





# Practical considerations on TDM

Azucena Aldaz Pharm D, Ph D.









#### CONFLICT OF INTEREST

#### There are no conflicts of interest to declare

in collaboration with Roger Jelliffe, founder of LAPK (University of Southern California)





# **Outline**

- The Therapeutic Drug Monitoring Process
- Request form
- Laboratory measurement
- Pharmacogenetics-guided TDM
- Overview of commonly monitored drugs

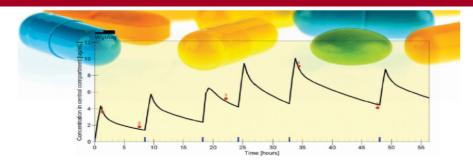




# Therapeutic drug monitoring process

#### **GOAL**

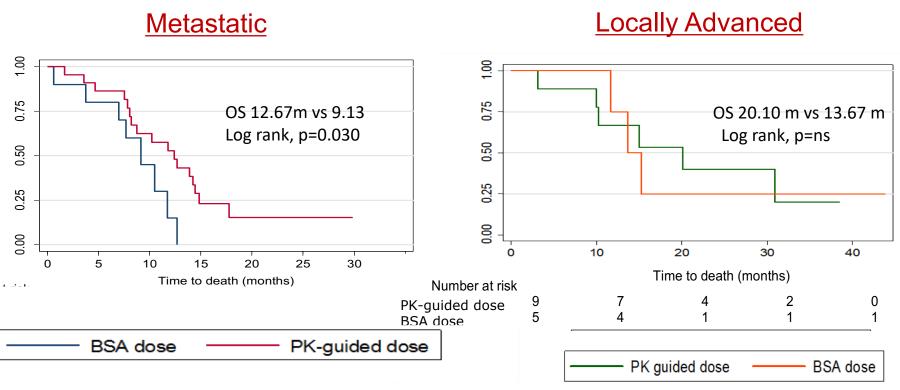
Improve health outcomes in a cost-effective manner, by optimizing individual pharmacotherapy







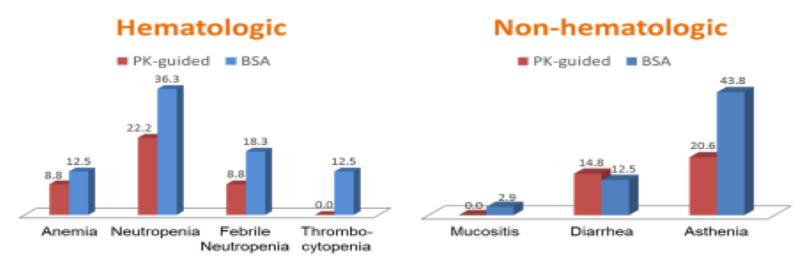
#### Clinical Outcomes in Pancreas Cancer



Authors: Azucena Aldaz et al, 2017







Neutropenia and asthenia, the most common adverse events, were significantly lower in the PK-guided group





### **HOW???**





**Tools** 

Knowledge





# Therapeutic drug monitoring process

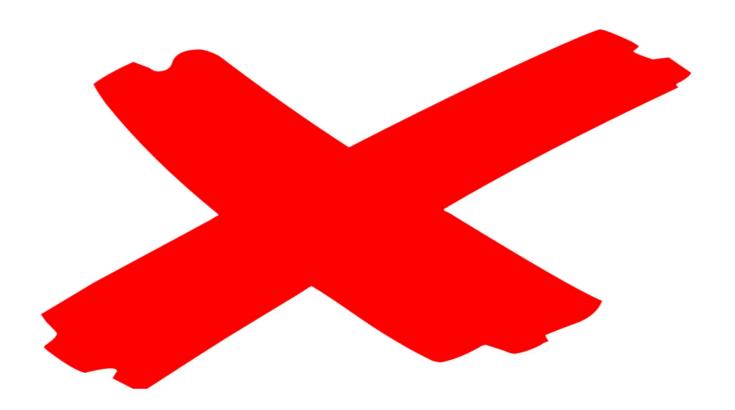
- Monitoring request
- Assessment of request suitability
- Biological samples: optimal sampling strategies
- Concentration data: Bioanalytical methods for TDM
- Estimation of individual pharmacokinetics parameters
- Analysis of clinical information
- Pharmacokinetic report



# Learning Objectives

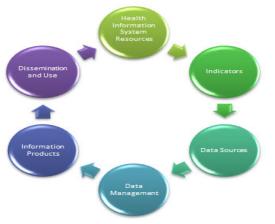
- Learn the different phases of the TDM process
- Know the advantages and disadvantages of the different analytical methods
- Understand the importance of quantifying concentrations below LLOQ
- Determine the assay error
- Assess the Bayesian adjustment methods





#### Co-medication





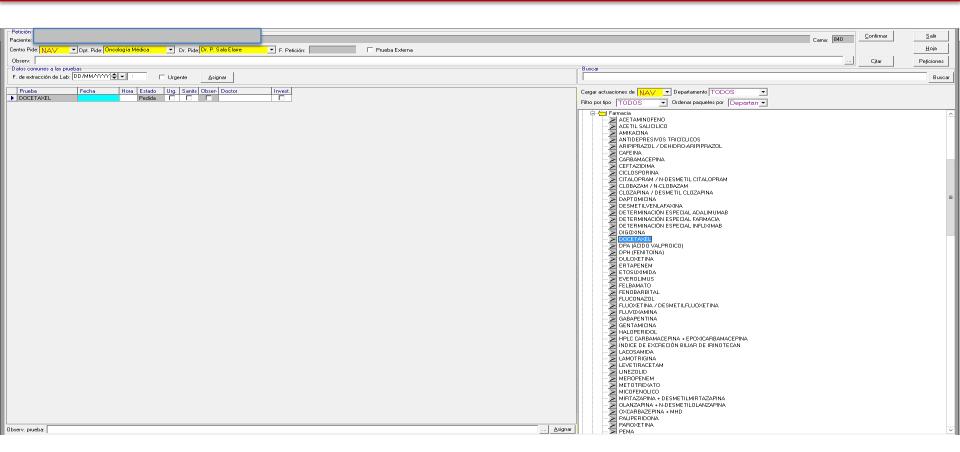
# HIS Health Information System

Figure 1 The six components of a Health Information System

Nowadays, professionals directly access in the electronic patient record to the drug monitoring application included by the Clinical Pharmacokinetic Unit in the HIS (Health Information System) of the







