

# **PHARMACOGENETICS OF BREAST CANCER**

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# Outlines

- Breast cancer therapeutic situation
- Pharmacogenetics of antiestrogen therapy
  - Tamoxifen
- Pharmacogenetics of cytotoxic agents
  - Cyclophosphamide
  - 5-FU
  - Paclitaxel
- Trastuzumab pharmacogenetics
- Summary

# Breast Cancer Facts

- The most common type of cancer and leading cause of cancer death in American women
- Two thirds of invasive breast cancer cases are ER-positive



*Gonzales-Angulo et al 2007; Nicholson et al. 2003*

# BREAST CANCER: Statistics

Estimated New Breast Cancer Cases and Deaths Year 2006			
	Female	Male	Both
Estimated New Cases	212,920	1,720	214,640
Estimated Deaths	40,970	460	41,430

*Jemal A, et al. 2006*

# Treatment Options

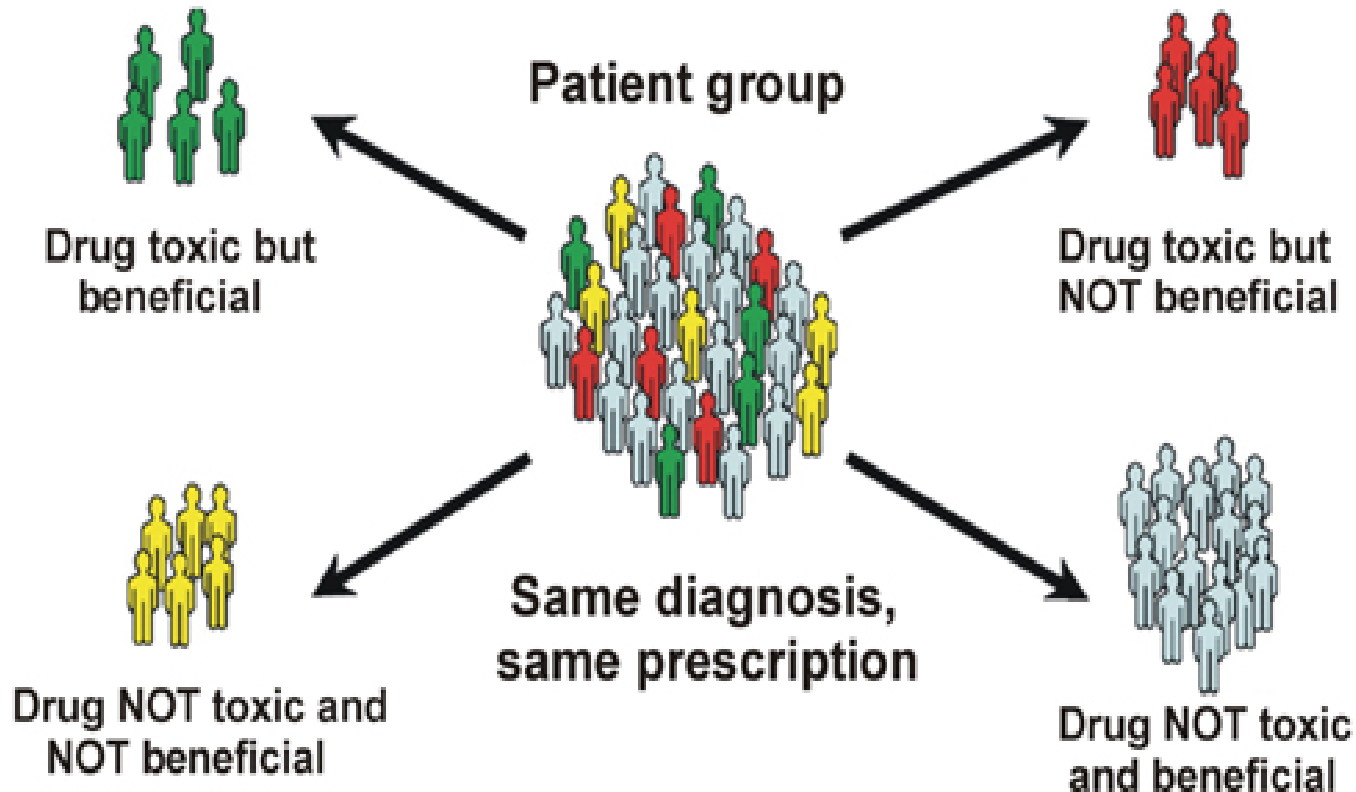
- Antihormonal treatment
  - Tamoxifen, aromatase inhibitors, fulvestrant
- Chemotherapy
  - Mono- poli-chemotherapy
  - Different schemes and schedules (CMF, taxanes, anthracyclines)
- Monoclonal antibodies
  - Herceptin
- Other transuctional pathways inhibitors
  - tyrosine kinase inhibitors, mTOR inhibitors, VEGF inhibitors

