

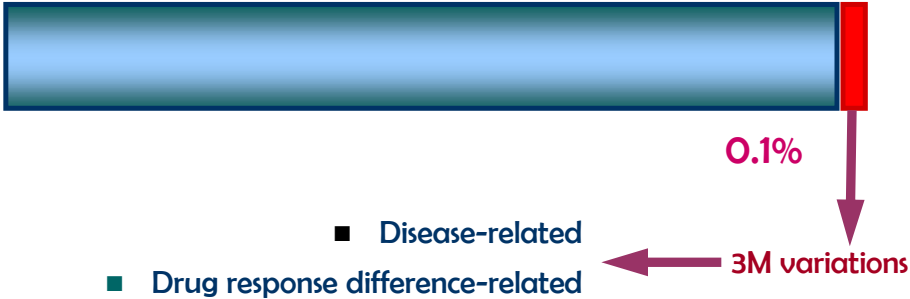
## Personalized Medicine in China

Hong-Hao Zhou, MD & Professor  
Institute of Clinical Pharmacology  
Pharmacogenetics Research Institute  
Central South University

### Genetic variations of human individuals

- The human genome contains approximately 3 billion of base pairs
- That code over thirty thousand genes

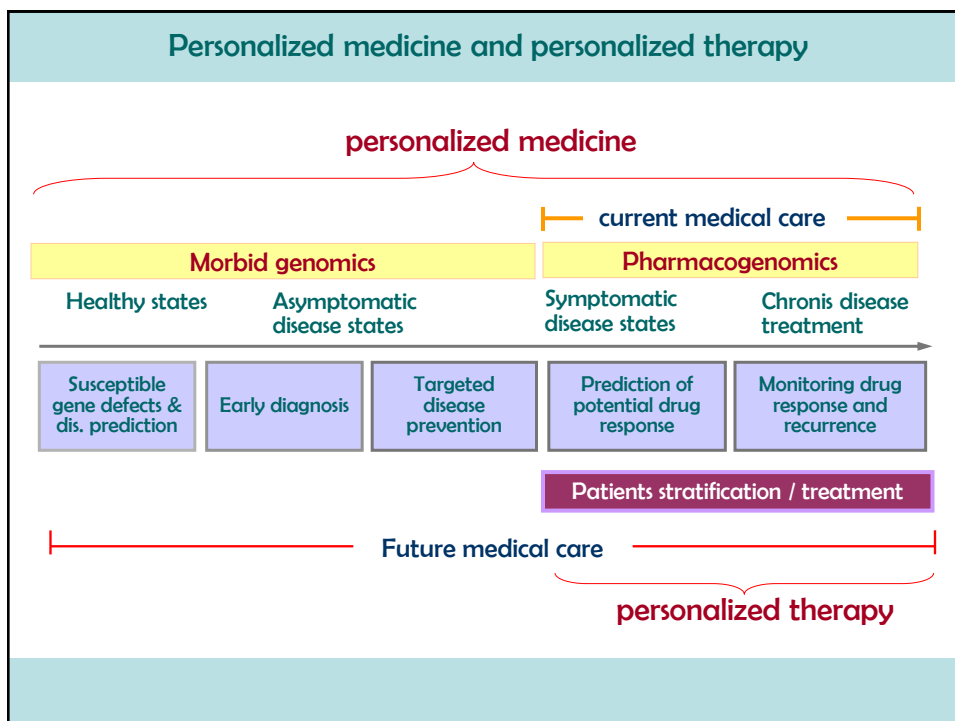
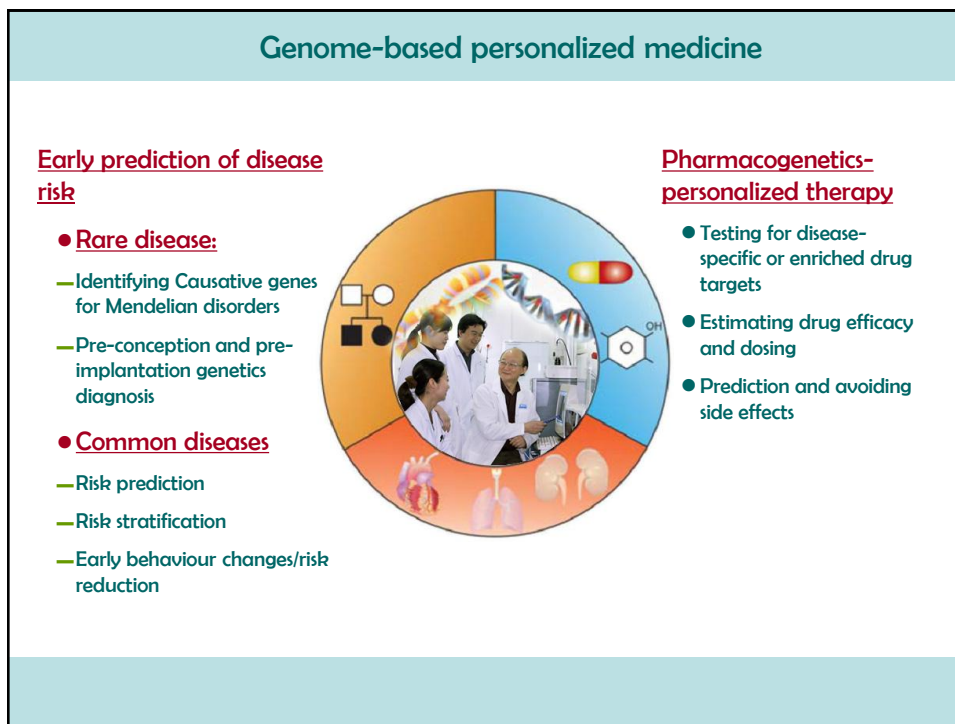
Individual DNA sequence: 99.9% identical