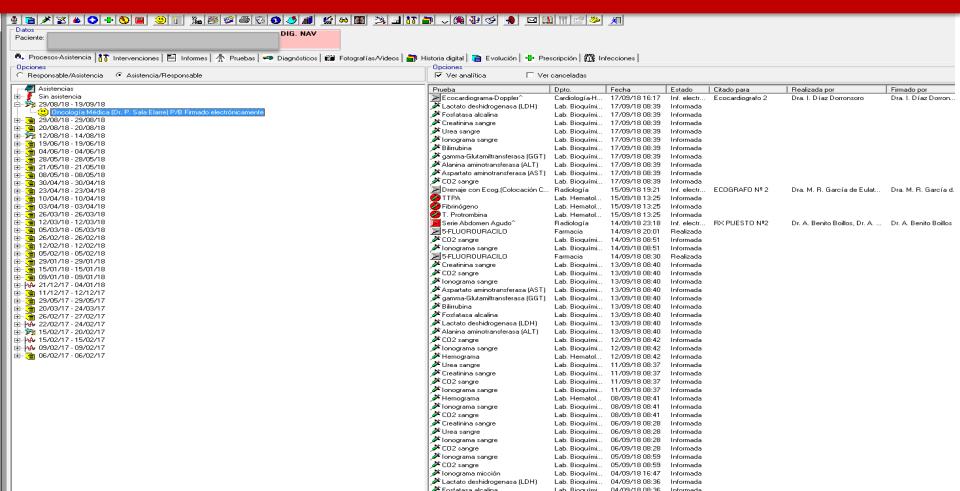




#### Clínica Universidad de Navarra - FARMACIA - Dra. A. Aldaz Pastor Archivo Puestos Carpetas Validación Consulta C. Calidad Trazabilidad Registros Documentación Informes Resultados Ventana × 63 Sistema Informático de Laboratorios - Clínica Universitaria \*\*\*\*\* PRUEBAS PENDIENTES \*\*\*\*\* Seleccion de Pruebas Pruebas demoradas Pruebas de CITOSTATICOS. Primera prueba pendiente desde 21/09/18 08:30 Carpeta ANALGESICOS Muestra (C. Barras) ANTI THE ANTIBIOTICOS Cod. Muestra Muestra Historia Prueba Extracción Dem. Lista ANTIDEPRESIVOS ▶ BGW-4046事 5-FLUOROURACILO 21/09/18 08:30 No Hepar, 5-verde FAR 213730 ANTIEPILEPTICOS ANTIPSICÓTICOS × BETALACTAMICOS Datos de la actuación BRONCODILATADORES Paciente CARDIOTONICOS 05/12/1946 Cama 801 F. Nacimiento Anular Salir CITOSTATICOS Historia INMUNOSUPRESORES. Sexo HOMBRE Paciente PRUEBAS ESPECIALES F <u>F</u>inalizar **PSICOFARMACOS** Actuación Nº Act. Plan. 36691393 Carpeta CITOSTATICOS Lista realización ☐ Urgente 5-FLUOROURACILO □ Demorada Actuación Posición Dr. solicitante Dra. A. Chopitea Ortega Dpto. solicitante Oncología Médica Recibida Estado Fec. planific. 21/09/2018 08:30:00 Seguimiento Muestras Resultados Cuestionario Incidencias Trazabilidad Reclamaciones Sequimiento Mostrar Sólo Urgencias 19/09/2018 20:04:26 Dr. X. Abasolo Tamayo Solicitud Imprimir Etiquetas 21/09/2018 08:30:00 | Huarte Huarte, Isabel Extracción Recepción 21/09/2018 08:45:00 Huarte Huarte, Isabel









504258/2018215270

CONTROL

26/09/2018

F. Nac.: 23/07/1954

Paciente:

Lista realización: 4754653

Dpto. solicitante: MAD - Digestivo

Doctor solicitante

Determinación especial Farmacología

Concentración. Informe

MUESTRAS

BGW-8746

OBSERVACIONES

\* PETICIÓN: 13684211

Motivo pet. prueba: niveles de depakine

\* CUESTIONARIO:

Nombre de la prueba a realizar, niveles de depakine

Especímen, cantidad y condiciones de conservación: 300 ma 1-1-1

25/09/2018 8:09:00

Persona de contacto del departamento solicitante:



# Prediction of Future Serum Concentrations with Bayesian Fitted Pharmacokinetic Models: Results with Data Collected by Nurses Versus Trained Pharmacy Residents

Therapeutic Drug Monitoring 16:166-173 © 1994 Raven Press, Ltd., New York

B. Charpiat, \*V. Breant, †C. Pivot-Dumarest, ‡P. Maire, and §R. Jelliffe

**TABLE 1.** Patient demographic data for the RCD and NCD groups<sup>a</sup>

	RCD group	NCD group
Male	3 (30%)	7 (30.4%)
Female	7 (70%)	16 (69,6%)
Age (yr)	$79.1 \pm 7.9$	$81.3 \pm 7$
Height (m)	$1.6 \pm 0.15$	$1.6 \pm 0.08$
Weight (kg)	$58.8 \pm 14.4$	$60.5 \pm 12.6$
Initial estimated creatinine clearance		
(ml/mm/1.73 m <sup>2</sup> )	$41.9 \pm 15.6$	$42.5 \pm 15$
No. of patients with four		
levels in first cluster	10 <sup>b</sup>	$6^{b}$
No. patients with ≤three	_	
levels in first cluster	$\mathbf{o}_{\mathbf{p}}$	$17^{b}$
All doses given i.m.	3	4
All doses given i.v.	6	13

<sup>&</sup>lt;sup>a</sup> All ± data are listed as mean values ± SD.

TABLE 4. Percent of serum levels accurately predicted (within ±20%) in RCD and NCD groups using the MAP Bayesian one-compartment fitted model

Bi	RCD	NCD	p-value <sup>a</sup>	
PSLAP	80%	22%	<0.01	
TSLAP	30%	13%	NS <sup>b</sup>	

PSLAP = peak levels accurately predicted and TSLAP = trough levels accurately predicted.

RCD: resident- collected data NCD: nurse-collected data

 $<sup>^{</sup>b}\chi^{2} = 12.7, p < 0.001.$ 

a Comparing groups.

<sup>&</sup>lt;sup>b</sup> NS = not significant.