# How do we make treatment decisions currently?

- Tumor stage, lymph nodes involvement
- Patient's prior treatment history
- Concomitant illness
- General activity of the drugs and regimens
- ER status
- HER status

**Selection of a particular treatment is not based on the molecular characteristic of the tumor.**

# Therapeutic Situation



# Response Rates

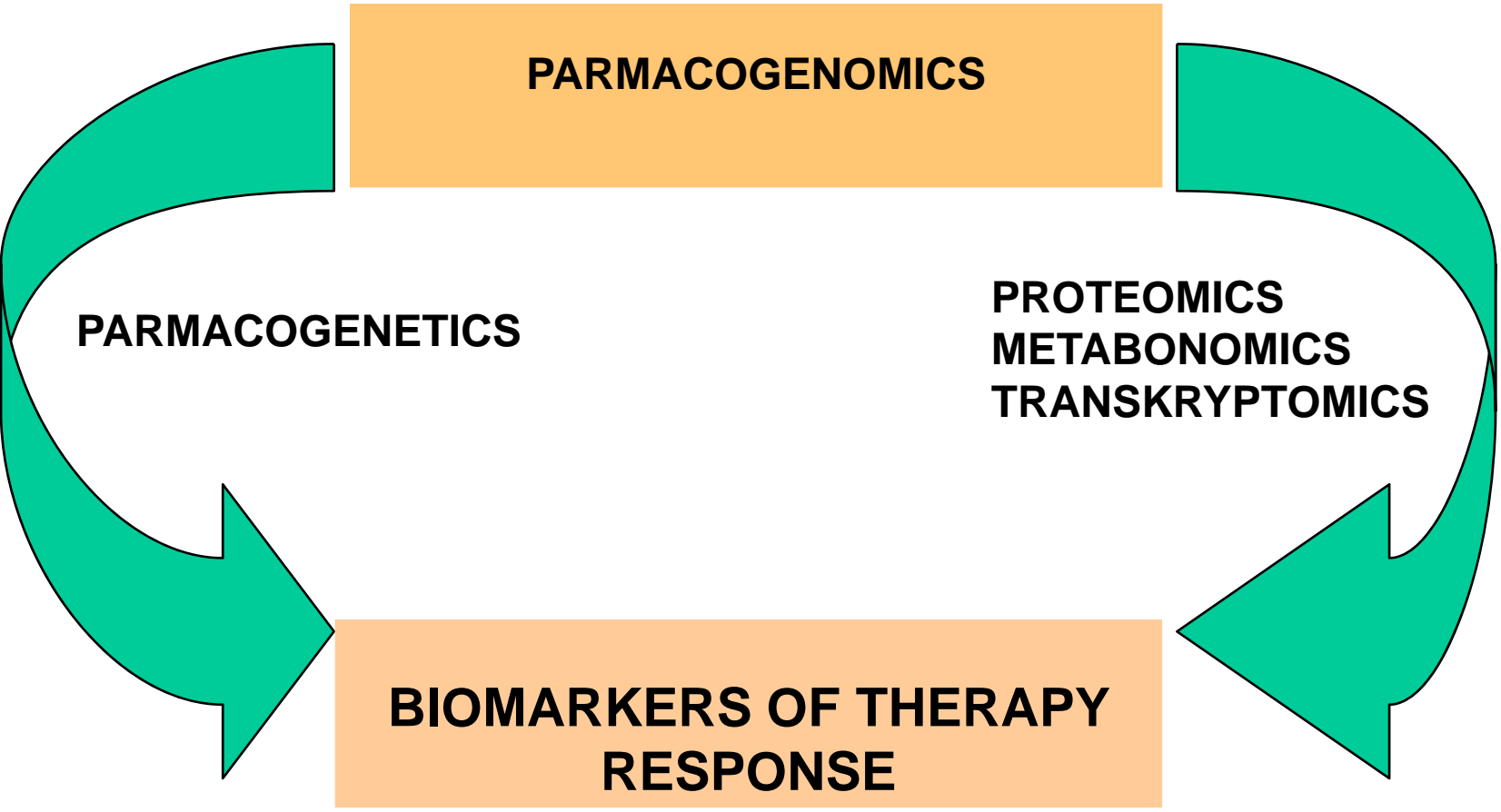
- Up to 65% of breast cancer that initially respond to tamoxifen (TAM) acquire resistance during treatment
- CMF chemotherapy yield a 10-year disease-free survival rate of 28%
- Anthracycline based chemotherapy improves survival rate by 38% in pre- and 20% in post-menopausal patients