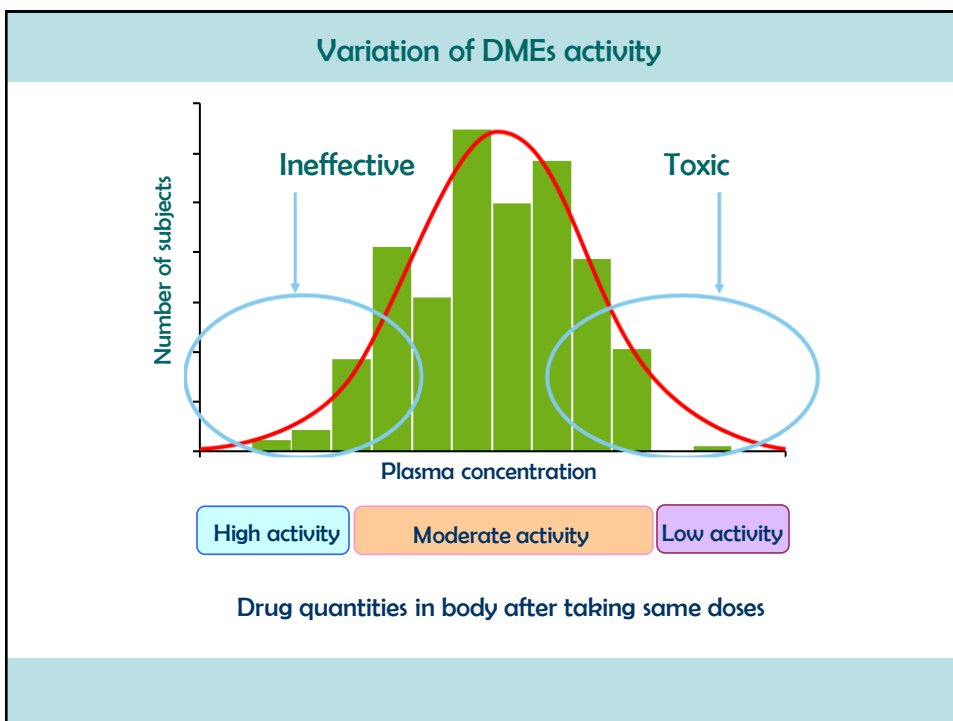
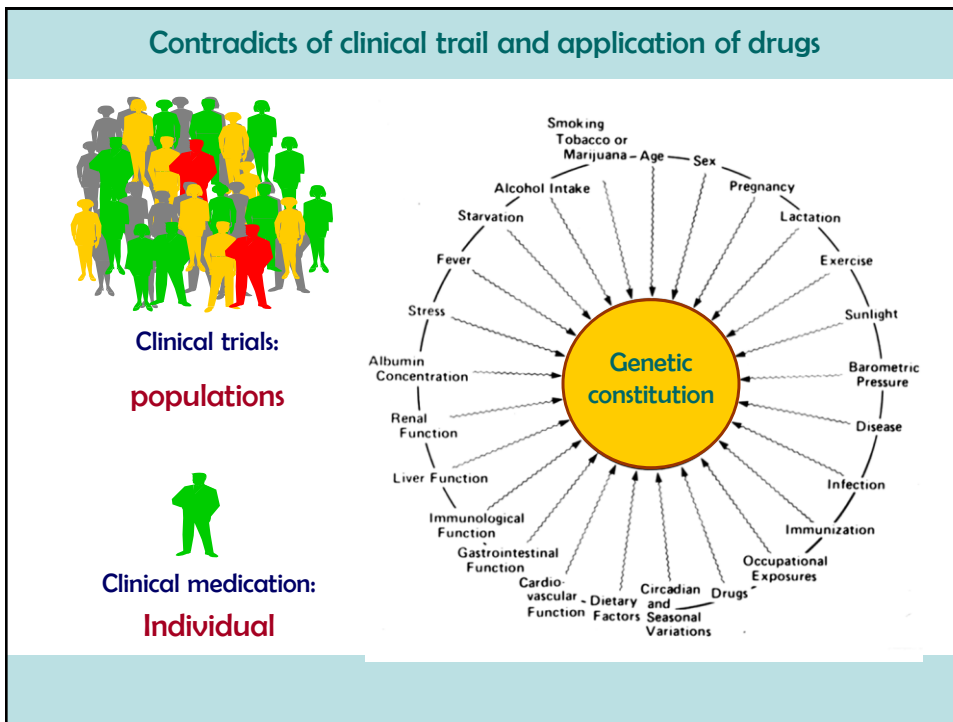


0.1%

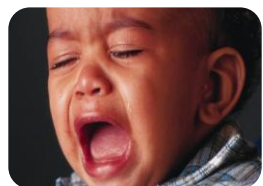
3M variations

- Disease-related
- Drug response difference-related

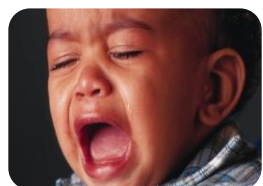




## Genetic variation of drug receptors induce Interindividual difference in drug effect and safety



Both have acute asthma



→ Asthma ends

β2 adrenergic receptor agonist  
Albuterol  
Salmeterol

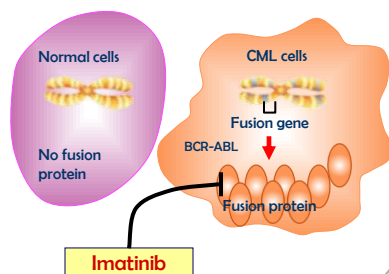
Variation in β2 adrenergic receptor (Arg/Arg16)

Death ?

## Personalized medication application 1: Testing for disease-specific or enriched drug targets

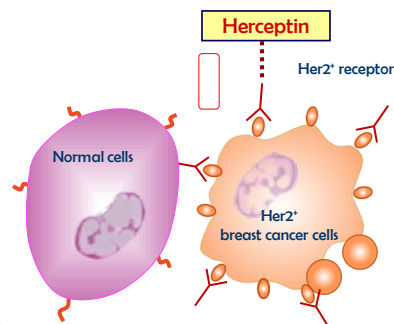
### Specific targets

- Imatinib, BCR-ABL tyrosine kinase inhibitor, used in the treatment of bcr-abl fusion gene-positive chronic myeloid leukemia



### Overexpression targets

- Herceptin, humanized Her-2 monoclonal antibody, high affinity to Her-2, used in the treatment of Her-2 receptor overexpressed breast cancer.

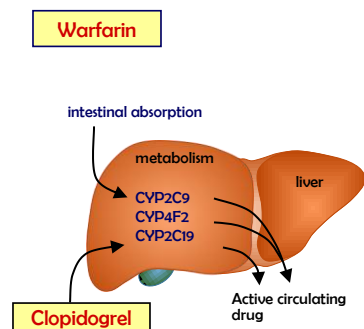


Salari K, Eur Heart J, 2012; 33, 1564–70

## Personalized medication application 2: prediction of drug effect and dosage

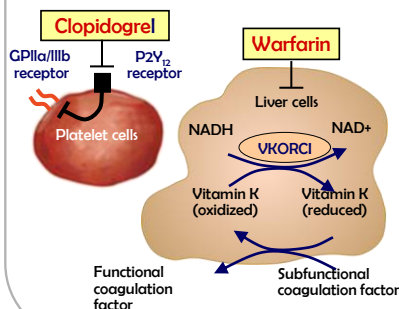
### Pharmacokinetics

- Clopidogrel (prodrug): CYP2C19 polymorphism
- Warfarin: CYP2C9 polymorphism



### Pharmacodynamics

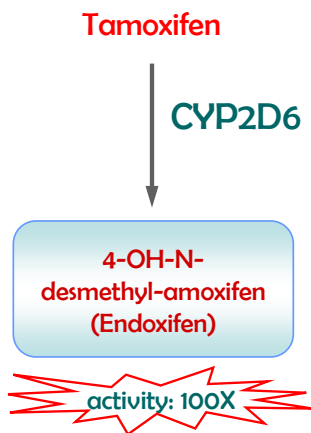
- Platelet membrane glycoprotein (GP)II $\alpha$ /III $\beta$  and membrane protein P2Y $_12$  receptor polymorphism: clopidogrel
- VitK epoxidation reductase polymorphism: warfarin