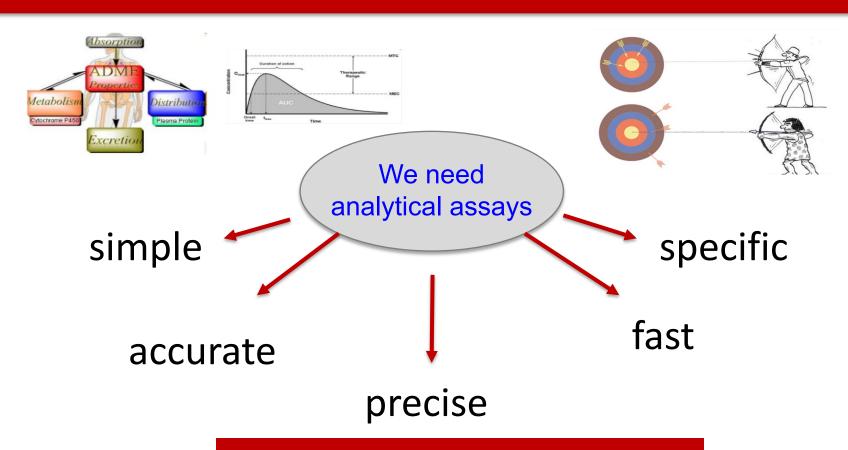


# Bioanalytical methods for TDM

The most precise methods for dosage adjustment use drug concentrations measured in biological samples, drawn at determined times







We need to control the results







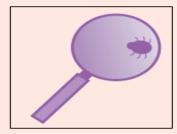
Hits the right target

#### **Precise**



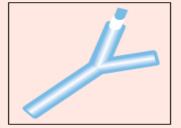
Hits the same target

#### Sensitive



Ability to exclude false *negatives* 

#### **Specific**



Ability to exclude false positives



















#### **IMMUNOASSAYS**



Are bioanalytical methods in which the quantitation of the analyte depends on the reaction of an antigen and an antibody (immune complex)

Homogeneous and Heterogeneous

Competitive and Non-competitive





#### **IMMUNOASSAYS**

# Reagents

**Antibodies** 

Monoclonal Polyclonal

Signal-generating labels

- Radioactive atoms
- Enzimes
- Fluorescens probes
- Chemiluminiscens
- Metal
- Liposomes

Separation matrices

Charcoal
Polyethilene glicol
Second antibody
Microbeads
Microwell plate





#### **Chemiluminescent Magnetic Immunoassay—CMIA**

**Electrochemiluminescence Immunoassay- ECLIA** 

Fluorescence Polarization Immunoassay - FPIA

Microparticle Enzyme Immunoassay - MEIA

**Antibody-conjugated magnetic immunoassay- ACMIA** 

**Quantitative Microparticle System- QMS** 

Cloned enzyme doner immunoassay- CEDIA





# Characterization of cross reactivity by carbamazepine 10,11-epoxide with carbamazepine assays

Sijiu Shena, Ronald J. Elinb, Steven J. Soldinc,d,\*

ization immunoassay. The cross-reactivity of the enoxide

Measurement of carbamazepine and carbamazepine 10, 11-epoxide by different analytical systems

Systems	Z01*			Z02*		Differ	T value	P value	
	N	Mean	STD	N	Mean	STD	(Z02-Z01)		
Abbott axsym	1512	7.24	0.26	1514	8.34	0.30	1.10	107.77	< 0.001
Abbott TDX/TDXFLX	230	7.28	0.36	233	8.32	0.39	1.04	29.81	< 0.001
BDI opus/plus/magnum	22	10.69	0.57	23	11.55	0.62	0.86	4.84	< 0.001
Beckman synchron RGT	324	6.31	0.47	325	6.69	0.48	0.38	10.19	< 0.001
Cedia	136	8.41	0.41	137	8.74	0.39	0.33	6.81	< 0.001
Chiron ACS:180	42	6.51	0.52	41	6.70	0.52	0.19	1.66	0.100
Dade ACA	299	7.75	0.29	299	9.96	0.44	2.21	72.52	< 0.001
Dade dimension	356	7.03	0.33	359	11.71	0.63	4.68	124.27	< 0.001
Roche cobas integra	116	7.89	0.34	117	8.41	0.36	0.52	11.33	< 0.001
Syva emit	11	7.83	0.73	11	7.85	0.81	0.02	0.06	0.952
Syva emit 2000	59	7.44	0.50	59	7.37	0.45	-0.07	0.80	0.426
Technicon immuno-1	68	7.32	0.80	68	7.4	0.76	0.08	0.60	0.551
Vitros	110	5.14	0.41	109	5.14	0.42	0.00	0.00	1.000
All methods	3339	7.21	0.74	3349	8.56	1.53	1.35	45.91	< 0.001

<sup>\*</sup> The target value for Z01 is 7.5 mg/L carbamazepine, for Z02 is 7.5 mg/L carbamazepine plus 5.0 mg/L carbamazepine 10, 11-epoxide. The mean concentrations are given in mg/L. Student's t test is used for statistical analysis.



#### Estimation of Carbamazepine and Carbamazepine-10,11-Epoxide Concentrations in Plasma Using Mathematical Equations Generated With Two Carbamazepine Immunoassays

Gwendolyn A. McMillin, PhD, 1,2 JoEtta M. Juenke, MT(ASCP), Gertie Tso, MT(ASCP), and Amitava Dasgupta, PhD<sup>4</sup>

Cross-Reactivity of Carbamazepine-10,11-Epoxide With the PETINIA, ADVIA Centaur, and CEDIA\*

Epoxide (µg/mL)	PETINIA	ADVIA Centaur	CEDIA
0.0	ND (—)	ND (—)	ND (—)
1.0	0.96 (96.0)	ND (—)	ND (—)
1.5	1.3 (86.6)	ND (—)	ND (—)
2.5	2.3 (92.0)	ND (—)	ND (—)
3.5	3.7 (105.7)	ND (—)	ND (—)
5.5	5.4 (98.2)	0.37 (6.8)	0.52 (9.4)
7.5	7.3 (97.3)	0.47 (6.3)	0.62 (8.2)
10.0	10.3 (103.0)	0.64 (6.4)	1.27 (12.7)
18.0	16.8 (93.3)	0.97 (5.3)	2.09 (11.6)
25.0	23.4 (93.6)	1.92 (7.7)	2.65 (10.6)





# **Immunoassays**





Automated

Standarized methods

Low sample volume

No sample preparation (or minimum)

Less time of analysis

Fairly low cost (instruments, reagents)

Easy to use. Personal minimally qualified

Less sensitivity (heterogeneous > homogeneous)

Less specificity

Total dependence on provider

Limited supply of assays

No availability for new drugs



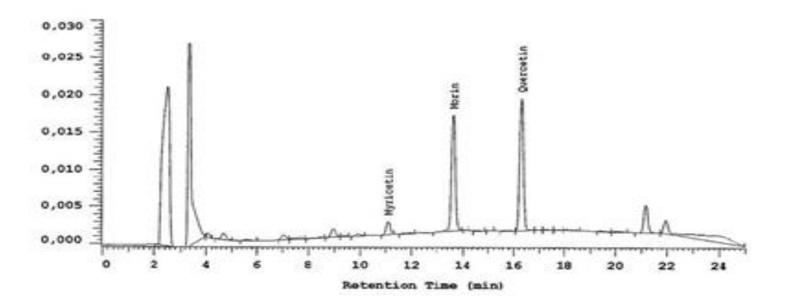


# Chromatography













#### Chromatographic methods





Standarized and in-house methods Good or high sensitivity (LC-MS/MS) Good or high specificity (LC-MS/MS)

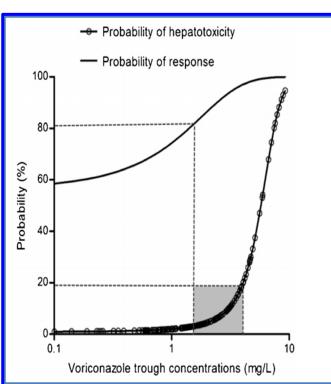
HPLC high sample volume required
Sample preparation required
Lengthy analysis (HPLC)
Moderate analysis length (LC-MS/MS)
Expensive instrumentation
Qualified and experienced staff





Which drugs may require therapeutic drug monitoring?

