Pharmacogenetics

Percision Medicine



Dr. Mohammed Al-Sbou Professor of Clinical Pharmacology Faculty of Medicine-Mutah Uni The human genome project has led to an explosion of genetic information that is freely available to identify polymorphisms that may determine drug response

Ex=> PCR, Molecular techniques in general

Advances in molecular genetics and genotyping technologies during the last two decades have led to identification of many polymorphisms in phase I and phase II drug metabolising enzymes, drug targets, and in drug transporters

@ Imperical therapy => giving the drug based on it's genetity

E Polymorphism may occur in receptors and ph. kinetics, ph. dynamics.

, Varia (Joyrs in single base or Whole Sequence in DNA. Individual Variation in Response to Drugs

- How individuals in a population are expected to respond to a fixed dose of drug?
- Inter-individual variability:
 - Some show less than usual response
 - Most show usual response
 - Others show more than usual response





Factors Determine Response to Drugs & it's very new concept.

Pharmacogenetics/Pharmacogenomics

- Pharmacogenetics: is study of variation in drug response due to heredity & is used in relation to genes determining drug metabolism
- Pharmacogenomics is a more general term; it refers to research area that comprises all genes in the human genome that may determine drug response

Benefits of Pharmacogenetics/Pharmacogenomics The concept <u>"The right medicine to the right</u> patient" is the basis of pharmacogenetics (personalised or individualised medicine) Ultimate goals are to improve clinical therapeutic outcome by: - Increasing drug efficacy - Increasing safety of drugs e.g. reducing incidence of ADRs

8

Personalised or Individualised Medicine



Pharmacogenomic approach to personalized medicine. Drug therapy is chosen for each patient based on their <u>particular genetic profile</u>



Polymorphisms can occur in any gene that encode:

- Drug metabolising enzymes
- Drug transporters
- Drug targets and receptors

<u>Genetic polymorphisms of drug</u> <u>metabolising enzyme genes</u>

The majority of phase I and phase II drug. metabolising enzymes are polymorphic ■ The cytochrome P450 (CYP) enzymes are the most important group of phase I enzymes Polymorphisms in cytochrome P450 genes can cause enzyme products with abolished or reduced or increased enzyme activity

Cytochrome P450 enzymes

All genes that encode for families 1-3 are polymorphic & their capacity to metabolise drugs depends on the functional importance and frequency of variant alleles

B Cyp4502 C9 - Subarbarine Metabolism

& Prequency of Allels & Common or Rare.

CYP2D6 = Metabolism of 25% of All drugs sortium CYP2D6 = Metabolism of 25% of All drugs sortium Exceptor In Visitor Malgesice.

<u>CYP2D6</u>

CYP2D6 contributes to metabolism of large of medications about 25% of all drugs, including:
 Antidepressants (TCAs, SSRIs)
 Antiarrythmics
 Analgesics

CYP2D6 Phenotypes

- Poor metabolisers(PM): lack functional enzyme
- Intermediate metabolisers (IM): carry two alleles that cause reduce activity - needs by reduce
- Extensive metabolisers (EM): have two normal alleles -> Uovna (Netabolism
- Ultra-rapid metabolisers (UM): multiple gene copies = Migh Metabolism
 - Normal dose => Non transponders - Needs Higher dose

Poor metabolisers can experience adverse effects when treated with standard dose
 Ultra-rapid metabolisers require high doses of drugs