Salari K, Eur Heart J, 2012; 33, 1564–70

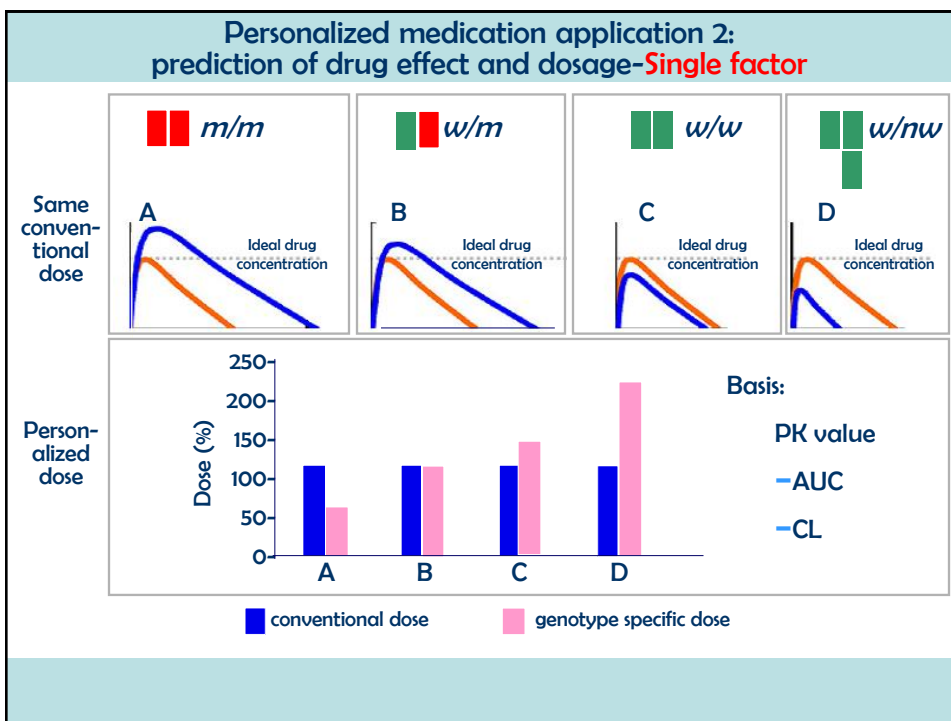
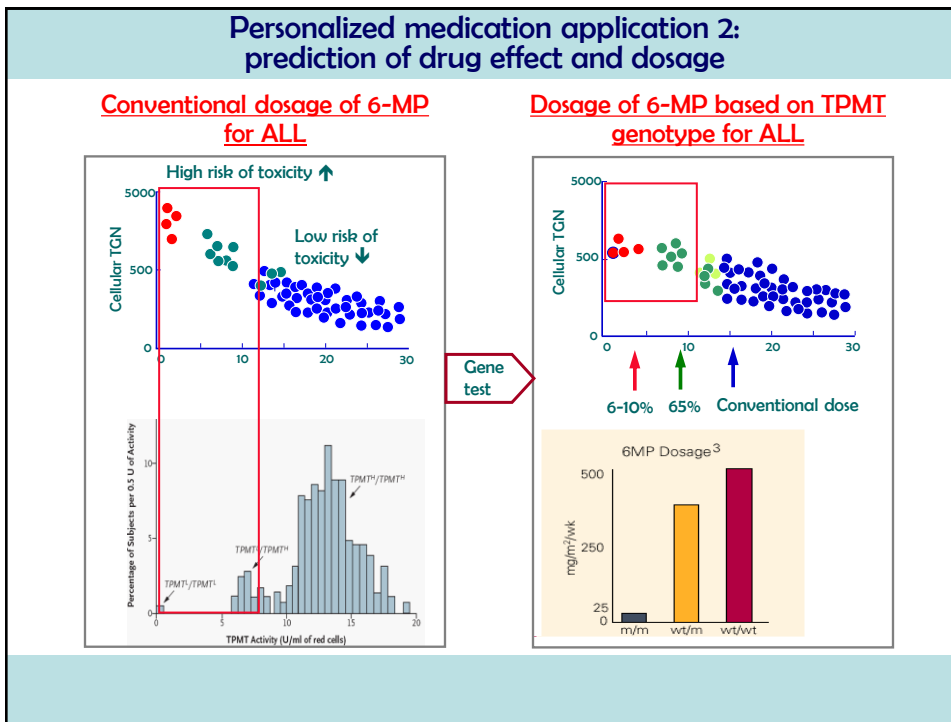
## Personalized medication application 2: prediction of drug effect and dosage

### -Tamoxifen: selective estrogen receptor regulator for breast cancer

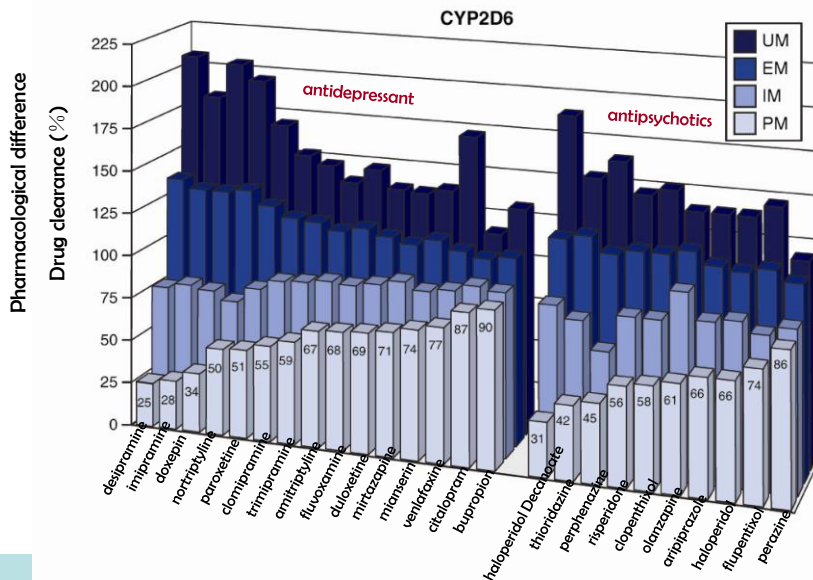


- Poor metabolizer: significant increase in the recurrence of breast cancer, lower survival rate
- When combined with CYP2D6 inhibitor, plasma concentration of highly active metabolite desmethyltamoxifen decrease by 65-75%

- EMA Pharmacovigilance Working Party (PhVWP) warns:
- CYP2D6 poor metabolizer may have lower response to tamoxifen
- CYP2D6 inhibitor combination should be avoided

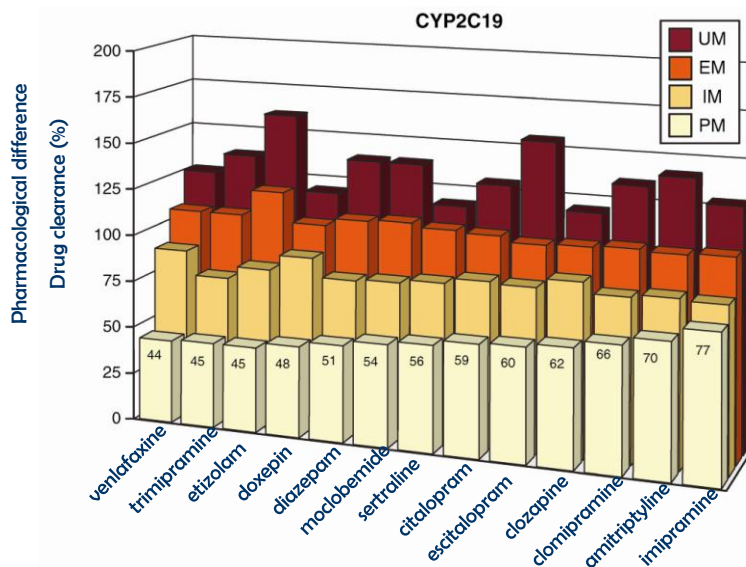


Personalized medication application 2:  
prediction of drug effect and dosage-**Single** factor



Molecular Psychiatry (2012), 1 -- 15

Personalized medication application 2:  
prediction of drug effect and dosage-**Single** factor



Molecular Psychiatry (2012), 1 -- 15

### Personalized medication application 2: prediction of drug effect and dosage-**Multiple factor**

**Factors for warfarin dosing**

● Pharmacogenetic dosing algorithm (international multiple center containing 9 countries and 4043 subjects)

Warfarin dose (mg/d) = [5.6044 - 0.2614(age) + 0.0087(height in cm) + 0.0128 (weight in kg) - 0.8677(VKORC1-1639 A/G) - 1.6974(VKORC1-1639 A/A) - 0.4854(VKORC1 genotype unknown) - 0.5211(CYP2C9\*1/\*2) - 0.9357(CYP2C9\*1/\*3) - 1.0616(CYP2C9\*2/\*2) - 1.9206 (CYP2C9\*2/\*3) - 2.3312 (CYP2C9\*3/\*3) - 0.2188 (CYP2C9 genotype unknowns) - 0.1092(Asian race) - 0.2760(Blacks) - 0.1032(Mixed race) + 1.1816(Enzyme inducers) - 0.5503(Amidarone) ]<sup>2</sup>/7

The International Warfarin Pharmacogenetics Consortium, N Engl J Med 2009;360:753-64.