- Narrow and established target concentration range
- Significant interindividual pharmacokinetic variability
- No high intraindividual pharmacokinetic variability
- A better relationship between concentration and response that between dose and response
- Therapeutic benefit demonstrate by TDM
- Possibility of having analytical methods to quantify the drug







# Methods of dosage optimization

#### Methods "a priori"

Based on population pharmacokinetics parameters

#### Methods "a posteriori"

Regression methods

Use data of TDM concentrations

#### Bayesian methods

Use population pharmacokinetics parameters + data of TDM concentrations



# Methods "a priori"



All V Q

All - A B C D E F G H I J K L M N O P Q R S T U V W X Y Z

HOME ABBREV CALCULATORS MED TERMINOLOGY DILUTIONS DRUGS I.DX. LABS MEDICAL FINANCIAL

ONCOLOGY RENAL RX LIST





	PHARMACOKINETIC DOSING  AMINOGLYCOSIDE-VANCOMYCIN DOSING			
	< Link to other Dosing Calculators / Internal Medicine >> 地面			
Patient Name	e: Location: Conventional dosing \$			
Select drug: Tol	ntamicin bramycin nikacin ncomycin			
	Program Hints -			
	Need dosing information for once daily dosing?			
	Gent-tobra: Mild (4 mg/kg)/ Amikacin (10 mg/kg)			
	Age: Weight: Kg \$ Gender: Male \$			
	SCR: Height: Centimeters ♦			
Desir	ed peak: Desired trough: Infusion time: 0.5 \$\diangle\$ hours			
	Volume of distribution: Please Select Value ♦ L/kgo			
	Usual range: aminoglycosides: 0.25 - 0.35 Vancomycin: 0.65 - 0.9			



# Algorithms

(example Lithium)

Dose = 
$$6.21 \, \text{C}_{\text{goal}}^{\text{ss}} - 1.93_{\text{form}} - 2.84 \, \text{ADT-} \ 0.07 \, \text{age}$$
  
+  $1.88 \, \text{sex+} \ 0.08 \, \text{weight} + 5.14$ 

*N* = 100 patients with bipolar disorder





These methods have limited utility and should only be used for initial dosing, provided that the population information on which they are based is well known and validated in routine clinical practice.



## Methods "a posteriori"

Linear regression

$$y = a.x + b \Rightarrow ln Cp = ln(D/Vd) - Ke.t$$

$$a = Ke = \frac{\sum (t) \cdot \sum (\ln Cp) - n \sum (t \cdot \ln Cp)}{\left(\sum (t)\right)^{2} - n \sum (t^{2})}$$

$$b = \ln \frac{D}{Vd} = \frac{\sum(t).\sum(t.\ln Cp) - n\sum(t^{2})\sum(\ln Cp)}{(\sum(t))^{2} - n\sum(t^{2})}$$





## Advantages of linear regression

- Simplicity
- Speed
- No computer is needed

## Limitations

- Implicit errors in linearization
- Does not allow the use of all information



## Methods "a posteriori"

Non- Linear regression

$$SS = \sum_{i=1}^{n} Wi.(Ci, t - f(Pm, t))^{2}$$



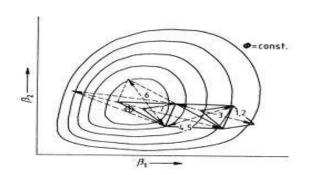


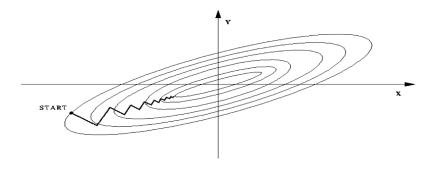
## Direct search algorithms

Simplex Nelder-Mead

## Gradient algorithms

Steepest Descent Marquart-Levenberg Gauss-Newton







## Advantages of Non-linear regression

- Data weighting
- Use of complex equations, including kinetic and clinical parameters
- Does not require additional population information

## Limitations

- Need a greater number of concentrations (≥3)
- Need a computer
- Requires experience to detect false solutions



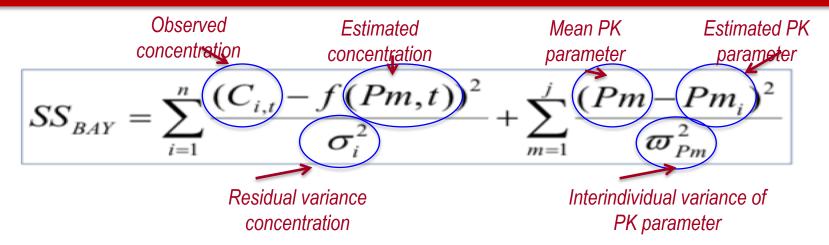
## Bayesian methods

Bayes' theorem, is used in many different ways.

In TDM, it provides a way to revise existing predictions or probabilities given new or additional evidence, using either parametric or (better) nonparametric approaches

$$P(A|B) = \frac{P(B|A) P(A)}{P(B)}$$
Thomas Bayes
1702-1761





#### Information "a priori":

PK parameters
Interindividual variance of each PK parameter
Residual variance of concentrations

#### Current information:

Observed concentrations

#### Information "a posteriori":

Estimated PK bayesian parameters
Estimated concentrations using PK bayesian parameters





## Advantages of Bayesian methods

- Minimum experimental information
- Flexibility of sampling time
- Consistent results
- Application to different PK models

## Limitations

- Clinical training
- Validate the PK model in the own population







## All estimation methods require a total control of :

- time of administration
- time of the previous doses or at least of the usual schedule in the case of ambulatory patients





## How to do Optimal TDM and Individualize Drug Therapy Optimally

We need good data, <u>accurately recorded</u>, precisely measured, with a good quantitative *index* of its <u>credibility</u>.

1 - WHEN was it drawn? Look at the clock on the wall, or your watch. Record the time as military time, to the minute, from 0000 to 2359 hrs. This is MOST IMPORTANT!

