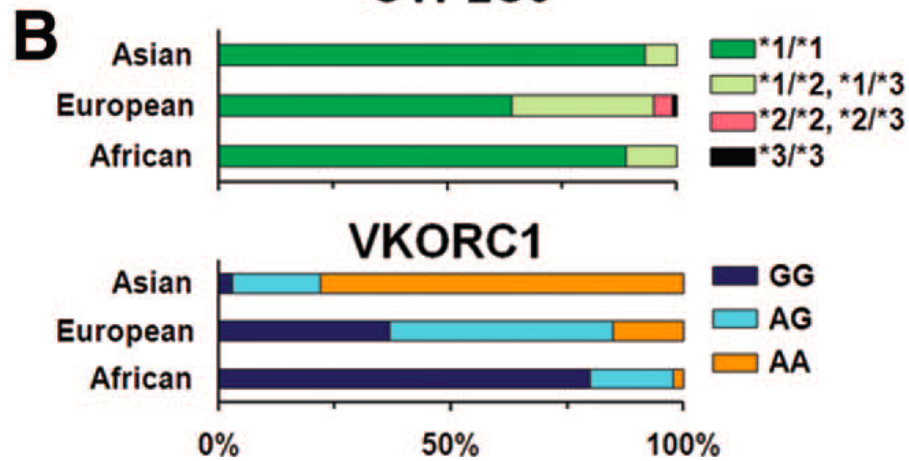
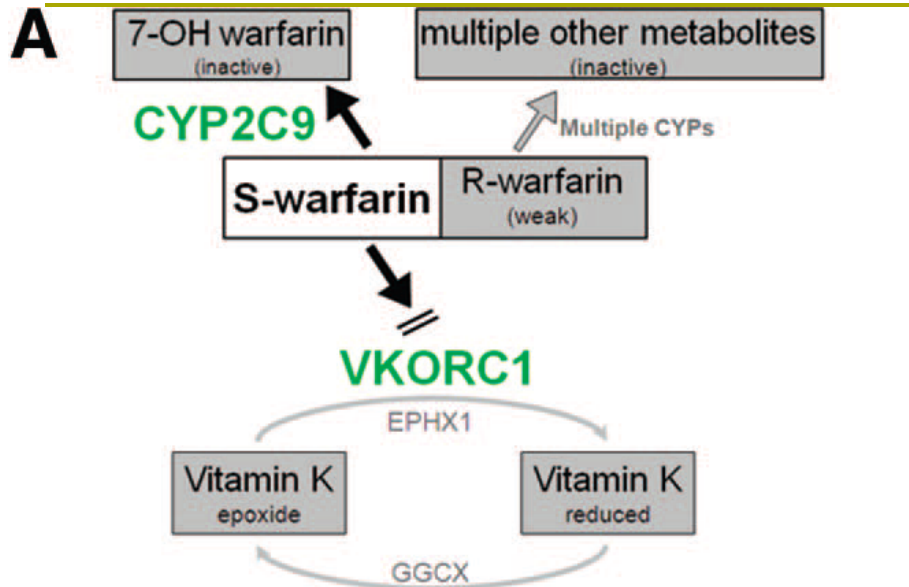


# A polymorphism within a conserved $\beta_1$ -adrenergic receptor motif alters cardiac function and $\beta$ -blocker response in human heart failure





