

Bumrungrad Personalized Medication Review

Test Data

HN 987654321



# Personalized Medication Review

Name Test Data

HN 987654321

Date of birth: 1<sup>st</sup> Jan 1989

Collected Date: 1<sup>st</sup> Aug 2024

Reported Date: 7<sup>th</sup> Aug 2024

## Personalized Medicine report

### Genetic Testing for Drug Allergy

#### Pharmacogenomics for *HLA-A* and *HLA-B* \*\*

In reference to the current database of drug-gene association, your results of *HLA-A* and *HLA-B* testing, which are related to drug allergy suggest that,

- You have a **normal/ low risk of drugs induced severe cutaneous reaction if the following medications are used,**
  - (1.) Carbamazepine (antiepileptic)
  - (2.) Oxcarbazepine (antiepileptic)
  - (3.) Co-trimoxazole (antibiotic)
  - (4.) Dapsone (antibiotic)
  - (5.) Abacavir (antiviral)
  - (6.) Nevirapine (antiviral)
  - (7.) Allopurinol (anti-hyperuricemia)

Therefore, the medications can be used per standard dosing guideline. However, if you are a naive person, other common reactions, such as hives or rashes should also be monitored.



Bumrungrad  
International  
HOSPITAL

# Personalized Medication Review

Name Test Data

HN 987654321

Date of birth: 1<sup>st</sup> Jan 1989

Collected Date: 1<sup>st</sup> Aug 2024

Reported Date: 30<sup>th</sup> Aug 2024

## Personalized Medicine report

### Genetic Testing for Drug Metabolism Pharmacogenomics panel (11 genes)

#### Current medications

**Azathioprine (Imuran®)** is a medicine used for treatment of irritable bowel syndrome (IBS). It is metabolized via TPMT (Thiopurine-S-methyltransferase) and NUDT15 (Nucleoside diphosphate-linked moiety X motif 15) enzyme. Since you have a normal function of *TPMT* and *NUDT15* gene, the medication can be used as per standard dosing guideline. However, its side effects, such as nausea, vomiting, low white blood cell count and increase risk of infection should be observed. Moreover, blood testing will be routinely monitored by the doctor while using this medication.

Regarding your **Mesalazine (Pentasa®)** and supplements which are **MTV** and **Omega-3**, the response including the efficacy and side effects of them could not be predicted from the genetic testing for drug metabolism (PGx panel-BH MedGene).

Bring the lists of your current medication and supplements to the hospital every visit, so that the doctor and/or a pharmacist can recheck the possible side effects and interaction among them.

Note: The prediction is based on simply the genetic profile. Other factors such as patient's current condition, kidney and liver function, and drug interaction among drugs, etc. may also be considered by the doctor if your medication treatment would be adjusted.

Reviewed by R. Ph 1

Pharmacy License No. 87654

Approved by: R. Ph 2

Pharmacy License No. 45678



ห้องปฏิบัติการเภสัชพันธุศาสตร์  
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PHARMACOGENOMICS AND PERSONALIZED MEDICINE REPORT

Name-Surname :		Age :		Gender :	
Hospital Number :		Hospital/Source :			
Specimen :		Phone/Fax :			
Ethnicity :		Requested date :			
Physician :		Reported date :			

HLA genotyping test for Carbamazepine profile (TEST CODE 410074)