### Personalized medication application 2: prediction of drug effect and dosage-**Single factor**

**Xiang-Ya PGx dosing algorithm**

Warfarin stable dose (mg/d) = [2.140 - 0.370 × (VKORC1-1639G>A) - 0.332 × (CYP2C9\*3) + 0.324 × (BSA) - 0.004 × (age in decades) - 0.231 × (number of increasing INR drugs) + 0.105 × (smoking habit) - 0.135 × (preoperative stroke history) - 0.108 × (hypertension)]<sup>2</sup>

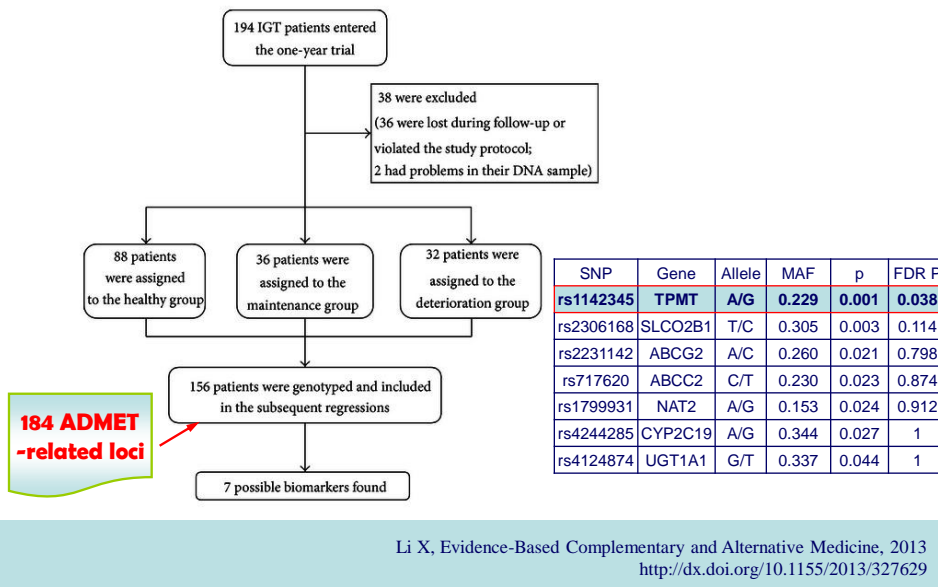
**Algorithms**

The dosing calculated via Xiang Ya dosing algorithm is more close to the actual required for the individual patient from South Central area of China

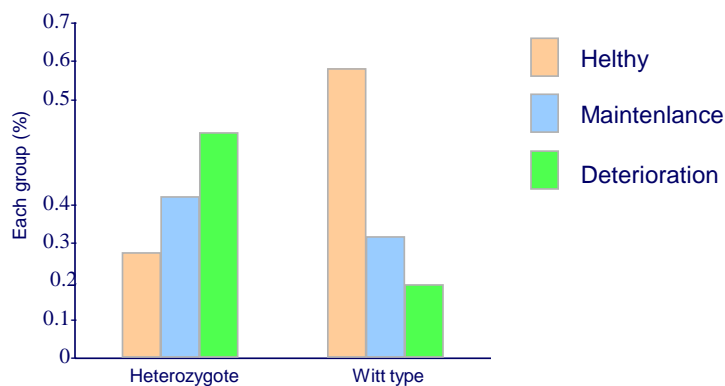


## Personalized medication application 2: prediction of drug effect and dosage

### PGx research of TCM Tianai hypoalcyemic capsule



## Personalized medication application 2: prediction of drug effect and dosage



Percentage of each group in rs1142345 heterozygote (AG) and wild-type (AA) patients.

Personalized medication application 3:  
prediction and prevention of drug toxicity

**HLA-B\*5801 and surrogate SNP rs9263726 are associated with allopurinol-SCAR in mainland Chinese**

● **Allopurinol**

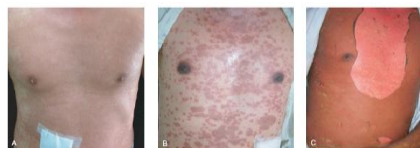
- Primary and secondary gout patients, hyperuricemia

■ **HLA-B\*5801**

Allele positive: 85/90 (94.4%)

■ **Rs9263726 A**

Allele positive: 82/90 (91.1%)



SJS, Stevens-Johnson Syndrome SCAR

SCAR

- erythema multiforme, fever, herpes, purpura, necrosis;
- Mortality 30-40%



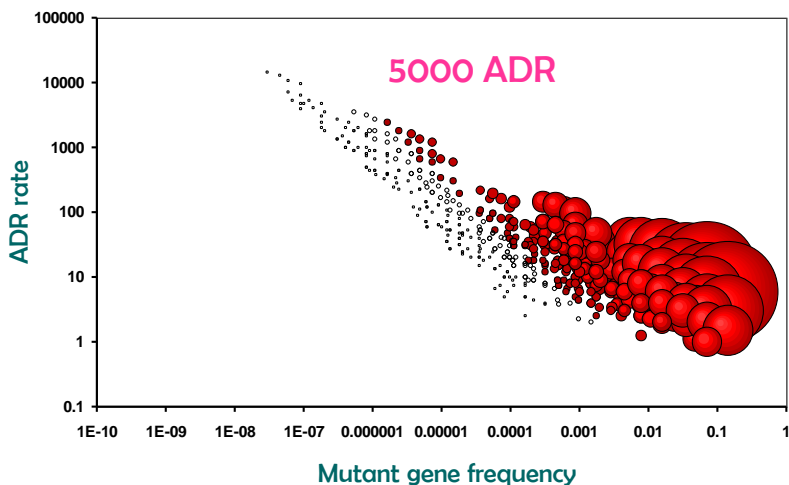
Personalized medication application 3:  
prediction and prevention of drug toxicity

**HLA-B\*5801 and surrogate SNP rs9263726 are associated with allopurinol-SCAR in mainland Chinese**

Patients	Allele positive /total (%)	OR (95% CI)	Sensitivity /specificity	P
<b>HLA-B*5801</b>				
Allopurinol-SCAR	85/90 (94.4%)	399.5 (74.6~2139.4)	94.4% /95.9%	7.10X10 <sup>-26</sup>
Allopurinol-Tolerated	2/49 (4.1%)			
<b>Rs9263726 A</b>				
Allopurinol-SCAR	82/90 (91.1%)	240.9 (49.1~1181.7)	91.1% /95.9%	7.10X10 <sup>-26</sup>
Allopurinol-Tolerated	2/49 (4.1%)			

### Personalized medication application 3: prediction and prevention of drug toxicity

Adverse drug reaction rate is related to mutant gene frequency



- 59% drugs with ADRs are metabolized by polymorphic DMEs

## Guidance for Industry Pharmacogenomic Data Submissions (2005)

Pharmacogenomic data must be submitted to the IND under if ANY of the following apply (these required pharmacogenomic information may influence FDA's decision to these drugs):

1. The test results are used for making decisions pertaining to a specific clinical trial, or in an animal trial used to support safety.
2. A sponsor is using the test results to support scientific arguments pertaining to, for example, the pharmacologic mechanism of action, the selection of drug dosing and dosing schedule, or the safety and effectiveness of a drug.
3. Test results constitute a known valid biomarker for physiologic, pathophysiologic, pharmacologic, toxicologic, or clinical states or outcomes in humans, or the test is a known valid biomarker for a safety outcome in animal studies (CYP2D6).