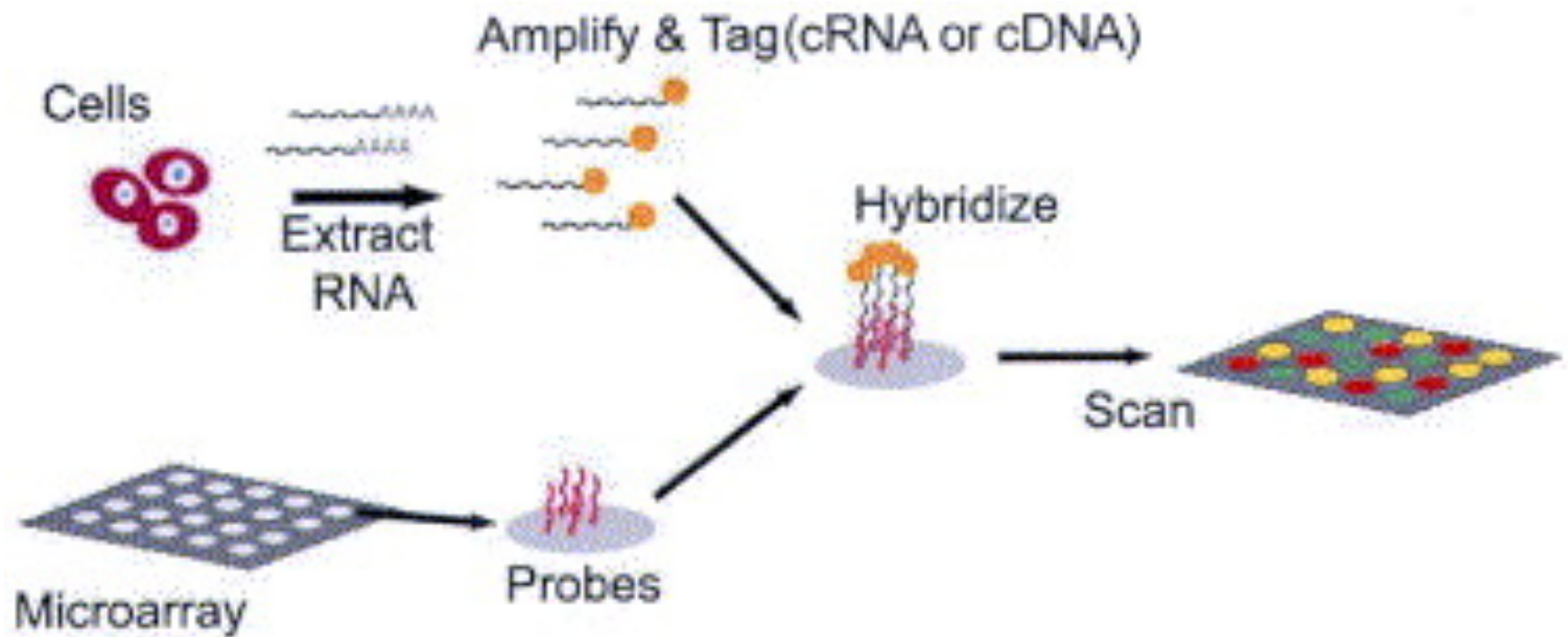
# PK and PD variants and warfarin dose



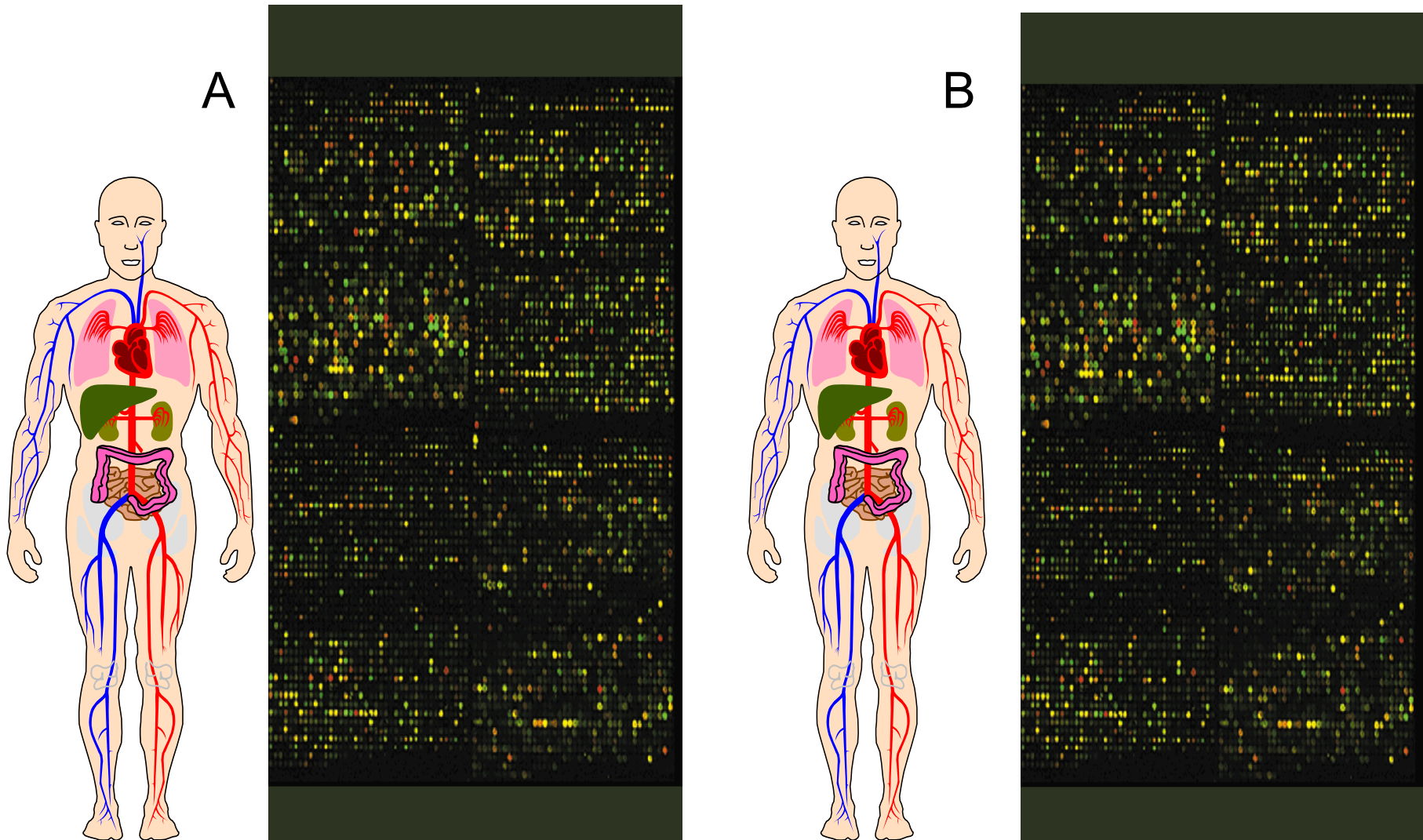


# Gene expression analysis

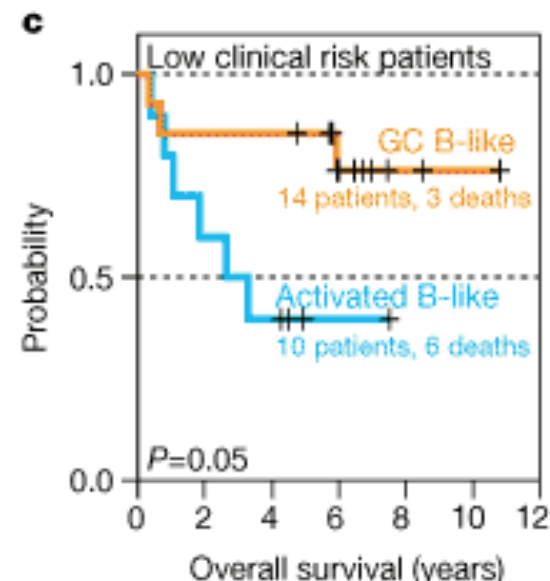
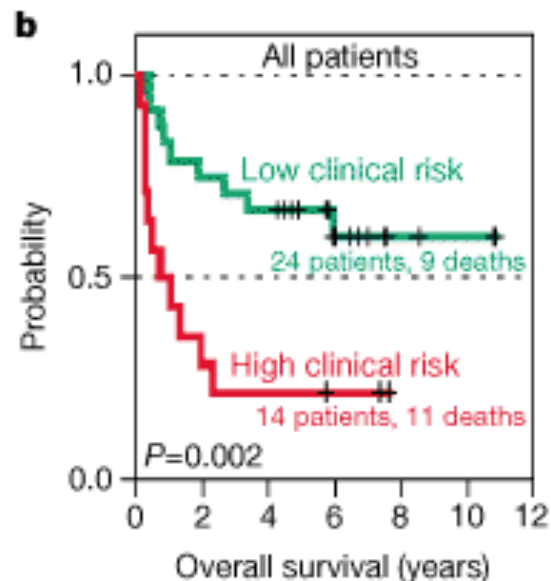
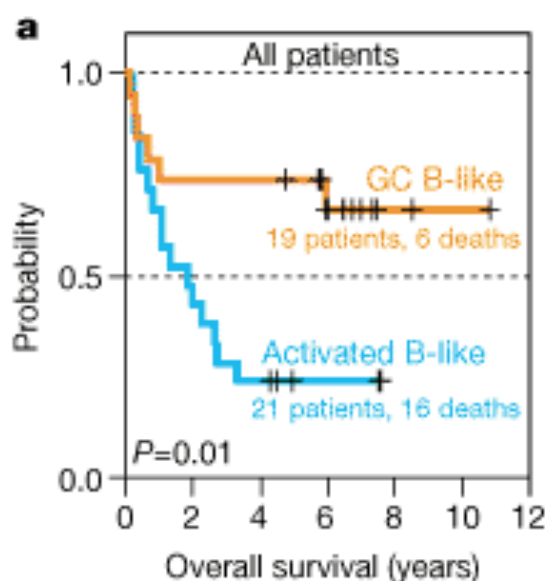
---



# Genetics can be used for more accurate diagnosis of cancer type



# Survival according to gene expression



# The NEW ENGLAND JOURNAL of MEDICINE

ESTABLISHED IN 1812

OCTOBER 28, 2010

VOL. 363 NO. 18

## BACKGROUND

Oncogenic fusion genes consisting of *EML4* and anaplastic lymphoma kinase (*ALK*) are present in a subgroup of non–small-cell lung cancers, representing 2 to 7% of such tumors. We explored the therapeutic efficacy of inhibiting *ALK* in such tumors in an