**PARMACOGENOMICS**

**PARMACOGENETICS**

**PROTEOMICS  
METABONOMICS  
TRANSCRIPTOMICS**

**BIOMARKERS OF THERAPY  
RESPONSE**



### What is an SNP?

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



### What is an SNP map?

Location of SNPs on human DNA



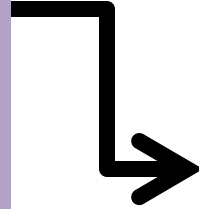
Human DNA

### How can an SNP map be used to predict medicine response?



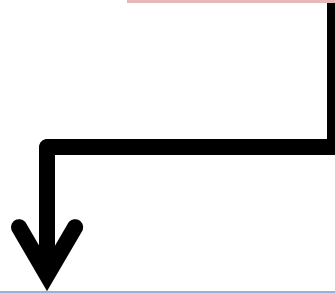
## Mechanism Upstream

- Drug Transport
- Metabolism Phase I
- Metabolism Phase II



## Interaction Drug-Target

- DNA Biosynthesis and Metabolism
- DNA Repair

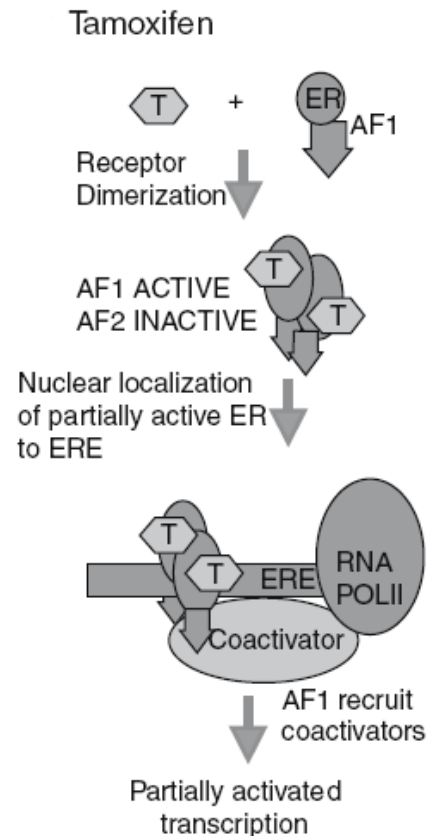
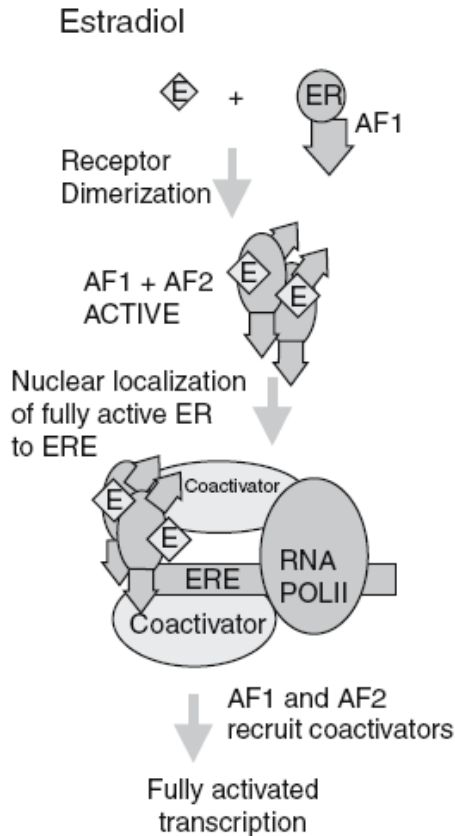