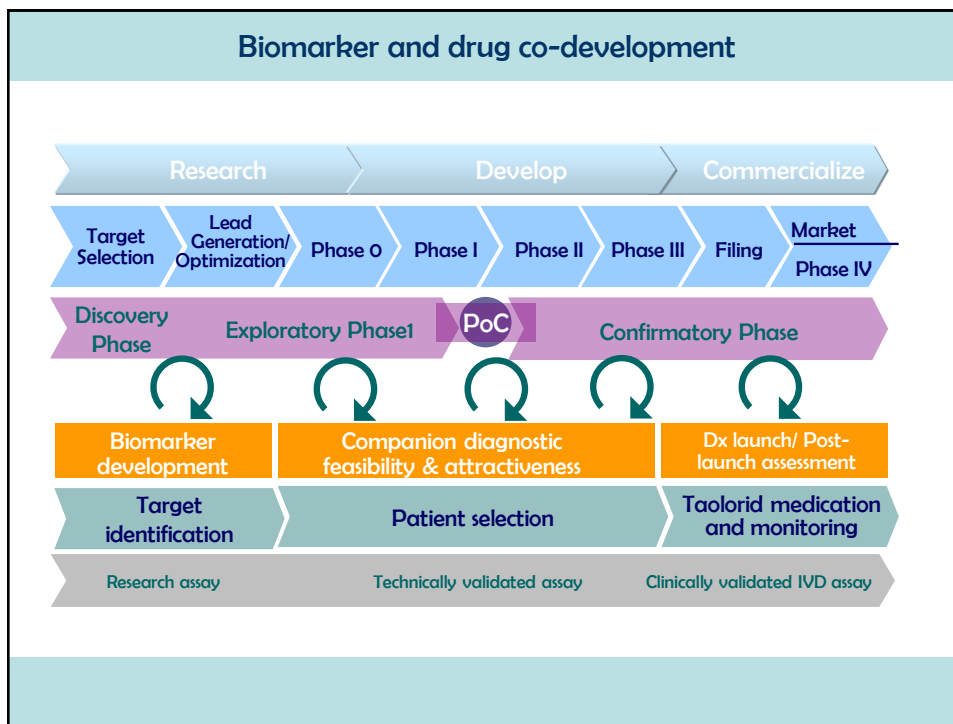
Pharmacogenomic information of most other drugs, which are not part of the investigational or marketing submissions, are not required but such data would be welcome on a voluntary basis.

## Genetic variations in FDA approved drug labels

	Biomarkers	Drugs or representative drugs
1	<i>CCR-5</i>	maraviroc (anti-retroviral agents)
2	<i>EGFR</i> expression	cetuximab, panitumumab, gefitinib
3	Her2/neu overexpression	herceptin
4	Philadelphia chromosome positive reaction	dasatinib
5	<i>C</i> protein deletion (inherited or acquired)	warfarin
6	<i>TPMT</i> variation	azathioprine
7	<i>UGT1A1</i> variation	irinotecan
8	<i>HLA-B*1502</i> allele	carbamazepine
9	<i>UCD</i>	valproic acid
10	<i>CYP2C9</i> mutant	warfarin
11	<i>VKORC1</i> variation	warfarin
12	familial hyperlipoproteinemia LDL receptor deletion or mutant	atorvastatin
13	<i>G6PD</i> deletion	rasburicase
14	<i>HLA-B*5701</i> allele	abacavir

## Genetic variations in FDA approved drug labels

	Biomarkers	Drugs or representative drugs
15	C-KIT expression	imatinib mesylate
16	<i>PML/RAR(α)</i> expression (retinoic acid receptor effective/ineffective)	retinoic acid
17	<i>UGT1A1</i> variation	nilotinib
18	<i>CYP2C19</i> mutant	voriconazole
19	<i>CYP2C9</i> mutant	celecoxib
20	<i>CYP2D6</i> variation	tomoxetine
21	<i>CYP2D6</i> and other variation	fluoxetine hydrochloride
22	gap gene deletion on the long arm of chromosome 5	lenalidomide
23	<i>DPD</i> deletion	capecitabine
24	<i>EGFR</i> expression	erlotinib
25	<i>EGFR</i> expression	gefitinib (head and neck cancer)
26	<i>G6PD</i> deletion	primaquine
27	<i>NAT</i> variation	isoniazide, busulfan
28	Philadelphia chromosome positive reaction	busulfan



### Examples of parallel development projects

#### Anticarcinogen used in subpopulation with specific gene mutant, approved by FDA in 2011

**Xalkori (crizotinib)**

- tyrosine kinase inhibitor, used in the treatment of anaplastic lymphoma kinase (ALK)-positive locally advanced or metastatic non-small cell lung cancer (NSCLC).
- FDA also approved Vysis ALK Break-Apart FISH kit from Abbott, ALK-positive NSCLC patients detected by this kit can be treated with Xalkori.

**Zelboraf (vemurafenib)**

- BRAF inhibitor, used in the treatment of advanced metastatic or unresectable melanoma.
- FDA also approved the first diagnostic reagent used for detection of cobas 4800 BRAFV600 mutant.

## Application and promotion of personalized medicine in Central South University



Mid 1990: Firstly proposed gene-directed telorid therapy in China



2004-4: Personalized Medicine Consultant Center



2006-2: Personalized Medicine Genetic Analysis Center



2010-9: Xiangya Medical Laboratory, CSU



2013-8: National Health and Family Planning Commission  
Personalized Medicine Testing and Training Base

## Xiangya Medical Laboratory

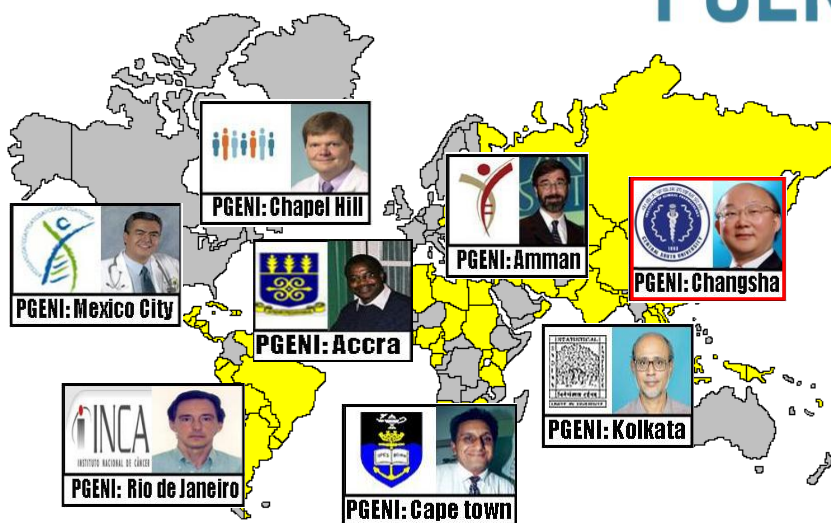


- Founded on the basis of 30-year clinical scientific research and in combination with the strong technical force of the 3 affiliated hospitals of CSU, XYML is a testing organization with third-party arbitration qualification.