HLA allele :		
Possibility genotype <sup>†</sup> :	<i>HLA-A*02:07/02:03</i>	<i>HLA-B*46:01/55:02</i>
Drug-Gene	Genotype	Therapeutic recommendation
Abacavir ( <i>HLA-B*57:01</i> )	Negative <i>HLA-B*57:01</i>	Abacavir can be used per standard dosing guideline with a normal/low risk of Abacavir hypersensitivity.
Allopurinol ( <i>HLA-B*58:01</i> )	Negative <i>HLA-B*58:01</i>	Allopurinol can be used per standard dosing guideline with a normal/low risk of Allopurinol-induced MPE, SJS, TEN, DRESS.
Carbamazepine ( <i>HLA-B*15:02/HLA-B75 family/HLA-A*31:01</i> )	Negative <i>HLA-B*15:02</i> Negative <i>HLA-A*31:01</i>	Carbamazepine can be used per standard dosing guideline with a normal/low risk of Carbamazepine-induced MPE, DRESS and SJS/TEN.
Co-trimoxazole ( <i>HLA-B*13:01, HLA-B*13:02</i> )	Negative <i>HLA-B*13:02</i> Negative <i>HLA-B*13:01</i>	Co-trimoxazole can be used per standard dosing guideline with a normal/low risk of Co-trimoxazole-induced SJS, TEN, DRESS.
Dapsone ( <i>HLA-B*13:01</i> )	Negative <i>HLA-B*13:01</i>	Dapsone can be used follow standard guideline with a lower risk of Dapsone-induced drug hypersensitivity.
Nevirapine ( <i>HLA-B*35:05</i> )	Negative <i>HLA-B*35:05</i>	Nevirapine can be used per standard dosing guideline with a normal/low risk of Nevirapine-induced skin rash.
Oxcarbazepine ( <i>HLA-B*15:02</i> )	Negative <i>HLA-B*15:02</i>	Oxcarbazepine can be used per standard dosing guideline with a normal/low risk of Oxcarbazepine -induced SJS, TEN.

Notes: Abbreviation; CBZ = carbamazepine, OxCBZ = oxcarbazepine, SJS = Steven-Johnson's syndrome, TEN = toxic epidermal necrolysis, MPE = maculopapular exanthema DRESS = drug reaction with eosinophilia and systemic symptoms

<sup>†</sup>The *HLA-A\*XX:XX* and *HLA-B\*XX:XX* allele were determined by PCR-SSOP method which is intermediate resolution *HLA* typing. This report is based on The Common and Well-Documented (CWD) database.

Interpreted by 

# PERSONALISED MEDICATION REPORT

For

Date of birth:	myDNA ID:	Pathology No:	Sample type:
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Collected:	Received:	Reported:	Doctor:
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# Personalised Medication Report

for

Unless instructed by their doctor, patients are advised not to alter the dose or stop any medications.

Name:	<input type="text"/>	DOB:	<input type="text"/>	Collected:	<input type="text"/>
Address:	<input type="text"/>	myDNA ID:	<input type="text"/>	Received:	<input type="text"/>
Doctor:	<input type="text"/>	Pathology No:	<input type="text"/>	Reported:	<input type="text"/>

Sample type and quality: **Buccal. The sample quality was assessed and deemed to be satisfactory according to the laboratory's acceptance criteria.**

Clinical Notes:



## REPORT SUMMARY

### MEDICATIONS OF INTEREST OVERVIEW

MEDICATIONS THAT DO NOT HAVE PRESCRIBING CONSIDERATIONS BASED ON myDNA TEST

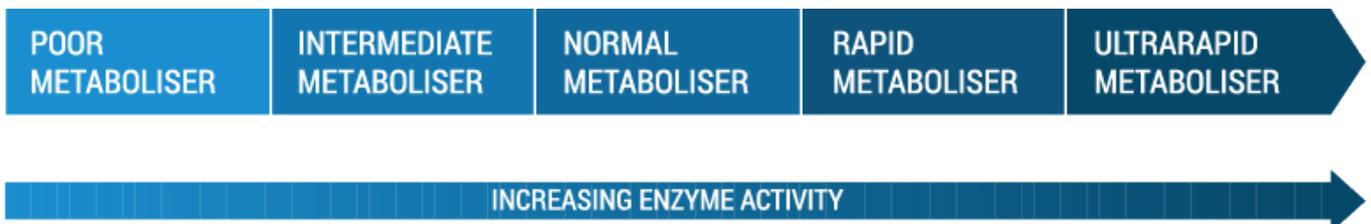
mesalazine, multivitamin, omega 3, tobacco smoke

### GENETIC TEST RESULTS OVERVIEW

GENE	GENOTYPE	PREDICTED PHENOTYPE	GENE	GENOTYPE	PREDICTED PHENOTYPE
CYP2D6	*1/*1	Normal metaboliser	CYP3A5	*3/*3	Poor metaboliser
CYP2C19	*1/*1	Normal metaboliser	SLCO1B1	*1/*1	Normal transporter function
CYP2C9	*1/*2	Intermediate metaboliser	CYP2B6	*1/*6	Intermediate metaboliser
VKORC1	AG	Moderately reduced VKORC1 enzyme level	OPRM1	AA	Normal mu opioid receptor expression
CYP1A2	*1F/*1F	Ultrarapid metaboliser (with inducer present)	ABCG2 (rs2231142)	CC	Normal transporter function
CYP3A4	*1/*1	Normal metaboliser			

Detailed interpretations of genetic test results are provided in the *pharmacogenomic interpretation* section below.