## Mechanism Downstream

- Apoptosis
- Chemokine Activation

# **I. Pharmacogenetics of Antiestrogen Treatments**

# Tamoxifen

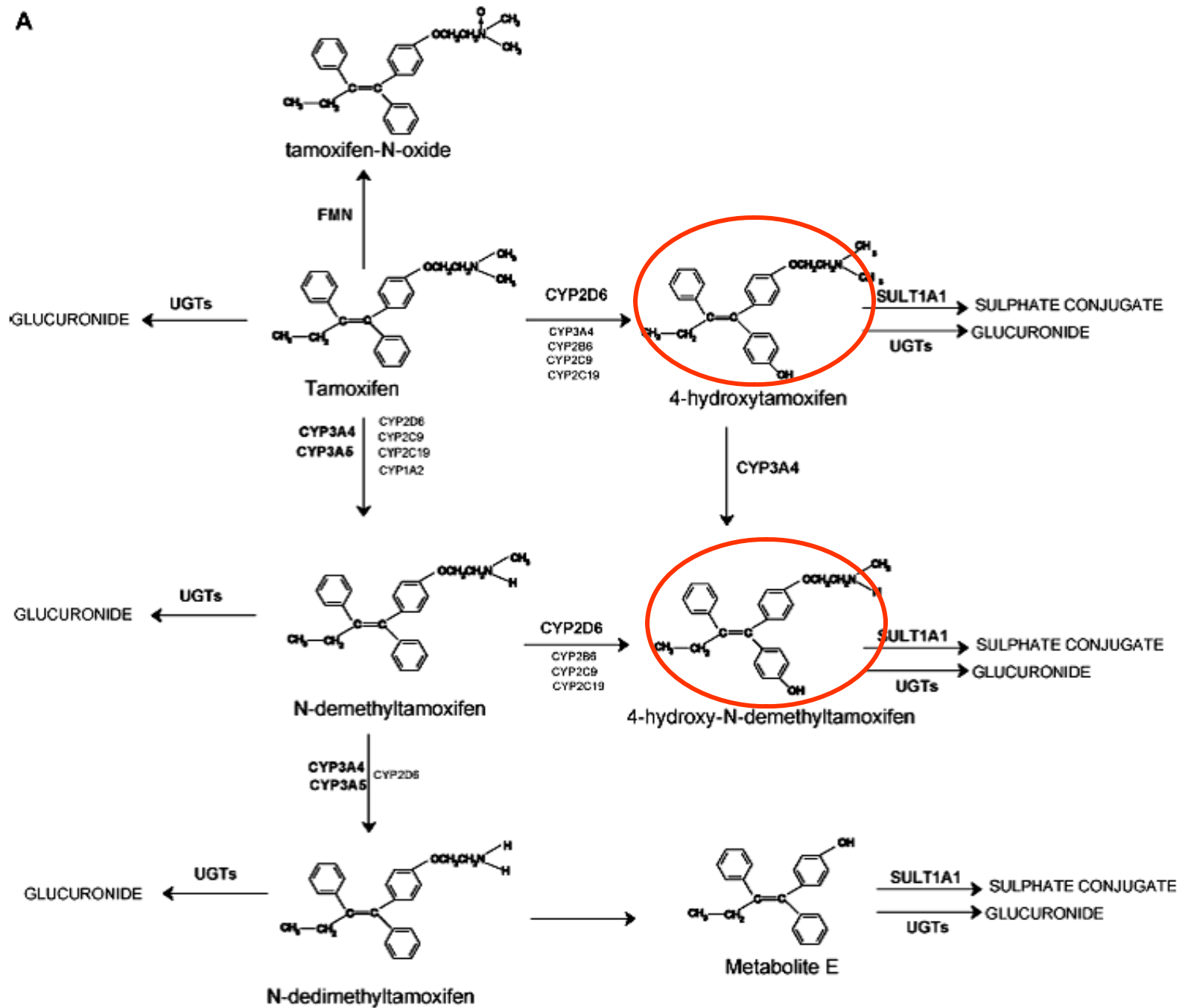


- SERM
- Use by pre- and post-menopausal breast cancer patients
- Prodrug- requires activation