GeneSight® Psychotropic COMBINATORIAL PHARMACOGENOMIC TEST enesiaht Questions about report interpretation? Patient, Sample DOB: 7/22/1984 Contact our Medical Information team Order Number: 219 05.891.9415 Report Date: 8/5/2020 El medinfo@essureshealth.co Classian fiample Clinician Reference 1458CIP PATIENT GENOTYPES AND PHENOTYPES PHARMACOKINETIC GENES CYP1A2 Extensive (Normal) Metabolizer CYP2D6 Poor Metabolizer *1/*1 *10/*10 This genotype is most consistent with the extensive inormali-CYP2D6*10 allele enzyme activity: Reduced metabolizer phenotype. CYP2D6*10 allele enzyme activity. Reduced This grnotype is most consistent with the poor metabolizer CYP2B6 Extensive (Normal) Metabolizer phenotype. This patient may have reduced enzyme activity as *1/*1 compared to individuals with the normal phenotype. CYP2B6*1 allele enzyme activity: Normal CYP2B6*1 allele enzyme activity. Normal CYP3A4 Extensive (Normal) Metabolizer 1/11 This genotype is most consistent with the extensive (normal) CYP3A4*1 able enzyme activity: Normal metabolizer phenotype. CYP3A4*1 allele enzyme activity. Normal This genotype is most consistent with the extensive (normal) CYP2C19 Ultrarapid Metabolizer metabolizer phenotype. *17/*17 CYP2C19*17 allele enzyme activity: Increased UGT1A4 Extensive (Normal) Metabolizer CYP2C19*17 allele enzyme activity. Increased *1/*1 This genotype is most consistent with the ultrarapid metabolizer UGT1A4*1 allele enzyme activity. Normal phenotype. This patient may have increased enzyme activity as UGT1A4*1 allele enzyme activity. Normal compared to individuals with the normal phenotype This genotype is most consistent with the extensive (normal) metabolizer phenotype. The patient is expected to have normal CYP2C9 Intermediate Metabolizer enzyme activity. *1/*2 CYP2C9"1 allele enzyme activity: Normal UGT2B15 Intermediate Metabolizer CYP2C9*2 allele enzyme activity: Reduced +2/2

This genotype is most consistent with the intermediate metabolizer phenotype. This patient may have reduced enzyme activity as compared to individuals with the normal phenotype.

> This genotype is most consistent with the intermediate metabolizer phenotype. This patient may have reduced enzyme activity as compared to individuals with the normal phenotype.

UGT2B15*2 allele enzyme activity: Reduced

UGT2B15*2 allele enzyme activity: Reduced

Pharmacokinetic Genes

Pharmacokinetic genes provide information on the metabolism of medications.

Activate Wi



HLA-A*3101

T/T

This patient is homozygous for the short promoter polymorphism of the serotonin transporter gene. The short promoter allele is reported to decrease expression of the serotonin transporter compared to the homozygous long promoter allele. The patient may have a decreased likelihood of response to selective serutonin reuptake inhibitors due to the presence of the short form of the gene and may benefit from medications with an alternative mechanism of action.

HTR2A

Increased Sensitivity

GIG

dife

This individual is homozygous variant for the G allele of the -1438G>A polymorphism for the Serotonin Receptor Type 2A. They carry two copies of the G allele. This genotype has been associated with an increased risk of adverse drug reactions with certain selective serotonin reuptake inhibitors. This patient does not carry the HLA-8*1502 allele or a closely related *15 allele. Absence of HLA-8*1502 and the closely related *15 alleles suggests lower risk of senous dermatologic reactions including toxic epidermal necrolysis (TEN) and Stevens-Johnson syndrome (SJS) when taking certain mood stabliggers.

Higher Risk

This patient is homozygous for the T allele of the rs1061235 A>T polymorphism indicating presence of the HLA-A*3101 allele or certain HLA-A*33 alleles. This genotype suggests a higher risk of serious hygersensitivity reactions, including Stevens-Johnson syndrome (SJG), toxic epidermal neorolysis (TEN), maculopapular eruptions, and Drug Reaction with Eosinophilia and Systemic Symptoms when taking certain mood stabilizers.

Pharmacodynamic Genes

Pharmacodynamic genes provide information on how DNA may impact response to some medications.

Dit is adviced to make a genotypong test to make save that the drag works. (if possible)

Depression

- Tricyclic antidepressants are metabolised by CYP2D6
- Disposition of nortriptyline is related to number of active CYP2D6 alleles and
- Dose required to obtain same plasma drug concentrations varies between subjects with different CYP2D6 phenotypes

Ultra-rapid metabolisers needed a 10-fold larger dose of nortriptyline than poor metabolisers to achieve the same plasma concentration

Ultra-rapid metabolisers require 500 mg of doses compared to 50 mg in poor metabolisers