The following diagram provides the range of enzyme activity predicted by the myDNA test.



### OTHER INFORMATION

Please note that the myDNA medication test does not cover TPMT genotyping, used to predict the dose and risk of myelosuppression of thiopurine drugs (azathioprine and 6-MP). TPMT genotype testing can be ordered through myDNA for Australian patients and may be Medicare subsidised. For non-Australian patients, consult your local pathology provider for TPMT genotype testing.

## POTENTIAL DRUG INTERACTIONS

The effect of drug-drug interactions can be additive to the effect of genotype on drug metabolism. Inhibitors can decrease and inducers can increase metabolism, leading to changes in drug concentration and clinical effects.

Comments in the medications of interest and future medications sections only consider the effects of the patient's genotype, not those due to interacting drugs. For the health professional's consideration, the table below identifies which of the patient's current drugs may inhibit or induce those enzymes tested by myDNA. The extent of the inhibition or induction depends on the dose and duration of the therapy. The overall effect on metabolism by a specific enzyme may be estimated by considering both the genetic finding and the potential interacting drug.

MEDICATION	INHIBITOR – MODERATE	INHIBITOR - STRONG	INDUCER
Tobacco Smoke			CYP1A2



## FUTURE MEDICATIONS

The following tables outline personalised recommendations for future medications.

*NOTE: These tables do not account for the effect of any inhibitors or inducers. The table is not an all-inclusive list of medications but includes many commonly prescribed medications.*

### MEDICATIONS WITH MAJOR PRESCRIBING CONSIDERATIONS

MEDICATION (DRUG CATEGORY)	INTERPRETATION	RECOMMENDATION
Acenocoumarol (Anticoagulants)	<b>VKORC1 - Moderately reduced VKORC1 enzyme level</b> <b>CYP2C9 - Intermediate metaboliser:</b> Slightly reduced metabolism of acenocoumarol by CYP2C9 is predicted. Reduced amount of VKORC1 present (the enzyme inhibited by acenocoumarol). Overall increased sensitivity to acenocoumarol, an increased risk of both supratherapeutic INR and bleeding, and a lower dose requirement are predicted.	Based on the CYP2C9 and VKORC1 genotypes, DPWG <sup>1,2</sup> states that no specific action is required for dosing of acenocoumarol. Genetic variation may lead to a decrease in the required maintenance dose, however there is insufficient evidence that this causes problems when therapy is initiated as usual, i.e. with frequent INR monitoring.
Warfarin (Anticoagulants)	<b>VKORC1 - Moderately reduced VKORC1 enzyme level</b> <b>CYP2C9 - Intermediate metaboliser:</b> Slightly reduced metabolism of warfarin by CYP2C9 is predicted. Reduced amount of VKORC1 (the enzyme warfarin inhibits). The combined CYP2C9 and VKORC1 results predict increased warfarin sensitivity and increased risk of supratherapeutic INR.	CYP2C9 and VKORC1 - For patients already taking warfarin (e.g. more than 5 doses), dose adjustment is guided by INR.  For patients initiating warfarin, there are CPIC <sup>3</sup> recommendations to reach the therapeutic dose. The summary of CPIC recommendations include consideration of the use of validated published pharmacogenetic algorithms <sup>4,5</sup> available at warfarindosing.org that take into account clinical details as well as genetic findings. See CPIC guidelines for further details. If the patient identifies to be of African ancestry, CPIC provides recommendations for special dosing requirements for warfarin.
Efavirenz (Antivirals)	<b>CYP2B6 - Intermediate metaboliser:</b> Reduced metabolism of efavirenz and higher dose-adjusted trough	CPIC and DPWG <sup>6,7</sup> provide a moderate recommendation to consider initiating efavirenz with decreased dose of 400 mg/day.