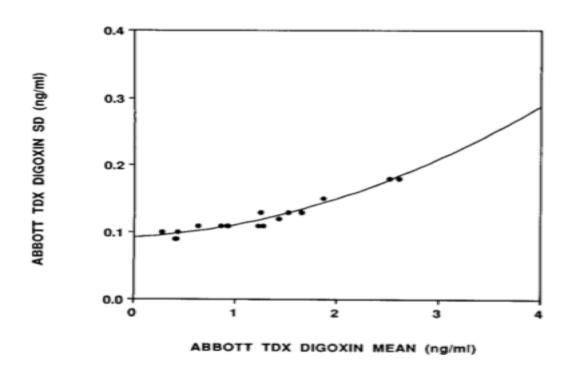
Same for the doses – even more important.

2 – How CREDIBLE is the assay?





We need to know the assay error over the entire range of the assay, down to and including the zero blank







## **Assay Error Pattern**

## Not all data have equal credibility.

There have been various empirical weighting schemes such as:

- unity weighting
- weighting by the reciprocal of the measured concentration (or of its squared value)
- the use of a constant coefficient of variation
- and others.

We need to know the relative amount of information contained in a data point, to give more quantitative importance to one that is known with good precision and less to one known with less precision (greater measurement error).

Use 1/variance (Fisher information) as the correct error description





# Fisher Information quantifies the credibility of a lab measurement

- Fisher Information = 1/Variance
- So need to know, or have a good estimate, of the SD of every serum level.
- Labs always get SD anyway, to get the CV%.
- Then, Variance = SD<sup>2</sup>
- And Credibility = Fisher info = 1/Var.

## Assay CV% versus correct weight, Fisher Info

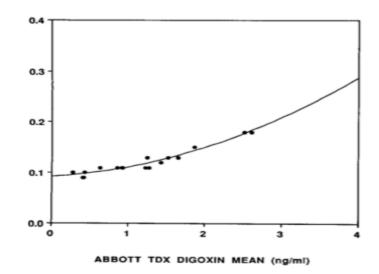
- Assume, for example, 10% assay CV
- If conc = 10, SD = 1, var = 1, weight = 1
- If conc = 20, SD = 2, var = 4, weight = ½ Aha!
- So a constant linear % error (the assay CV%) is NOT the correct measure of credibility!
- As conc approaches zero, CV% approaches infinity.
- But assay SD, var, weight are always finite. Fisher info is the correct measure of assay precision.
- Also, no need for LLOQ any more!



## Determining the Assay SD polynomial

- Measure blank, low, medium, high, and very high samples in at least quintuplicate.
- Get mean conc + SD for each sample.
- Fit a polynomial to the mean and SD data.
- SD =  $A_0C^0 + A_1C^1 + A_2C^2 + A_3C^3$
- Then can weight each single measurement by the reciprocal of its variance (Fisher Info)
- No lower detectable limit (LOD) or limit of quantification (LOQ)!
- No need to censor (withhold) any data.









# Different weighting schemes lead to different parameter values being found in fitted pharmacokinetic models

This is well known, and has been explicitly shown in pharmacokinetic studies



Therapeutic Drug Monitoring 16:552-559 © 1994 Raven Press, Ltd., New York

## Population Pharmacokinetic Models: Effect of Explicit Versus Assumed Constant Serum Concentration Assay Error Patterns upon Parameter Values of Gentamicin in Infants on and off Extracorporeal Membrane Oxygenation

\*†Warren F. Dodge, ¶Roger W. Jelliffe, ‡Joseph B. Zwischenberger, Renee A. Bellanger, §James A. Hokanson, and \*†Wayne R. Snodgrass



SD (
$$\mu$$
g/ml) = 0.170455 - (0.043038  $C$ )  
+ (0.017003  $C^2$ )

Polynomial equation

SD (
$$\mu$$
g/ml) =  $A + (0.00 C) + (0.00 C^2)$ 

A = Constant SD of 0.5 ug/ml

The two population models differ only in the laboratory assay SD pattern used.

IQR (interquartile range) = 75th percentile - 25th percentile = central 50th percentile of values. DF50 = the dispersion factor covering the central 50% of the distributions found = (75th percentile - 25th percentile) divided by 1.32.

<sup>a</sup> An assay SD error pattern that assumes a constant SD of 0.5 μg/ml at all serum gentamicin concentrations.

b The explicitly determined laboratory assay SD error pattern.



**TABLE 2.** Gentamicin pharmacokinetic parameter values ( $V_d$  and  $K_{el}$ ) for 11 neonates undergoing extracorporeal membrane oxygenation

	Constant assay SD error pattern <sup>a</sup>	Explicit assay SI error pattern <sup>b</sup>
V <sub>d</sub> (L/kg)		
Mean	0.602	0.813
SD	0.053	0.253
25th	0.551	0.609
50th	0.602	0.696
75th	0.602	0.875
IQR	0.051	0.266
DF50	0.039	0.202
$K_{ei} (h^{-1})$		
Mean	0.105	0.084
SD	0.026	0.012
25th	0.086	0.069
50th	0.096	0.075
75th	0.119	0.093
IQR	0.033	0.024
DF50	0.025	0.018



TABLE 4. Performance evaluation of two gentamicin population models for neonates undergoing extracorporeal membrane oxygenation

	Predictor <sup>a</sup>	
	Explicit assay SD pattern and median pharmacokientic parameter values	Constant assay SD error pattern and median pharmacokinetic parameter values
Precision		
Mean squared error (MSE)	1.296 (0.786, 1.806)	1.404 (0.855, 1.951)
Root mean squared error (RMSE)	1.138 (0.886, 1.343)	1.185 (0.925, 1.397)
Relative to each other (ΔMSE for predictor explicit weighting minus predictor general weighting)		-0.249, 0.033)
Bias		
Mean error (ME)	-0.369(-0.583, -0.607)	-0.645(-0.843, -0.447)
Relative to each other (ΔME for predictor explicit weighting minus	,	
predictor general weighting)	0.276 (0.124, 0.338)	

The two models differ only in the laboratory assay SD error pattern used (explicit assay SD error pattern and assumed constant assay SD error pattern) with median pharmacokinetic parameter values.

<sup>a</sup> Values given are point estimates (95th confidence interval).