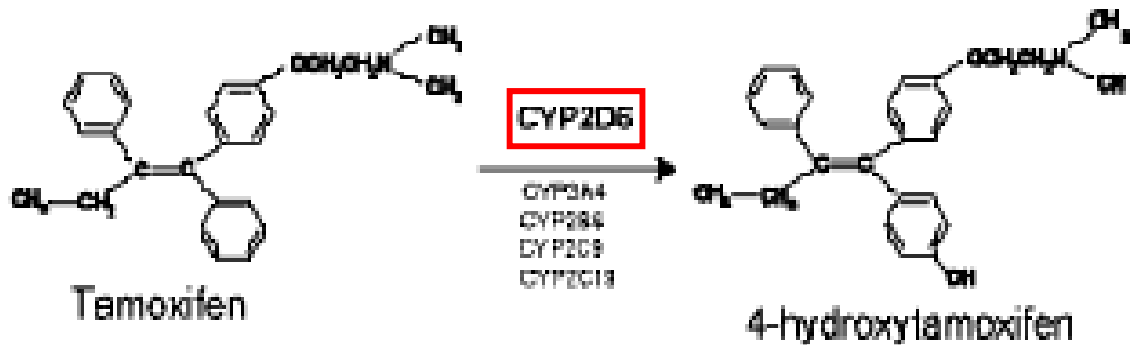
# Tamoxifen- Clinical Assessment

<b>Indications</b>	<b>Year of Approval</b>
<b>Metastatic Breast Cancer (postmenopausal)</b>	<b>1977</b>
<b>Adjuvant Breast Cancer (postmenopausal node +)</b>	<b>1986</b>
<b>Metastatic Breast Cancer (premenopausal)</b>	<b>1989</b>
<b>Adjuvant Breast Cancer (postmenopausal node -)</b>	<b>1990</b>
<b>Metastatic Breast Cancer (male)</b>	<b>1993</b>
<b>Reduction in Breast Cancer Incidence</b>	<b>1998</b>
<b>Ductal Carcinoma in Situ (DCIS)</b>	<b>2000</b>