**MEDICATIONS WITH MAJOR PRESCRIBING CONSIDERATIONS**

MEDICATION (DRUG CATEGORY)	INTERPRETATION	RECOMMENDATION
	<p>concentrations compared with normal metabolisers is predicted. This has been associated with an increased risk of concentration-dependent adverse effects, including CNS adverse events.</p>	<p>If therapeutic drug monitoring is available and a decreased dose of efavirenz is prescribed, consider obtaining steady-state plasma efavirenz concentrations to ensure they are in the suggested therapeutic range. The potential benefits and risks of the reduced dose and pill number should be considered.</p>
Fluvastatin (Statins)	<p><b>SLCO1B1 - Normal transporter function</b> <b>CYP2C9 - Intermediate metaboliser:</b> This SLCO1B1 genotype is associated with typical statin exposure and myopathy risk.<sup>8</sup></p> <p>This CYP2C9 genotype predicts increased fluvastatin exposure as compared with normal metabolisers, which may translate to increased myopathy risk.<sup>8</sup></p> <p>Other factors that may further increase this myopathy risk include higher doses, certain co-administered drugs, female sex, patient frailty, renal failure, hypothyroidism, advanced age, low BMI, intense physical exercise and/or African ancestry.</p>	<p>CPIC guidelines<sup>8</sup> provide a moderate recommendation to prescribe less than or equal to 40mg daily as a starting dose and adjust doses based on disease-specific guidelines. If doses &gt;40mg are required for desired efficacy, consider an alternative statin or combination therapy (i.e. fluvastatin plus non-statin guideline directed medical therapy).</p>

**MEDICATIONS WITH MINOR PRESCRIBING CONSIDERATIONS**

DRUG CATEGORY	MEDICATION	GENE(S) INVOLVED	POTENTIAL CLINICAL ISSUES	PUBLISHED GUIDELINES
ADHD - miscellaneous agents	Atomoxetine	CYP2D6	Reduced / inadequate response	CPIC <sup>9</sup> , FDA <sup>10</sup>
Angiotensin receptor blockers	Irbesartan	CYP2C9	Increased therapeutic and/or adverse effects	-
Antidepressants - other	Agomelatine	CYP1A2	Reduced / inadequate response	-
	Bupropion	CYP2B6	Altered response	-
	Mirtazapine	CYP2D6 CYP1A2	Reduced / inadequate response	-
Antidepressants - SNRIs	Duloxetine	CYP2D6 CYP1A2	Reduced / inadequate response	CPIC <sup>11</sup>
Antidepressants - SSRIs	Sertraline	CYP2B6 CYP2C19	Increased therapeutic and/or adverse effects	CPIC <sup>11</sup>
Antidiabetics	Gliclazide	CYP2C9 CYP2C19	Increased therapeutic and/or adverse effects	DPWG <sup>12</sup>
	Glimepiride	CYP2C9	Increased therapeutic and/or adverse effects	DPWG <sup>13</sup>
	Glyburide	CYP2C9	Increased therapeutic and/or adverse effects	DPWG <sup>14</sup>
Antiepileptics	Fosphenytoin	CYP2C9	Adverse effects	CPIC <sup>15</sup>
	Phenytoin	CYP2C9	Adverse effects	CPIC <sup>15</sup>

MEDICATIONS WITH <u>MINOR</u> PRESCRIBING CONSIDERATIONS				
DRUG CATEGORY	MEDICATION	GENE(S) INVOLVED	POTENTIAL CLINICAL ISSUES	PUBLISHED GUIDELINES
Antipsychotics	Clozapine	CYP2D6 CYP1A2	Reduced / inadequate response	DPWG <sup>16</sup>
	Olanzapine	CYP1A2	Reduced / inadequate response	DPWG <sup>16</sup>
Antivirals	Nevirapine	CYP2B6	Adverse effects	-
Beta blockers	Propranolol	CYP2D6 CYP1A2	Altered response	-
Haemostatic agents	Avatrombopag	CYP2C9	Altered response	FDA <sup>17</sup> , TGA <sup>18</sup>
Miscellaneous	Dronabinol	CYP2C9	Adverse effects	-
	Lesinurad	CYP2C9	Adverse effects	-
NSAIDs	Celecoxib	CYP2C9	Increased therapeutic and/or adverse effects	CPIC <sup>19</sup>
	Flurbiprofen	CYP2C9	Adverse effects	CPIC <sup>19</sup>
	Ibuprofen	CYP2C9	Adverse effects	CPIC <sup>19</sup>
	Lornoxicam	CYP2C9	Adverse effects	CPIC <sup>19</sup>
	Meloxicam	CYP2C9	Adverse effects	CPIC <sup>19</sup>
	Piroxicam	CYP2C9	Adverse effects	CPIC <sup>19</sup>
	Tenoxicam	CYP2C9	Adverse effects	CPIC <sup>19</sup>
Opioid Analgesics	Methadone	CYP2B6	Altered response	-
Proton pump inhibitors	Dexlansoprazole	CYP2C19	Reduced / inadequate response	CPIC <sup>20</sup>
	Lansoprazole	CYP2C19	Reduced / inadequate response	CPIC <sup>20</sup>
	Omeprazole	CYP2C19	Reduced / inadequate response	CPIC <sup>20</sup>
	Pantoprazole	CYP2C19	Reduced / inadequate response	CPIC <sup>20</sup>