## Report

The pharmacokinetic report should be written in simple, clear language, avoiding the use of complicated pharmacokinetic terms for clinicians

### It should include:

- Patient data
- Drug data
- Concentration data
- Interpretation and comments



#### DIGOXINA

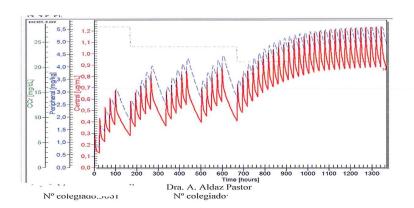
~	-	4	618368
DIGOXINA			
Solicita:	Cirugía Cardiaca	Dr. G. Rábago Juan Aracil	11/09/18
Realiza: Referencia:	Farmacia		18/09/18

La concentración sérica de digoxina en la muestra obtenida a las 08:36 h del 18/09/18 ha sido de 0.86 ng/mL.

En el gráfico adjunto se muestra el óptimo ajuste de la concentración sérica medida con la simulación realizada en función de la pauta seguida y la evolución renal y ponderal de la paciente. El punto rojo representa la concentración medida.

Se han representado 4 semanas de la antigua pauta de 0,125 mg diario excepto dos días semanales y la actual pauta diaria (0,125 mg diario de Digoxina® comprimidos).

La disfunción renal reduce significativamente el volumen de distribución aparente de digoxina reduciendo su unión a tejido y en la paciente la concentración máxima en miocardio que es la asociada con la respuesta es próxima a 5,5 mcg/kg tejido, que es el mínimo terapéutico.





#### SERTRALINA

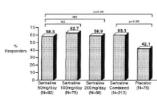
Sertralina						
Solicita:	Psiquiatría Médica	у	Psicología	Dr. P. Molero Santos	486	26/02/18
Realiza:	Farmacia			Dra. A. Aldaz Pastor		27/02/18
Referencia:	AAP/aap					28/02/2018

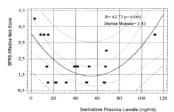
La concentración sérica de sertralina (SER) y su metabolito activo N-sertralina (N-SER) en la muestra obtenida a las 8 h del 27/2/18 han sido, respectivamente, de 158,6 ng/mL y 157,4 ng/mL, siendo la relación N-SER/SER de 0,99.

La pauta que recibe la paciente, desde al menos su ingreso el día 16/2/18, es de 200 mg en el desayuno (Aremis®).

Ya se ha comentado en los informes de otros pacientes los trabajos de Mauri et al sobre la naturaleza curvilínea de la relación entre la concentración de sertralina y la respuesta clínica. En sus trabajos este grupo muestra que los mejores resultados se observan entre 50 y 70 ng/mL. Además, ya en 1995 Preskorn, de cuyo trabajo se muestra una imagen en este informe) advertía sobre el mejor balance beneficio/riesgo de dosis bajas de sertralina sobre dosis más elevadas.

Un aspecto a destacar en los valores medidos es la relación N-SER/SER cuya mediana habitualmente es de 2 y el rango de valores es de 1,7 a 3,4. Sin embargo, en esta paciente es de 0,99. Esto puede reflejar alguna interacción, porque teóricamente han transcurrido ya 10 días desde que se incrementó a la pauta actual y por tanto ya debería haberse alcanzado el estado de equilibrio estacionario en ambas moléculas, aunque no se puede asegurar dada la elevada variabilidad interindividual. Se recomienda valorar la reducción de la dosis actual, al menos a 150 mg en un primer estadio.





Dra. A. Aldaz Pastor Nº colegiado:929





# Can pharmacogenomics be used for posology adjustments?

- Avoid adverse drug reactions (5-fluorouracil, HDMTX)
- Predicts drug efficacy (antineoplastics (RAS, EGFR mutations etc....)
- Predicts drug serum concentrations (Population PK parameters, NEVER individual predictions)
- Can be used to adjust initial posology (Need more tan one variable)



#### Population Pharmacokinetics and Pharmacogenetics of Imatinib in Children and Adults

Aurélie Petain, Darouna Kattygnarath, Julie Azard, et al.

Clin Cancer Res 2008;14:7102-7109. Published online November 3, 2008.

Purpose: The aim of this study was to explore the effect of several demographic, biological, and pharmacogenetic covariates on the disposition of imatinib and its main metabolite (CGP74588) in both adults and children.

Experimental Design: Thirty-three children with solid malignancies included in a phase II exploratory study and 34 adults with gastrointestinal stromal tumors received 340 mg/m² and 400 mg imatinib, respectively. Plasma imatinib and CGP74588 concentrations observed on day 1 and at steady-state were analyzed by a population pharmacokinetic method (NONMEM) to evaluate the effect of age, body weight, age, sex, albuminemia, plasma α1-acid glycoprotein (AGP), and eight polymorphisms corresponding to ABCB1, ABCG2, CYP3A4, CYP3A5, and AGP (pharmacogenetic data available for 46 of 67 patients).

Results: Analysis of the whole data set in 67 patients showed that apparent clearance (CL/F) of imatinib was positively correlated with body weight and albuminemia and negatively with AGP. By considering these three covariates, the interindividual variability on CL/F decreased from 47% to 19%. The apparent clearance of CGP74588 was similarly dependent on both body weight and AGP and significantly lower (30% reduction) at steady-state. By adding genotype status to the final covariate imatinib model, a 22% reduction in CL/F was observed in heterozygous compared with wild-type patients corresponding to ABCG2 c.421C>A (P < 0.05).

Conclusions: By considering morphologic and biological covariates, a unique covariate model could be used to accurately describe imatinib pharmacokinetics in patients ages 2 to 84 years. Morphologic and biological characteristics have a stronger influence than pharmacogenetics on imatinib pharmacokinetics.





## CPIC® Guideline for Fluoropyrimidines and DPYD

#### Most recent guideline publication:

Clinical Pharmacogenetics Implementation Consortium (CPIC) Guideline for Dihydropyrimidine Dehydrogenase Genotype and Fluoropyrimidine Dosing: 2017 Update (October 2017)

#### Updates since publication:

No updates on dosing recommendations since 2017 publication.

#### Tables provided in the main manuscript of the guideline:

Table 1. Assignment of likely DPD phenotype based on DPYD genotype

Table 2. Recommended dosing of fluoropyrimidines by DPD phenotype

Supplement to: <u>Clinical Pharmacogenetics Implementation Consortium (CPIC) Guideline for</u>
<u>Dihydropyrimidine Dehydrogenase Genotype and Fluoropyrimidine Dosing: 2017 Update (October 2017)</u>