early-phase clinical trial of crizotinib (PF-02341066), an orally available small-molecule inhibitor of the *ALK* tyrosine kinase.

## CONCLUSIONS

The inhibition of *ALK* in lung tumors with the *ALK* rearrangement resulted in tumor shrinkage or stable disease in most patients. (Funded by Pfizer and others; ClinicalTrials.gov number, NCT00585195.)

# Gene expression data and clinical practice

---

	<b>Application available now</b>	<b>Potential or emerging application</b>
<b>Oncology</b>		
Breast	Predict recurrence	Predict response to therapy
Lymphoma	Classify / guide therapy	Classify subtypes/ guide therapy
Unknown primary		Assess prognosis
Neuroblastoma		
<b>Infectious disease</b>		
HIV	Classify virus	Predict response to therapy
SARS		
Hepatitis C		
<b>Cardiovascular</b>	?	
Hypertension		
Atherosclerosis		
Cardiac Failure		



# Nobody is perfect

## How science has mastered the code

6bn

Number of DNA letters in the human genome, arranged in two sets of 3 billion

20,000

Approximate number of genes in the human genome. Genes are strings of DNA letters that make proteins

15m

Approximate number of known places where DNA spelling commonly varies

3m

Places in a genome where one DNA letter has an unusual spelling. The project will list about 95%

250-300

The genes in every person that work abnormally

Source: 1,000 Genomes Project



# What is a SNP

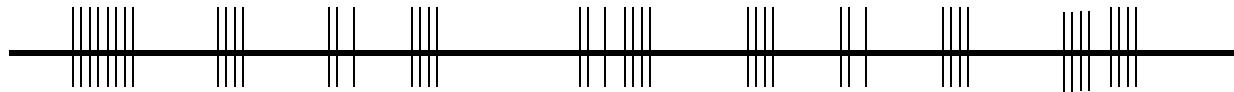
Different people can have a different nucleotide or base at a given location on chromosome



... G G T A A C T G ...