A



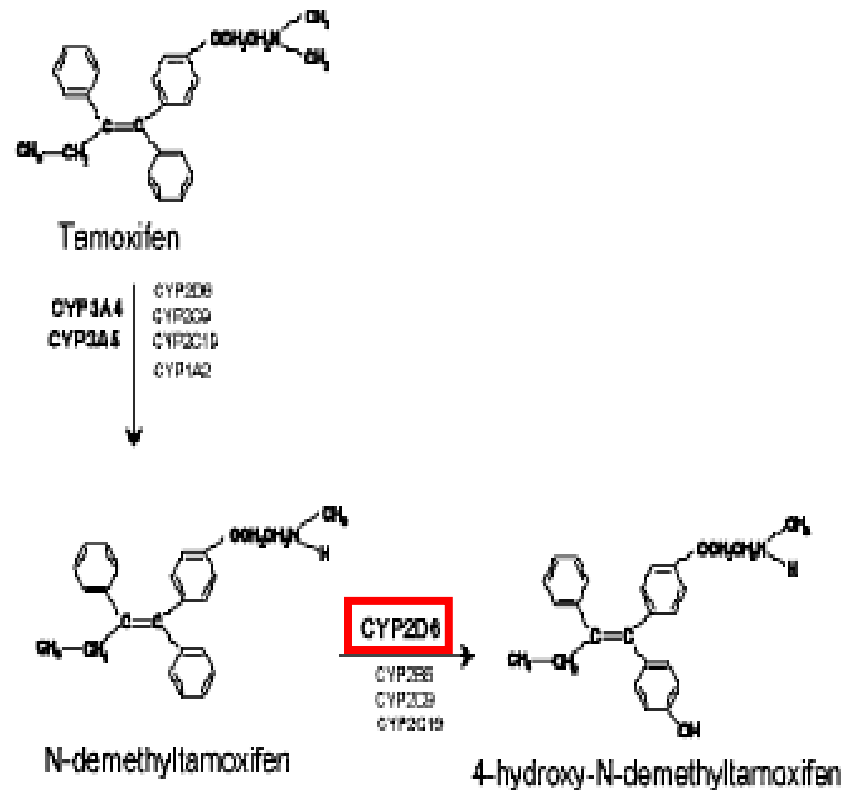
# 4-Hydroxytamoxifen



- 100- fold greater affinity than TAM towards ER
- 30- to 100-fold more potent inhibitor than TAM

# 4-Hydroxy-N-Demethyltamoxifen (Endoxifen)

- Equivalent potency to 4-OH-TAM, but 10-fold higher concentration in plasma
- Impact of CYP2D6 enzyme variant and inhibitors of CYP2D6 on endoxifen concentration.



# CYP2D6 Gene

- Encodes enzyme involved in biotransformation of many drugs
- Highly polymorphic: 63 allelic variants  
[www.cypalleles.ki.se](http://www.cypalleles.ki.se)
- Wide range of metabolic phenotypes: UM, EM, IM and PM
- *CYP2D6*\*4 the most frequent null allele in Caucasians (20-25%), causes 70-90% of all PM

# Metabolic Phenotype

- **Extensive Metabolizers (EM)** have two copies of functional alleles. They have a normal response to the standard dose of a particular drug.
- **Intermediate Metabolizers (IM)** may showed impaired metabolism, by missing one functional allele.
- **Poor Metabolizers (PM)** have problems processing the standard dose of a drug, because they miss two functional alleles. Require different dose.
- **Ultra Metabolizers (UM)** have one or more extra copy of the gene that produce the 2D6 enzyme, so they will require different dose.

# CYP2D6 Alleles

BEVERAGE ET AL.

**Table 1.** Frequencies of the Most Prevalent Alleles Across the Broad Ethnic Groups of Caucasians, Black Africans, and Asians

Allele Type	CYP2D6 Allele	Caucasian (%)	Black African (%)	Asian (%)
Functional	*1	33–40	28–50	23–42
	*2	22–34	11–78	9–20
Reduced function	*9	0–2.9	0	3.3
	*10	1.9–8	3.1–8.6	38–70
	*17	0.1–0.3	9–34	0.5
	*41 <sup>18</sup>	8	—	—
Nonfunctional	*3	1–3.9	0–0.5	0.8–1
	*4	12–23	1.2–7	0–2.8
	*5	1.6–7.3	0.6–6.1	4.5–6.1
	*6	0.7–1	0	—
Duplication	*1 × 2	0.2–0.5	3.3	0.5
	*2 × 2	0.7–1.6	1.6–2.5	0–1
	*4 × 2	0.1–0.2	0.9	—

# Impact of *CYP2D6*\*4 Allele

Study of 180 postmenopausal patients with 5 years of TAM treatment (*Goetz et al. 2007*)

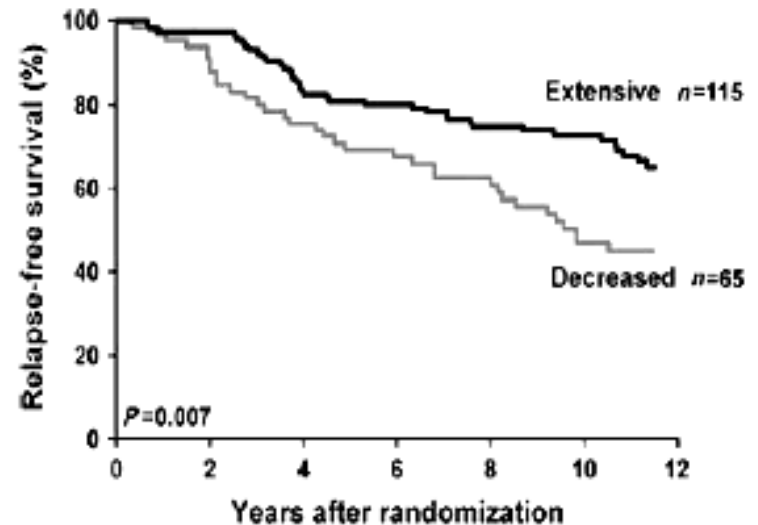
- Patients with decreased *CYP2D6* metabolism