MEDICATIONS WITH <u>USUAL</u> PRESCRIBING CONSIDERATIONS				
DRUG CATEGORY	MEDICATION	GENE(S) INVOLVED	POTENTIAL CLINICAL ISSUES	PUBLISHED GUIDELINES
ADHD - miscellaneous agents	Viloxazine	CYP2D6	No altered effect predicted by genotype	-
Angiotensin receptor blockers	Losartan	CYP2C9	No altered effect predicted by genotype	-
Antianginals	Perhexiline	CYP2D6	No altered effect predicted by genotype	-
Antiarrhythmics	Flecainide	CYP2D6	No altered effect predicted by genotype	-
	Propafenone	CYP2D6	No altered effect predicted by genotype	-
Anticholinergics (genitourinary)	Darifenacin	CYP2D6	No altered effect predicted by genotype	-
	Fesoterodine	CYP2D6	No altered effect predicted by genotype	-
	Tolterodine	CYP2D6	No altered effect predicted by genotype	-
Anticholinesterases	Donepezil	CYP2D6	No altered effect predicted by genotype	-
	Galantamine	CYP2D6	No altered effect predicted by genotype	-
Anticoagulants	Prasugrel	CYP2C19	No altered effect predicted by genotype	DPWG <sup>21</sup>
	Ticagrelor	CYP2C19	No altered effect predicted by genotype	DPWG <sup>22</sup>
Antidepressants - other	Mianserin	CYP2D6	No altered effect predicted by genotype	-

**MEDICATIONS WITH USUAL PRESCRIBING CONSIDERATIONS**

DRUG CATEGORY	MEDICATION	GENE(S) INVOLVED	POTENTIAL CLINICAL ISSUES	PUBLISHED GUIDELINES
	Hydrocodone	CYP2D6	No altered effect predicted by genotype	CPIC <sup>34</sup>
	Morphine	OPRM1	Associated with increased sensitivity to morphine	CPIC <sup>34</sup>
	Oliceridine	CYP2D6	No altered effect predicted by genotype	-
	Oxycodone	CYP2D6	No altered effect predicted by genotype	CPIC <sup>34</sup>
	Tramadol	CYP2D6	No altered effect predicted by genotype	CPIC <sup>34</sup>
Proton pump inhibitors	Esomeprazole	CYP2C19	No altered effect predicted by genotype	-
	Rabeprazole	CYP2C19	No altered effect predicted by genotype	-
Psychostimulants	Amphetamine	CYP2D6	No altered effect predicted by genotype	-
	Dextroamphetamine	CYP2D6	No altered effect predicted by genotype	-
	Lisdexamfetamine	CYP2D6	No altered effect predicted by genotype	-
Statins	Atorvastatin	SLCO1B1	No altered effect predicted by genotype	CPIC <sup>8</sup>
	Lovastatin	SLCO1B1	No altered effect predicted by genotype	CPIC <sup>8</sup>
	Pitavastatin	SLCO1B1	No altered effect predicted by genotype	CPIC <sup>8</sup>
	Pravastatin	SLCO1B1	No altered effect predicted by genotype	CPIC <sup>8</sup>
	Rosuvastatin	ABCG2 (rs2231142) SLCO1B1	No altered effect predicted by genotype	CPIC <sup>8</sup>
	Simvastatin	SLCO1B1	No altered effect predicted by genotype	CPIC <sup>8</sup>