Table 2 Recommended dosing of fluoropyrimidines<sup>a</sup> by DPD phenotype

Phenotype	Implications for phenotypic measures	Dosing recommendations	Classification of recommendations <sup>b</sup>
DPYD normal metabolizer	Normal DPD activity and "normal" risk for fluoropyrimidine toxicity.	Based on genotype, there is no indication to change dose or therapy. Use label- recommended dosage and administration.	Strong
DPYD intermediate metabolizer	Decreased DPD activity (leukocyte DPD activity at 30% to 70% that of the normal population) and increased risk for severe or even fatal drug tox- icity when treated with fluoropyrimi- dine drugs.	Reduce starting dose based on activity score followed by titration of dose based on toxicity or therapeutic drug monitoring (if available).  Activity score 1: Reduce dose by 50%  Activity score 1.5: Reduce dose by 25% to 50%	Activity score 1: Strong Activity score 1.5: Moderate
DPYD poor metabolizer	Complete DPD deficiency and increased risk for severe or even fatal drug toxicity when treated with fluoropyrimidine drugs.	Activity score 0.5: Avoid use of 5-fluorouracil or 5-fluorouracil prodrug-based regimens. In the event, based on clinical advice, alternative agents are not considered a suitable therapeutic option, 5-fluorouracil should be administered at a strongly reduced dose <sup>d</sup> with early therapeutic drug monitoring. <sup>c</sup> Activity score 0: Avoid use of 5-fluorouracil or 5-fluorouracil prodrug-based regimens.	Strong

<sup>\*5-</sup>fluorouracil or capecitabine. Bating scheme described in Supplement. Increase the dose in patients experiencing no or clinically tolerable toxicity in the first two cycles to maintain efficacy; decrease the dose in patients who do not tolerate the starting dose to minimize toxicities. If available, a phenotyping test (see main text for further details) should be considered to estimate the starting dose. In the absence of phenotyping data, a dose of <25% of the normal starting dose is estimated assuming additive effects of alleles on 5-FU clearance. Therapeutic drug monitoring should be done at the earliest timepoint possible (e.g., minimum timepoint in steady state) in order to immediately discontinue therapy if the drug level is too high.</p>





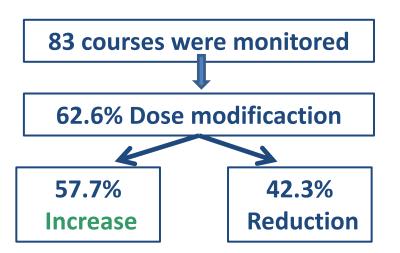
>35

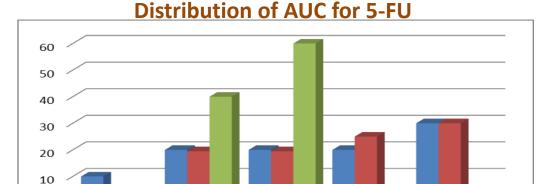
## OUR DATA ABOUT PHARMACOKINETICALLY GUIDE 5-FLUOURACIL DOSE-ADJUSTMENT 2016-2017

<20

20-25

1st course





25-30

2nd course

Dose increase (percentage) =  $13.75 \pm 9.75$ 

Dose reduction (percentage) =  $11.71 \pm 8.06$ 

BSA dosing: 30% null-low efficacy

30-35

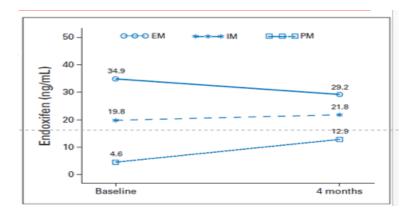
3rd course

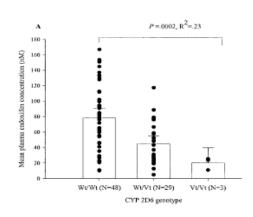


Hypothesis: The CYP2D6 polymorphic state slow metabolizer MAY AFFECT endoxifen plasma concentrations as well as disease-free survival in postmenopausal women ER + under adjuvant treatment with tamoxifen 20mg / day for 5 years

B)

A)







## CPIC® Guideline for Tamoxifen based on CYP2D6 genotype

Most recent guideline publication:

Clinical Pharmacogenetics Implementation Consortium (CPIC) Guideline for CYP2D6 and Tamoxifen Therapy (January 2018)

Updates since publication:

No updates on dosing recommendations since publication.

Tables provided in the main manuscript of the guideline:

Table 1. Assignment of likely CYP2D6 phenotypes based on diplotypes

Table 2. Dosing Recommendations for tamoxifen based on CYP2D6 phenotype

Supplement to: Clinical Pharmacogenetics Implementation Consortium (CPIC) Guideline for CYP2D6 and Tamoxifen Therapy (January 2018) [△]

#### **CPIC GUIDELINES**

Table 1 Assignment of likely CYP2D6 phenotypes based on genotypes

Phenotype <sup>a</sup>		Genotype	Examples of CYP2D6 diplotypes <sup>b</sup>	
Metabolizer	Activity score			
CYP2D6 ultrarapid metabolizer	> 2.0	An individual carrying duplications of functional alleles	*1/*1xN, *1/*2xN, *2/*2xN <sup>c</sup>	
CYP2D6 normal metabolizer	1.5 and 2.0	An individual carrying two normal function alleles or one normal func- tion and one decreased function allele	*1/*1, *1/*2, *1/*9, *1/*41, *2/*2,	
CYP2D6 normal metabolizer or inter- mediate metabolizer (controversy remains) <sup>2</sup>	1.0	An individual carrying two decreased function alleles or one normal function and one no function allele. An activity score (AS) of 1.0 is associated with decreased tamoxifen metabolism to endoxifen compared to those with an AS of 1.5 or 2.	*1/*4, *1/*5, *41/*41	
CYP2D6 intermediate metabolizer	0.5	An individual carrying one decreased function and one no function allele	*4/*10,*4/*41, *5/*9	
CYP2D6 poor metabolizer	0	An individual carrying only no func- tional alleles	*3/*4,*4/*4, *5/*5, *5/*6	

"See the CYP2D6 frequency table" for race-specific allele and phenotype frequencies. "For a complete list of CYP2D6 diplotypes and resulting phenotypes, see the CYP2D6 genotype to phenotype table." Note that genotypes with an activity score of 1 are classified as Nils in the online CYP2D6 genotype to phenotype table. "Where xit Prevales from the number of CYP2D6 genotype is prevaled to the number of CYP2D6 genotype is prevaled to the number of CYP2D6 genotype is prevaled to the number of CYP2D6 genotypes. For individuals with CYP2D6 duplications or multiplications, see supplemental data for additional information on how to translate diplotypes into phenotypes. See placetism with an activity score of 1.0 may be classified as intermediate metabolizers, some reference laboratories. A group of CYP2D6 genotype to phenotype table "3".