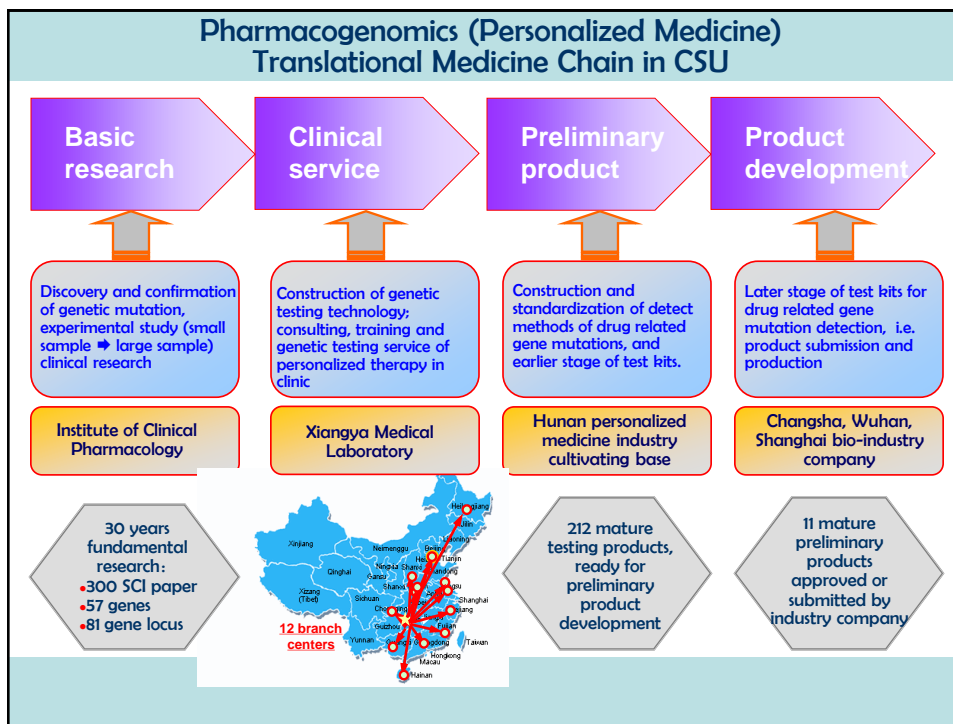
## PGENI local centers

Research centers with high reputation in pharmacogenomics study and application in the world



## Personalized medicine clinical service by Xiangya Medical Laboratory





### The world's first personalized medication gene chip

<http://app1.sfda.gov.cn/datasearch/face3/base.jsp>

Registration number	National food and drug administration 2012 no.3401324
Production unit	Hunan Honghao Gene Bio-technology limited company
Product name	CYP2D6*10、CYP2C9*3、ADRB1(1165G>C)、AGTRI(1166A>C)、ACE(I/D)testing kit(gene chip)
Product standard	VZB/PRC 4686-2012
Product performance, structure and component	Gene chip, wash buffer A, wash buffer B, positive control, negative control, hybridization solution, PCR reaction buffer 1, PCR reaction buffer 2, PCR reaction buffer 3, PCR reaction buffer 4, PCR reaction buffer 5, enzyme 1, enzyme 2, locate reference. Period of validity: A: 2-8℃; B: -20℃, 6 months. Accessory: registration product standard, instruction
Expire date	2016.10.28
Approval date	2012.10.29
Scope of application	Test 5 polymorphism: CYP2D6*10(CYP2D6*1/*1、CYP2D6*1/*10、CYP2D6*10/*10)、CYP2C9*3(CYP2C9*1/*1、CYP2C9*1/*3、CYP2C9*3/*3)、ADRB1(1165G/G、1165G/C、1165C/C)、AGTRI(1166A/A、1166A/C、1166C/C)、ACE(II、ID、DD)。
Specifications	20 runs/kit



## Personalized Medicine in Jiao Tong Univ. Shanghai

The genetic mutation databases of important P450 genes have been constructed for the Chinese population



CYP2D6 database



CYP2C19 database



CYP2C10 database



CYP2E1 database



Sequencing raw data



Alleles and genotypes data



Analysis results data



Primers and conditions data

### news feature

## A great leap forward

After helping to sequence the human genome, Chinese scientists are debating how best to continue toward world p. David C reports

China  
inter  
ject (



### news feature

ing has produced a homogeneous genetic make-up within each population, and the lack of emigration or immigration means that it is easy to construct large family pedigrees. These features have made Iceland and Finland hotspots for mapping the chromosomal location of genes predisposing to disease. "We have several Icelands and several Finlands," says Lin He of Shanghai Jiao Tong University, who coordinates a bank for blood, tumour and cerebrospinal fluid samples collected from populations across China.

JiaoTong University have compiled the world's largest sample band related to neuropsychiatric disease in a joint project with the Shanghai Institutes of Biological Sciences.

Xu Zhi-Hong, president of Peking University. "This is our challenge."

It is a challenge that both the government and researchers are taking on. In 1998, the government set up a programme alongside the HGP sequencing effort, which has estab-

### Nature

1 March 2001:10-12

### news feature

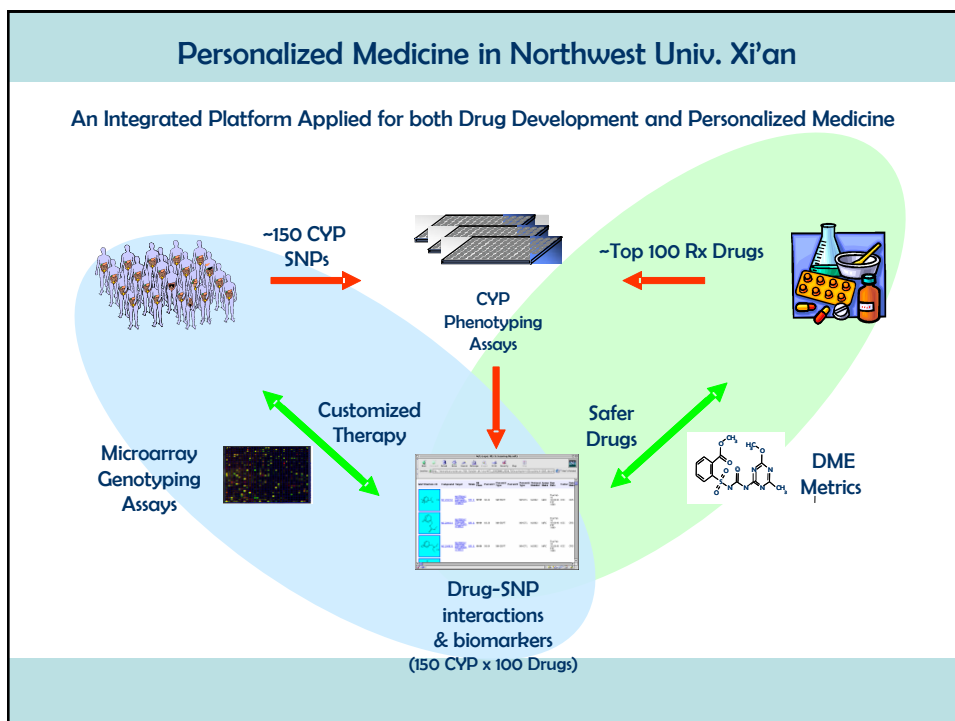
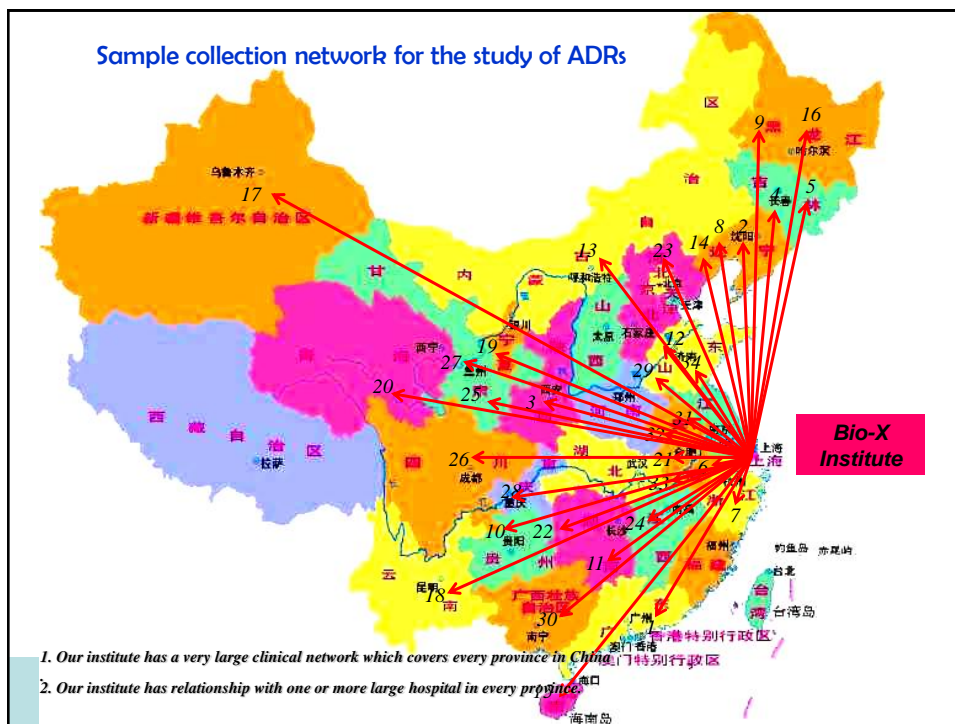
Tong University, meanwhile, He claims to have compiled the world's largest sample bank related to neuropsychiatric disease, in a joint project with the Shanghai Institutes of Biological Sciences.