- Shorter time to recurrence

p=0.034    HR=1.91

- Worse RFS

p=0.017    HR=1.74



# Replication Study for *CYP2D6*\*4

Study of 206 postmenopausal patients treated with TAM (*Schroth et al. 2007*)

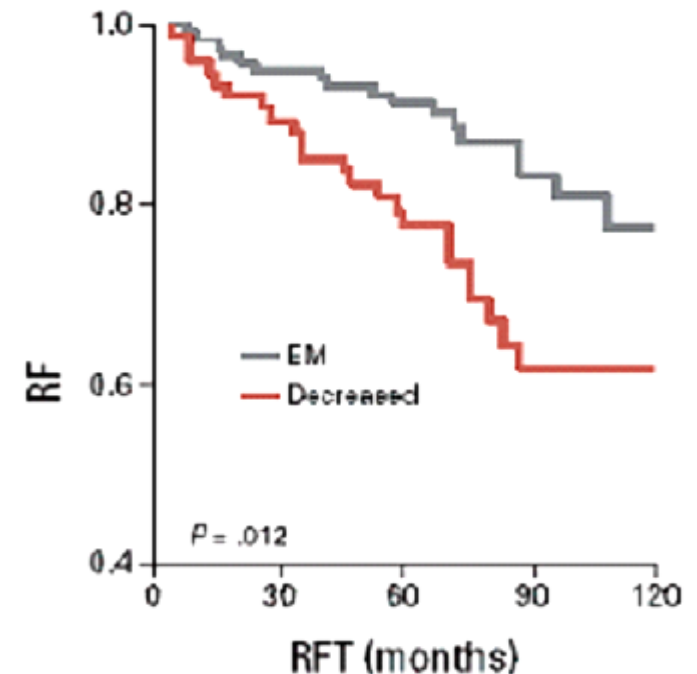
- Patients with decreased *CYP2D6* metabolism

-shorter RFT

p=0.012      HR=2.24

-worse EFS

p=0.02      HR=1.89



# How could CYP2D6 testing change clinical practice?

- Alter therapy course for postmenopausal patients, who are carriers of *CYP2D6\*4* allele
- Development of treatment algorithms for carriers of CYP2D6 alleles variants
- Avoiding co-administration of certain antidepressants with TAM, since they may act as CYP2D6 inhibitors

# Tamoxifen and an FDA Perspective



## FDA Recommends Relabeling Tamoxifen

On October 18, an FDA panel unanimously agreed that the 2D6 gene is a predictor of tamoxifen efficacy. They recommend relabeling tamoxifen to say that 2D6 poor metabolizers who take tamoxifen have a higher risk for breast cancer recurrence, and that testing is available.

# Roche AmpliChip™ CYP450



- FDA & EU cleared for diagnostic analysis of two critical drug metabolism genes: **CYP2D6** and **CYP2C19**
- Pharmacogenetics microarray- based test
- Recommended for cyclophosphamide (CYP2C19) and tamoxifen (CYP2D6)