... G G C A A C T G ...

# What is a SNP map



Location of SNPs on human DNA

# How a SNP map can be used to predict a response to a drug

## Genotype profile

Patients with good response to drug X



Patients with poor response to drug X



Predictive of efficacy

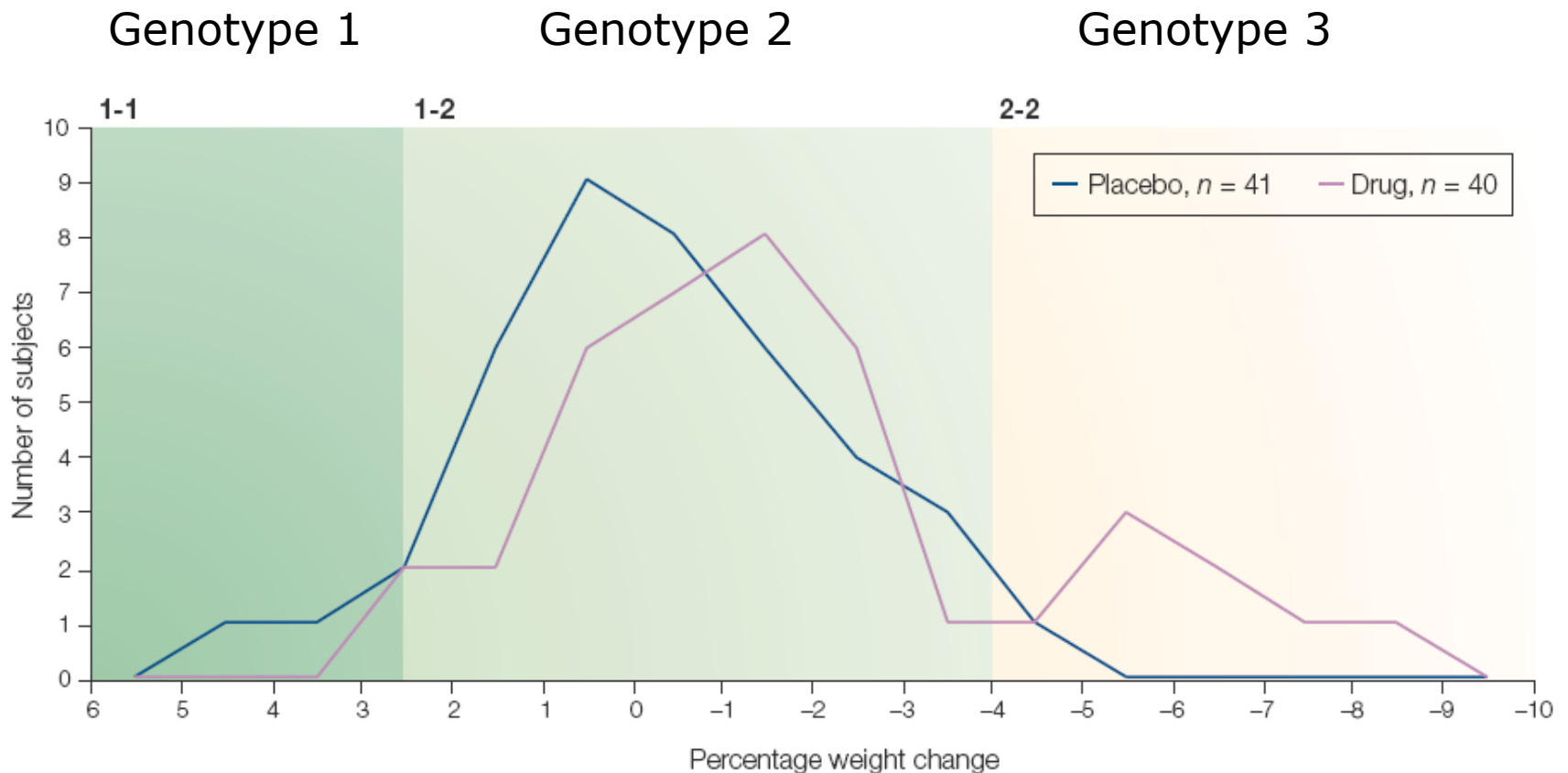


Predictive of a poor response





# Distribution of weight change by genotype



# Pharmacogenomics and adverse events

Drug	Adverse Drug Reaction		Genetic Risk Factor		
	Reaction	Prevalence	Risk Allele	Freq. <sup>1</sup>	Effect <sup>2</sup>
Clopidogrel	Cardiovascular events	0.13	<i>CYP2C19*2/3/4/5</i>	0.03	3
Gefitinib	Diarrhea	0.28	<i>ABCG2 Q141K</i>	0.07	5
Isoniazid	Hepatotoxicity	0.15	<i>CYP2E1*1 &amp; NAT2</i>	0.13 <sup>3</sup>	7
Augmentin	Hepatotoxicity	<0.001	<i>HLA-DRB1*1501</i>	0.20	10
Irinotecan	Neutropenia	0.20	<i>UGT1A1*28</i>	0.32	28
Ticlopidine	Hepatotoxicity (cholestatic)	<0.001	<i>HLA-A*3303</i>	0.14	36
Tranilast	Hyperbilirubinemia	0.12	<i>UGT1A1*28</i>	0.30	48
Flucloxacillin	Hepatotoxicity	<0.001	<i>HLA-B*5701</i>	0.04	81
Allopurinol	Severe cutaneous reaction	<0.001	<i>HLA-B*5801</i>	0.15	678
Abacavir	Hypersensitivity reaction	0.08	<i>HLA-B*5701</i>	0.04	>1000
Carbamazepine	Stevens-Johnson syndrome	<0.001	<i>HLA-B*1502</i>	0.04	>1000

# Drugs with pharmacogenomic tests in label

---

Antiviral	Abacavir Miraviroc	HLA-B*5701 CCR5
Cardiology	Warfarin	2Cp, VKCOR1
Neuropharm	Carbemezepine	HLA-B*1502
Oncology	Trastuzumab Irotectan Azothioprine Gefitinib Cetuximab Panitumumab	HER2 UGT1A*28 TMPT status EGFR status KRAS status
Pain	Codeine	2D6