### Rising stars

Studies of samples taken from Chinese populations are already yielding advances. The first success came in 1998, when a team led by Jia-hui Xia of the National Laboratory of Medical Genetics of China in Changsha, Hunan province, identified a gene for a form of neurological deafness. Then, last year, researchers led by He mapped the gene for brachydactyly type A-1 — a disease in which joints of the fingers are missing, mis-



And the pace looks set to pick up. Last month, researchers led by Yan Shen of the



## Personalized Medicine in Northwest Univ. Xi'an

### More CYP and CYP SNP Reagents

CYP	Companies							CYP	Companies						
	Host SNP ID	Lifegen	Cypex	Invitrogen	BD Gentest	BD Gentest	BD Gentest		Host SNP ID	Lifegen	Cypex	Invitrogen	BD Gentest	BD Gentest	BD Gentest
1A2	Prototype	Yes	Yes	Yes	Yes	Yes	Yes	Prototype	Yes	Yes	Yes	Yes	Yes	Yes	
2A6	Prototype	Yes	Yes				Yes	Prototype							
2C9	Prototype	Yes	Yes	Yes	Yes	Yes	Yes	E92D, I331V	Yes						
	R144C, R159L	Yes	Yes	Yes	Yes	Yes	Yes	R433W, I331V	Yes						
2D6	Prototype	Yes	Yes	Yes	Yes	Yes	Yes	R115Q, I331V	Yes						
	R299C	Yes	Yes					W120R, I331V	Yes						
	S480T	Yes	Yes					P144H, I331V	Yes						
	P345	Yes	Yes					E227L, I331V	Yes						
	N166D	Yes						R150H, I331V	Yes						
	L91M	Yes						R410C, I331V	Yes						
	H94R	Yes						L17P, I331V	Yes						
	H324P	Yes						I19L, I331V	Yes						
	G42R	Yes						R442C	Yes						
	G159R	Yes						R329H	Yes						
	T107I	Yes						S51G	Yes						
	R23C	Yes						D360N	Yes						
	A85V	Yes						A161P	Yes						
	I297L	Yes						A74T	Yes						
	R343G	Yes						F168L	Yes						
	I507F	Yes						W212C	Yes						
	V7M	Yes						E122A	Yes						
	Q151E	Yes						Prototype	Yes	Yes	Yes	Yes	Yes	Yes	
	V136M	Yes						S222P	Yes						
	V338M	Yes						M445T	Yes						
R440H	Yes						I118V	Yes							
E410K	Yes						P218R	Yes							
A237S	Yes						G56D	Yes							
V11M	Yes						R130Q	Yes							
R201H	Yes						V170I	Yes							
R30H	Yes						D174H	Yes							
G373S	Yes						T353M	Yes							
R173C	Yes						L373F	Yes							
L213P	Yes						P416L	Yes							
E413Q	Yes						L15P	Yes							
P325L	Yes						R152Q	Yes							
G212E	Yes						T185S	Yes							
L213S	Yes						F189S	Yes							
L421P	Yes						L295P	Yes							
							P467S	Yes							

## Personalized Medicine in Northwest Univ. Xi'an

### Lifegen Cypro-screen™

A Broad Range of Cytochrome P450s and Over Hundred CYP SNPs

High-throughput Analysis Service for Your Drug Candidates

<ol style="list-style-type: none"> <li><b>CYP Reaction Phenotyping</b> <ul style="list-style-type: none"> <li>CYP isoform identification</li> <li>Evaluate the drug metabolic differences caused by CYP SNPs</li> </ul> </li> <li><b>CYP Mediated Drug-Drug Interaction by High-throughput Fluorometric Assay and LC-MS</b> <ul style="list-style-type: none"> <li>CYP-based inhibition assay</li> <li>CYP SNPs-based inhibition assay</li> </ul> </li> </ol>	<p><b>Enzymes Available</b></p> <p>12 CYP prototype: 1A1 1A2 2A6 2A13 2B6 2C8 2C9 2C19 2D6 2E1 3A4 3A5</p> <p>10 SNPs of 2C9: R144C R150H L191 H251R E272G R335W I359L I359T D360E P489S</p> <p>20 SNPs of 2C19: A161P L17P I19L S51G D360N I331V E92D E122A F168L W120R W212C M74T R132Q R144H R150H R329H R410C R433W R442C P227L</p> <p>55 SNPs of 2D6: V7M V11M V136M V374M V338M R25W R26H R26C R173C R201H R329L R365H R296C R343G R380H P345 P325L P430L G42R G169R G212E G479R L91M H94R T107I F1201 A237S A122S A85V A90V A300G Q151E W152G E153K E156A N166D S168A S486T S311L L213P L213S L231P M279K I297L H324P E334A I369T G373S E410K E418Q H478V P345-S486T R296C-S486T</p> <p>19 SNPs of 3A4: S222P M445T I118V P218R G56D R130Q V170I D174H T363M L373F P416L L15P R162Q T185S F189S L293P P467S</p>
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# Personalized Medicine in Tsinghua University Univ. Beijing

## Mutation Screening for Neonatal Deafness

Used for the detection of 9 hot mutations of four genes, Mitochondrial 12S rRNA, GJB2, GJB3 and PDS, which are in conjunction with hereditary hearing loss. Especially, the 1555 A>G and 1494 C>T mutations at Mitochondrial 12S rRNA could be detected. Exposure to aminoglycoside antibiotics such as streptomycin can lead to severe-to-profound hearing loss when patient carry 12S rRNA the mutations.