Therapeutic recommendation <sup>b</sup>	Classification of recommendation <sup>a</sup>
id moderate and strong CYP2D6 inhibitors. Initiate therapy with rec- mended standard of care dosing (tamoxifen 20 mg/day).	Strong
id moderate and strong CYP2D6 inhibitors. Initiate therapy with rec- mended standard of care dosing (tamoxifen 20 mg/day).	Strong
sider hormonal therapy such as an aromatase inhibitor for post- hopeusal women or aromatase inhibitor along with ovarian function pression in premenopausal women, given that these approaches superior to tamoxifen regardless of CYP2DD genotype. <sup>43</sup> If aroma- inhibitor use is contraindicated, consideration should be given to a higher but FDA approved tamoxifen dose (40 mg/day). <sup>45</sup> Avoid 2D6 strong to weak inhibitors.	Optional <sup>b</sup>
sider hormonal therapy such as an aromatase inhibitor for post- iopanual women or aromatase inhibitor along with ovarian function pression in premenopausal women, given that these approaches superior to tamoxifen regardless of CYP2/D6 genotype. <sup>43</sup> If aroma- is inhibitor use is contraindicated, consideration should be given to a higher but FDA approved tamoxifen dose (40 mg/day). <sup>45</sup> Avoid 2D6 strong to weak inhibitors.	Moderate <sup>b</sup>
sider hormonal therapy such as an aromatase inhibitor for post- opausal women or aromatase inhibitor along with ovarian function pression in premenopausal women, given that these approaches superior to tamoxifen regardless of CYP2D6 genotype. <sup>43</sup> If aroma- inhibitor use is contraindicated, consideration should be given to a higher but FDA approved tamoxifen dose (40 mg/day). <sup>45</sup> Avoid 2D6 strong to weak inhibitors.	Moderate
ommend alternative hormonal therapy such as an aromatase inhibi- ior postmenopausal women or aromatase inhibitor along with ovar- function suppression in premenopausal women given that these roaches are superior to tamoxifen regardless of CYP2D6 geno- f <sup>13</sup> and based on knowledge that CYP2D6 poor metabolizers ched from tamoxifen to anastrozole do not have an increased risk sourrence. <sup>38</sup> Note, higher dose tamoxifen (40 mg/day) increases does not nomalize endoxifen concentrations and can be consid- tif there are contraindications to aromatase inhibitor therapy. <sup>45,56</sup>	Strong

normal metabolizer." However, in the case of tamoxifen, prescribing recommendations for those with an s a \*1.0 allele are provided a "moderate" recommendation. In contrast, prescribing recommendations cause the recommendation and primarily extrapolated from evidence generated from \*1.0 individuals

The remit of this review was narrow and specifically examined the role of CYP2D6. Recent data suggest that the metabolism of TAM is complex and may be related to the effects of more than one genotype. It may be necessary, therefore, for future research to examine other metabolic pathways. In the meantime, further examination of the link between endoxifen levels and clinical outcomes could be of value and could be a mechanism that is easily integrated into existing

Health Technol Assess. 2011 Sep;15(33):1-102. doi:

#### The clinical effectiveness and costeffectiveness of genotyping for *CYP2D6* for the management of women with breast cancer treated with tamoxifen: a systematic review

N Fleeman, 1\* C Martin Saborido, 2 K Payne, 3 A Boland, 1 R Dickson, 1 Y Dundar, 1 A Fernández Santander, 4 S Howell, 5 W Newman, 6 J Oyee 1 and T Walley 7

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<sup>2</sup>School of Nursing and Physiotherapy, Universidad Pontificia Comillas, Madrid, Spain

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<sup>4</sup>Department of Biomedical Sciences, Universidad Europea de Madrid, Madrid, Spain

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7Health Services Research, University of Liv

\*Corresponding author

#### **Conclusions**

#### Poor metaboliser plus intermediate metaboliser versus extensive metaboliser

In the four cohorts that explored OS between these groups of patients, there was no evidence of a difference between PMs + IMs and EMs. However, five out of eight cohorts reported significantly improved outcomes for relapse/recurrence in EMs. Interestingly, in one of these cohorts, reported only as an abstract, the significant differences were found only when using the AmpliChip\* (Roche Molecular Systems) to genotype for an extensive number of alleles and not when four common alleles were tested for.

#### Intermediate metaboliser versus extensive metaboliser

There was no evidence of a difference in OS or relapse/recurrence between IMs and EMs from the only cohort that compared outcomes for these two phenotypes.

#### Poor metaboliser versus extensive metaboliser plus intermediate metaboliser

There was no evidence of a difference in OS or of relapse/recurrence between PMs and EMs + IMs from any of the three cohorts that compared these outcomes in these groups of patients.

#### Poor metaboliser versus extensive metaboliser

From two cohorts, no evidence of a difference in overall survival (OS) between PMs and EMs was reported. However, there was evidence of improved outcomes in terms of relapse/recurrence (disease-free survival, recurrence-free survival or time to recurrence) in the three cohorts that compared these outcomes.

It has not been possible for this review to ascertain whether pharmacogenetic testing for CYP2D6 is clinically effective or cost-effective. Key issues include the fact that it is not clear which alleles should be tested for and how phenotypes should then be derived. Assuming we are able to resolve





## Drugs subject to monitoring

Antibiotics PKIPD Index

Antidepressants

Curvilineal relationship PK/Pogs:

3011eptics Anti-cancer Grogs:

5-Fluouracil

5-tecan Antiepileptics
Imprecision in Ti

**Antipsicotics** Curvilineal relationship PK/PD Irinotecan

Taxanes

Doxorrubicin

MTX

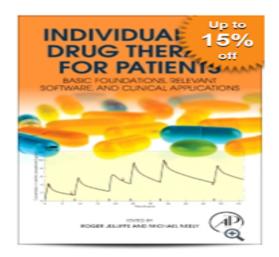
TKI

Monoclonal antibody

Immunotherapy Digestive Rheumatology Dermatology

**Cardiotonics** Need to control peripheral c.

## Individualized Drug Therapy for Patients, 1st Edition Basic Foundations, Relevant Software and Clinical Applications



Editor(s): Jelliffe & Neely
Expected Release 17 Nov 2016

Date:

Imprint: Academic Press Print Book ISBN: 9780128033487

Pages: 434

Dimensions: 235 X 191

This practical guide provides clinical pharmacologists, pharmacists, and physicians with a valuable resource to help move traditional drug therapy beyond a memorized ritual to being a thoughtful quantitative process aimed at optimizing therapy for each individual patient

If you wish to prescribe drugs to your patients with skill, optimal precision, and consideration for each patient's individual needs, to minimize poor outcomes from blunt, last century one-size-fits-all dosing for the fictitious average patient, consider this book.

More information from Googling "Individualized Drug Therapy" at Elsevier or Amazon.

## Take-home messages

- To have good control of every step of the TDM process is necessary for good dosing adjustment of drug therapy in each patient.
- You can't do drug dosing adjustments if you don't control all information, and if you don't use the information correctly.
- Until now, pharmacogenomics has demonstrated a contribution for pharmacodynamic purposes. For dosage optimization, pharmacokinetics is a much more potent tool.