Genetic polymorphisms of CYP2D6 gene may be associated with ADRs and clinical response

to antidepressants

- 30% of patients with ADRs to antidepressants were PMs, poor Metabolizers
- High incidence of <u>UMs</u> among <u>non-responders</u>
 (20%)

CYP2C9

- CYP2C9 metabolises a wide range of drugs
 Including drugs with narrow therapeutic indices such as:
 - Warfarin
 - Phenytoin

 Non-steroidal anti-inflammatory drugs (NSAIDs) including ibuprofen, diclofenac and celecoxib

Warfarin and Bleeding

- Warfarin is one of the most widely prescribed
 oral anticoagulant drugs
- It is used for:
 - Prophylaxis and treatment of venous thromboemolism
 - Treatment of deep vein thrombosis (DVT)
 - Atrial fibrillation (AF)
 - In patients with prosthetic heart valves

Warfarin and Bleeding

The main complication of warfarin therapy is haemorrhage -> Bleeding

Genetic polymorphisms in CYP2C9 gives rise to variants with altered enzymes activity Polyce Medere Two allelic variants CYP2C9*2 (Arg144Cys) and CYP2C9*3 (Ile359Leu) show 12% and 5% of enzyme activity of the wild type CYP2C9*1 allele, respectively, and are associated with decreased warfarin dose requirements & increased risk of bleeding

Quartarin clouic = genotyping at your genes to know which brag suits you!

Peptic Ulcer

- Proton pump inhibitors (PPIs) are used for treatment of gastric acid related diseases such as peptic ulcers, gastro-esophageal reflux disease (GERD) & in combination with antibiotics (amoxicillin & clarithromycin) for eradication of *Helicobacter pylori* (Hp)
- **CYP2C19** metabolises several PPIs including omeprazole and lanzoprazole
- Plasma concentrations of **omeprazole**, depend on patient's **CYP2C19 phenotype**

AmpliChip CYP450 Array

The AmpliChip CYP450
 Test provides comprehensive
 detection of gene variations
 including deletions and
 duplications for the
 CYP2D6 and CYP2C19
 genes



Genetic Polymorphisms of Drug Metabolising Enzyme Genes

With respect to phase II enzymes, the most important polymorphisms occur in *N*-acetyltransferase-2 (NAT-2) and thiopurine methyltransferase (TPMT)
 NAT-2 is involved in the metabolism of isoniazid and sulphamethoxazole

Acetylation

- Most individuals are either <u>rapid</u> or <u>slow</u> <u>acetylators</u>, but proportion varies between races
- The percentage of slow acetylators: " " and a
 - 90% in North African
 - 50% in Caucasian
 - 10% in Asian populations

Thiopurine S-methyltransferase (TPMT) TPMT catalyzes methylation of **thiopurine** drugs such as 6-mercaptopurine & azathioprine These drugs are commonly used in treatment of acute lymphoblastic leukaemia (ALL), autoimmune diseases, inflammatory bowel diseases, in organ & tissue transplantation Clinical testing for TPMT genetic polymorphisms is available

- It has been shown that:
 - 90% of population exhibit high TPMT activity
 - 10% show intermediate activity
 - 0.3% have low or absent enzyme activity

"I ormal dose? toxicity

Genetic Polymorphisms in Drug Transporters Transporters are membrane proteins that play crucial role in absorption, distribution & elimination of drugs Genetic polymorphisms can occur in transport proteins & may contribute to inter-individual variation in drug response MDR1 (multi-drug resistant) P-glycoprotein-Digoxin -> HF drug Serotonin transporter-antidepressant response

Genetic Polymorphisms in Drug targets and Receptors

Drug target genes including those coding for receptors, ion channels and specific enzymes are subject to genetic polymorphisms B2-adrenergic receptor: B2 agonist (salbutamol) Angiotensin converting enzyme (ACE): ACE inhibitors (lisinopril) Vitamin K epoxide reductase complex لا بت مرجعة جزاناج , و الكن التحلة جرن (VKORC): Warfarin 33

Practical Points

 Genetic is an important factor responsible for failure to therapy & occurrence of adverse drug reactions

The goal of PGx is to maximize efficacy & minimize toxicity, based on individual's genetic composition

Individual variation in response to drug (some may benefit, other fail to respond to treatment, others may develop adverse effects)