Kit

SFDA

BioMixer

SlideWasher

LuxScan

Software

Human Mutation, 29:306-14, 2008

## The Screening of the Deaf in Beijing

6 | 北京日报 2011年3月4日 星期五

### 两万多聋人免费查病因

基因检测可查出是否遗传下一代 医生建议健全人也可进行检测

昨天是全国“爱耳日”，本市聋人人群致聋基因筛查项目筹备启动，两万多名聋人聋家人可免费接受筛查，聋家人可查出患病基因，医生将提供其遗传病诊断及治疗建议。

#### 抽血两三毫升可查致聋原因

北京市聋人协会日前启动本市聋人人群致聋基因筛查项目。该项目旨在通过基因检测，找出聋人致聋的遗传原因。筛查过程简单，只需抽取两到三毫升血液，即可进行检测。筛查结果将帮助聋人了解自己的致聋原因，并为聋家人提供遗传病诊断及治疗建议。

#### 三色卡告聋人致聋原因

北京市聋人协会日前启动本市聋人人群致聋基因筛查项目。该项目旨在通过基因检测，找出聋人致聋的遗传原因。筛查结果将帮助聋人了解自己的致聋原因，并为聋家人提供遗传病诊断及治疗建议。



图为聋人儿童在北京市聋人协会进行听力检测。

#### 健全人也可进行检测

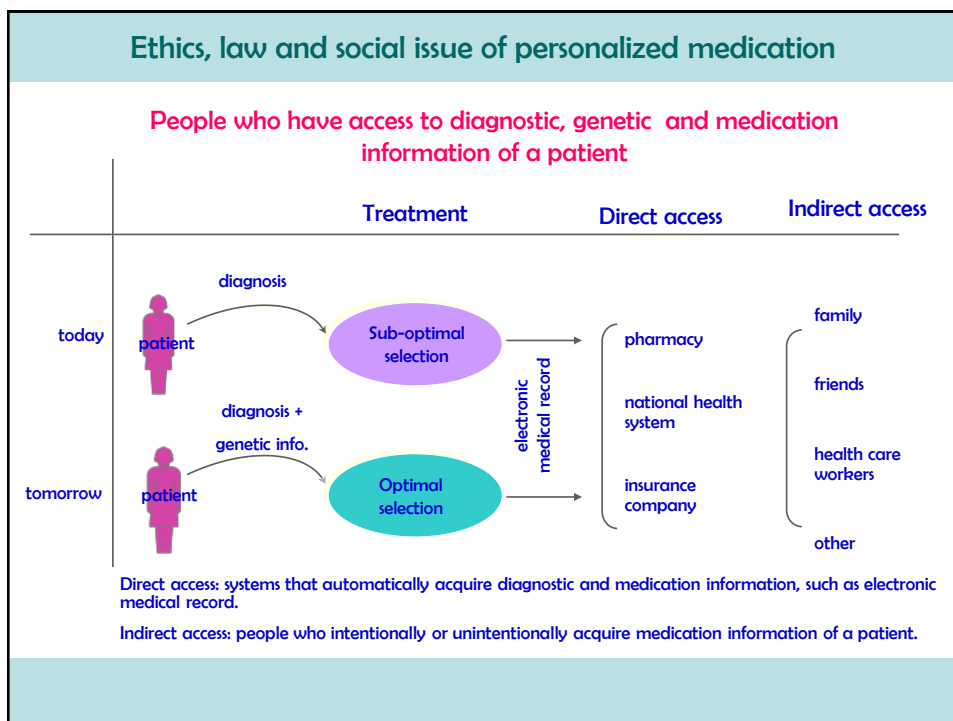
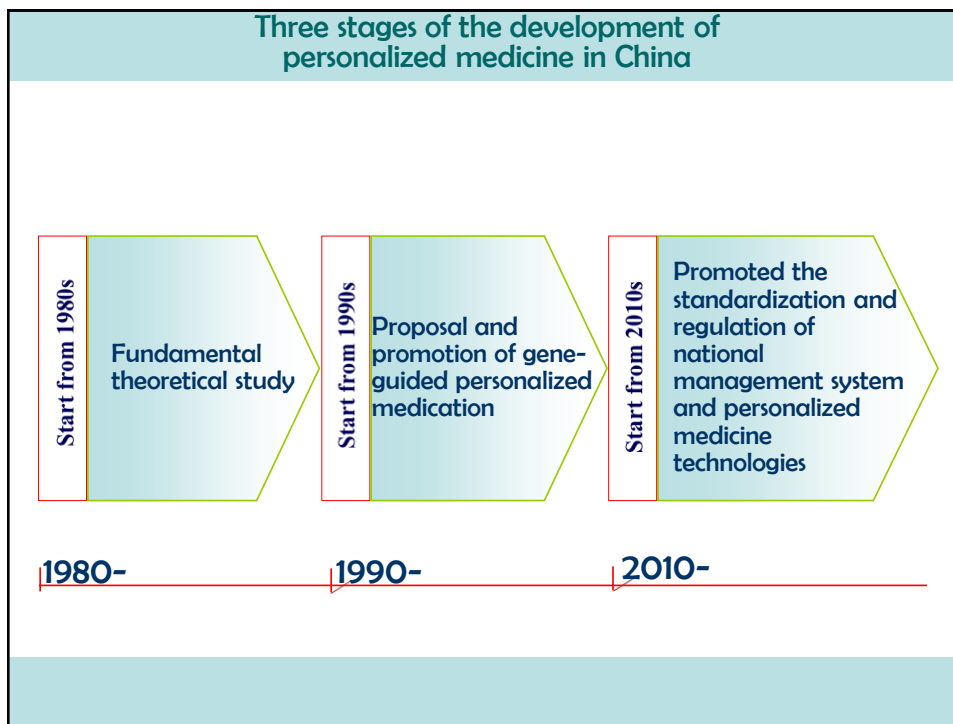
除了聋人，聋家人也可进行检测。医生建议健全人也可进行检测，以了解自己的遗传状况。筛查结果将帮助聋家人了解自己的遗传病诊断及治疗建议。

#### 本市每年至少150名新生儿听力障碍

北京市聋人协会日前启动本市聋人人群致聋基因筛查项目。该项目旨在通过基因检测，找出聋人致聋的遗传原因。筛查结果将帮助聋人了解自己的致聋原因，并为聋家人提供遗传病诊断及治疗建议。



图为北京市聋人群致聋基因筛查项目启动仪式暨“3.3爱耳日”主题宣传活动。



## Ethical, legal and social issues of personalized medication

### (1) Protect individual's genetic privacy

- Individual's genetic information should not be acquired without consent.
- Individual's decision of not to know his genetic information for any reason should be respected.
- Procedure and regulation of rational acquisition, transport and reservation of individual's genetic information should be assigned by gene testing units.

### (2) Respect individual's right to knowledge

- The aim, purpose, procedure, outcome and risk of genetic test should be informed to, and agreed by, the participant.
- Testing outcomes with individual's consent should be given to participant veritably. Risk evaluation of disease and positive significance in medication should be explained to participant scientifically, comprehensively and positively, to help patient understand his genetic information correctly and positively.

### (3) Against genetic discrimination

- Individual's genetic information might give rise to genetic discrimination. To protect people against genetic discrimination, firstly the privacy of an individual's genetic information should be protected, secondly advanced legal system and social moral system should be established.

Although genetic testing in personalized medication is not aimed for genetic disease detection and prediction, potential risk in society, psychology and economy might rise when gene mutation, potential disease susceptibility and prognosis information are known. Therefore, informed consent form should be signed by participant before sample collection, except for somatic mutation detection.

## National missions undertaken by CSU

Guided by the official letter of approving the regulation and management of personalized medicine testing by the general office of the Ministry of Health, ([2013]No195), the bureau of medical policy and management, National Health and Family Planning Commission has convened the first conference of personalized medicine committee in Beijing on May 13 2013.

- As a main member of the personalized medicine committee, the Ministry of Health, take part in the standardized management of personalized medicine molecular testing laboratory in China;
- Set the standard of personalized molecular testing laboratory;
- Draw up the guidelines of personalized medication-related gene (DMEs, transporters, receptors) molecular testing projects;
- National Health and Family Planning Commission Personalized Medicine Testing and Training Base (exclusive);
- National Pilot Standard Laboratory of Personalized Medicine Molecular Testing (one of the three).



*Acknowledgment*



*Thank you for your attention*