**LEGEND:**

CPIC = Clinical Pharmacogenetics Implementation Consortium  
 DPWG = The Royal Dutch Pharmacists Association – Pharmacogenetics Working Group

TGA = Therapeutic Goods Administration (Australia)  
 FDA = Food and Drug Administration (US)

CPIC and DPWG guidelines are available on the PharmGKB website [www.pharmgkb.org/view/dosing-guidelines.do](http://www.pharmgkb.org/view/dosing-guidelines.do)



## PHARMACOGENOMIC INTERPRETATION

**EXPLANATION OF GENETIC RESULTS**

GENE	GENOTYPE	PREDICTED FUNCTION
CYP2D6	*1/*1	<p><b>CYP2D6 - Normal metaboliser</b></p> <p>Due to the presence of two copies of normal function alleles, this individual is predicted to have a normal metaboliser phenotype. For a drug extensively metabolised by CYP2D6, drug exposure and clinical effects may be expected to lie within the normal range.</p>

**EXPLANATION OF GENETIC RESULTS**

GENE	GENOTYPE	PREDICTED FUNCTION
<b>CYP2C19</b>	*1/*1	<b>CYP2C19 - Normal metaboliser</b> Due to the presence of two copies of normal function alleles, this individual is predicted to have a normal metaboliser phenotype. For a drug extensively metabolised by CYP2C19, drug exposure and clinical effects may be expected to lie within the normal range.
<b>CYP2C9</b>	*1/*2	<b>CYP2C9 - Intermediate metaboliser</b> Due to the presence of one normal function allele and one decreased function allele, this individual is predicted to have an intermediate metaboliser phenotype. For a drug extensively metabolised by CYP2C9, drug exposure and clinical effects may either be increased (for an active drug) or decreased (for a prodrug). As the decreased function allele is associated with only a small reduction in enzyme function, this variation may only be significant for certain medications, with high dosages or if drug-drug interactions occur.
<b>VKORC1</b>	AG	<b>VKORC1 - Moderately reduced VKORC1 enzyme level</b> The VKORC1 enzyme is predicted to be present in moderately reduced amounts and the response to warfarin will be enhanced. The CYP2C9 genotype should also be considered together with the VKORC1 genotype for calculating the initial warfarin dose.
<b>CYP1A2</b>	*1F/*1F	<b>CYP1A2 - Ultrarapid metaboliser (with inducer present)</b> Due to the presence of two *1F alleles, this individual is predicted to have an ultrarapid metaboliser phenotype. Enzyme activity is highest in the presence of inducers, such as tobacco smoke, regular consumption of cruciferous vegetables or chargrilled meats, and certain drugs. For a drug extensively metabolised by CYP1A2, drug exposure and clinical effects may either be reduced (for an active drug) or increased (for a prodrug).
<b>CYP3A4</b>	*1/*1	<b>CYP3A4 - Normal metaboliser</b> The *22 allele is not present and this individual is expected to have a normal metaboliser phenotype. Whilst many drugs are known to be metabolised by CYP3A4, relatively few genetic variations have been found that affect metabolism of a limited number of these drugs.
<b>CYP3A5</b>	*3/*3	<b>CYP3A5 - Poor metaboliser</b> Due to the presence of two no function alleles, this individual is predicted to have a poor metaboliser phenotype (CYP3A5 non-expressor). CYP3A5 is known to metabolise certain drugs, including tacrolimus. Note that this individual's phenotype is the most common one amongst Caucasians.
<b>SLCO1B1</b>	*1/*1	<b>SLCO1B1 - Normal transporter function</b> The decreased function *5 allele is not present and this individual is predicted to have normal function of the SLCO1B1 encoded transporter. The transporter is important for the clearance of certain drugs, including simvastatin.
<b>CYP2B6</b>	*1/*6	<b>CYP2B6 - Intermediate metaboliser</b> This individual is predicted to have an intermediate metaboliser phenotype due to the presence of one normal function allele and one decreased function allele. Due to technical difficulties in unambiguously determining this genotype, the individual's other possible genotype is *4/*9 which also predicts an intermediate metaboliser phenotype. For a drug extensively metabolised by CYP2B6, drug exposure and clinical effects may either be increased (for an active drug) or decreased (for a prodrug). This individual is at risk of experiencing adverse effects (active drug) or therapeutic failure (prodrug).