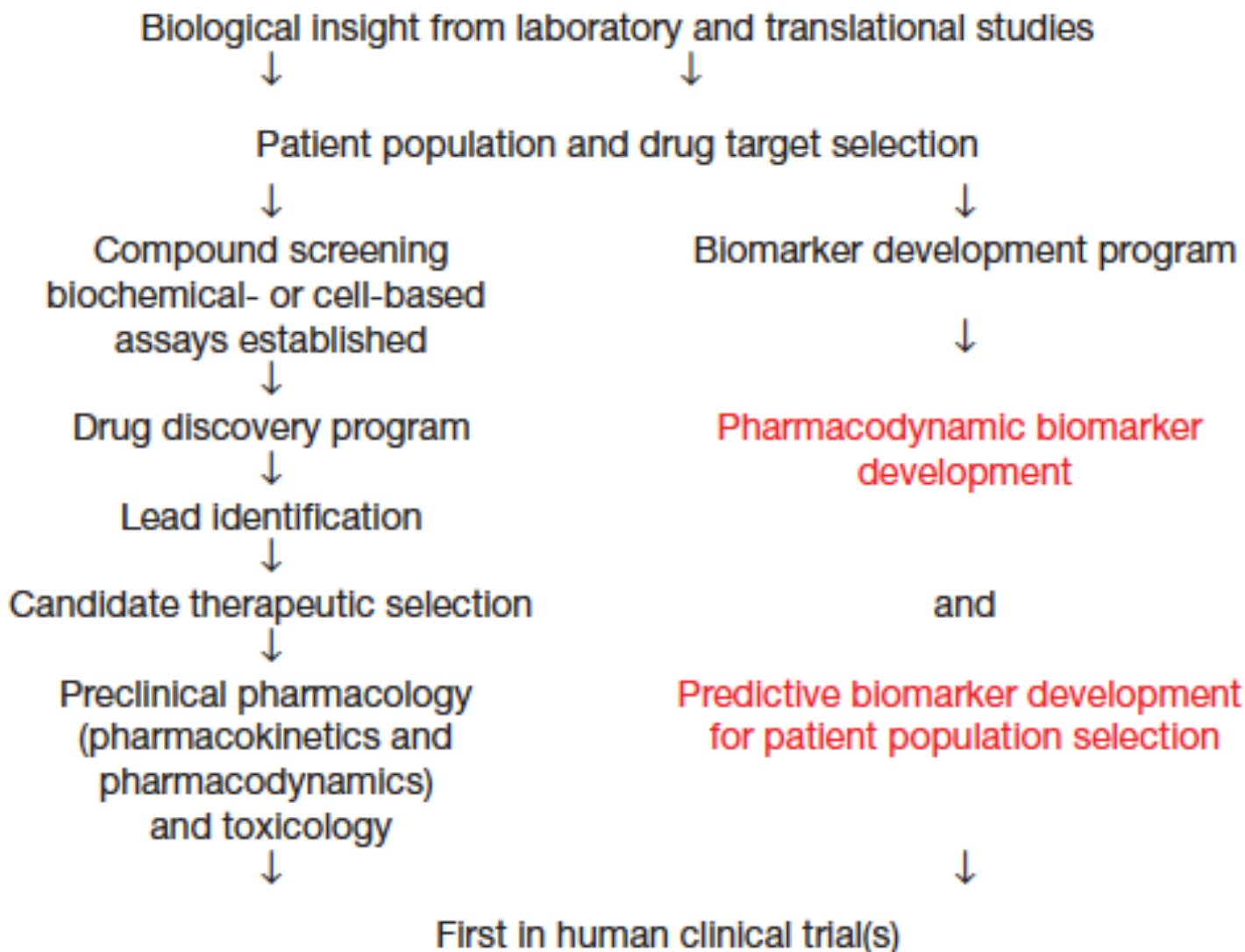
# Genomics: A powerful tool for drug target identification and validation

---

- 7,000 rare diseases worldwide
- Medicines for only 80-90 of them
- Know the cause, have the target, can recruit the right patients

# Recommended schema for anti-cancer drug development

---



# Hurdles to implementation

---

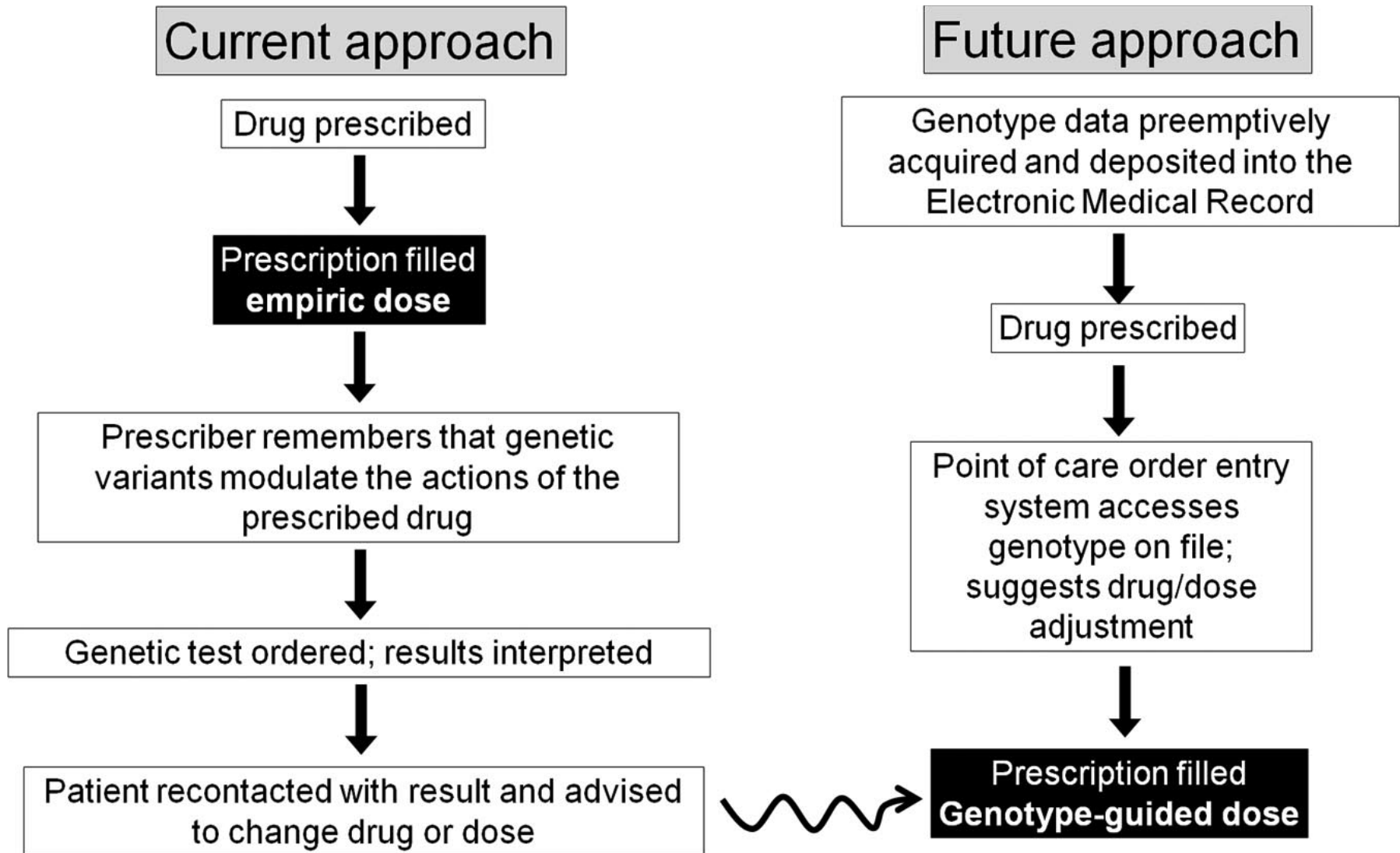
- Physicians
  - Education - number of variants growing
  - Needs a laboratory test – inconvenience and associated costs
- Regulators
  - The level of evidence required to adopt a specific genetic test as a guide to practice is not established.
- Pharmaceutical companies
  - Dissects the market
- Patients
  - Suspicion and sensitivity to genetic testing

# Francis Collins

---

“The limiting factor right now is that oftentimes, if you are ready to write a prescription, you do not want to wait a week to find out the genotype before you decide whether you’ve got the right dose and the right drug. But if everybody’s DNA sequence is already in their medical record and it is simply a click of the mouse to find out all the information you need, then there is going to be a much lower barrier to beginning to incorporate that information into drug prescribing. If you have the evidence, it will be hard, I think, to say that this is not a good thing. And once you’ve got the sequence, it’s not going to be terribly expensive. And it should improve outcomes and reduce adverse events.”

# Current and future approaches





# References

---

- Roden et al Pharmacogenomics: The Genetics of Variable Drug Responses Circulation 2011; 123: 1661

# Use of pharmacogenomics to deliver right medicine to right patient

---