Tag-It™ Mutation Detection Kit for  
P450-2D6 v2

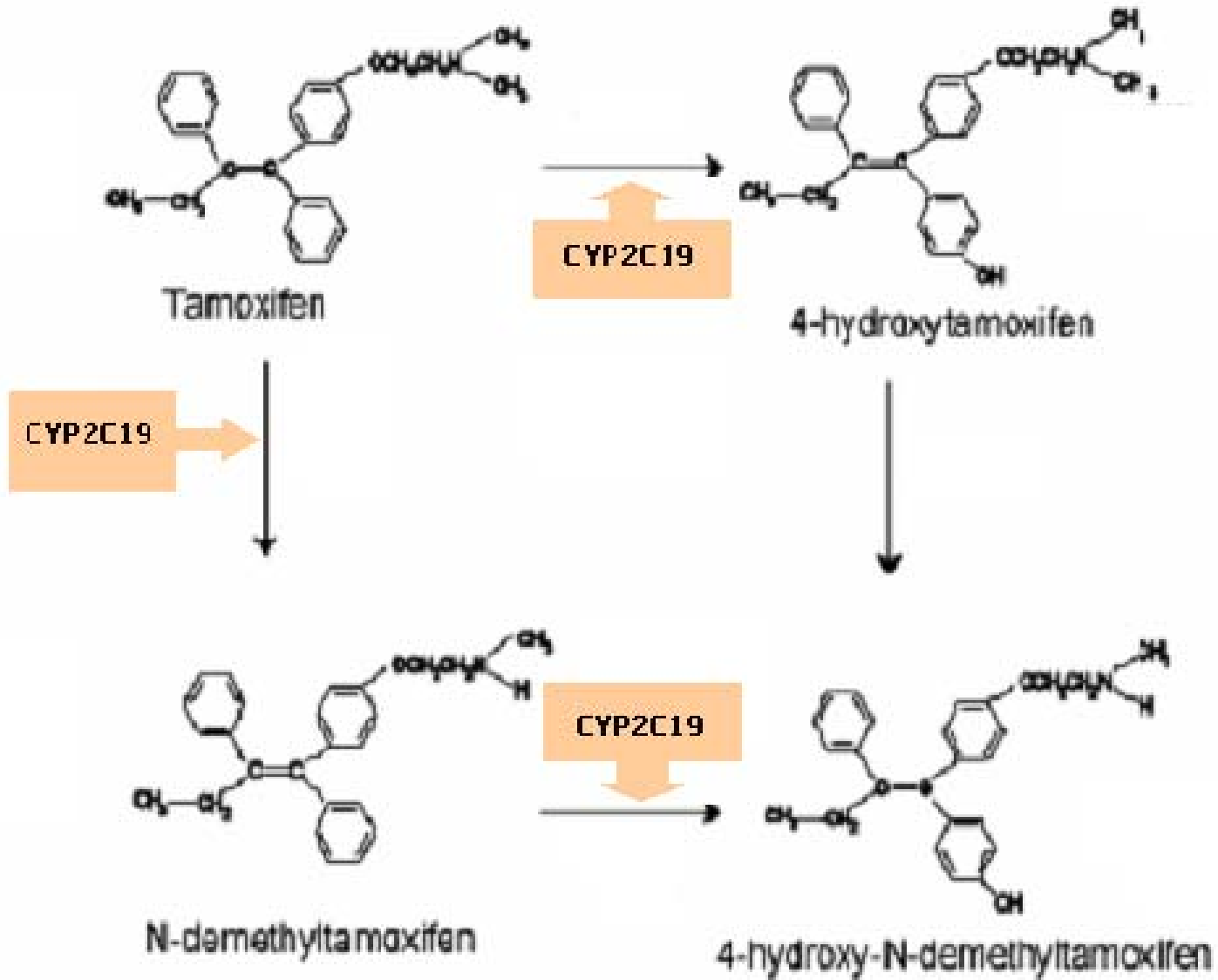
For use with Luminex® 100 or 200 xMAP® System

Variant <sup>†</sup>	* Designation	Effect
/	Gene Duplication	↑ Activity
/	*5 Gene Deletion	No Gene
-1584C>G	*2A Promoter	Expression
100C>T	*4 (A-L) <sup>a</sup> *10 (A,B)	P34S
124G>A	*12	G42R
138InsT	*15	Frameshift
883G>C	*11	Splicing
1023C>T	*17	T107I
1661G>C <sup>b</sup>	*2	Silent
1707T>del	*6 (A-D)	Frameshift
1758G>T	*8	Stop Codon
1758G>A	*14	G169R
1846G>A	*4 (A-L) <sup>a</sup>	Splicing
2549A>del	*3 (A,B)	Frameshift
2613-2615 delAGA	*9	K281Δ
2850C>T <sup>b</sup>	*2, *17	R296C
2935A>C	*7	H324P
2988G>A	*41	mRNA Stability
4180G>C <sup>b</sup>	*2	S486T

<sup>†</sup> Nomenclature derived from CYP2D6 Allele Nomenclature Database<sup>3</sup>

<sup>a</sup> For \*4, no \*4I designation exists

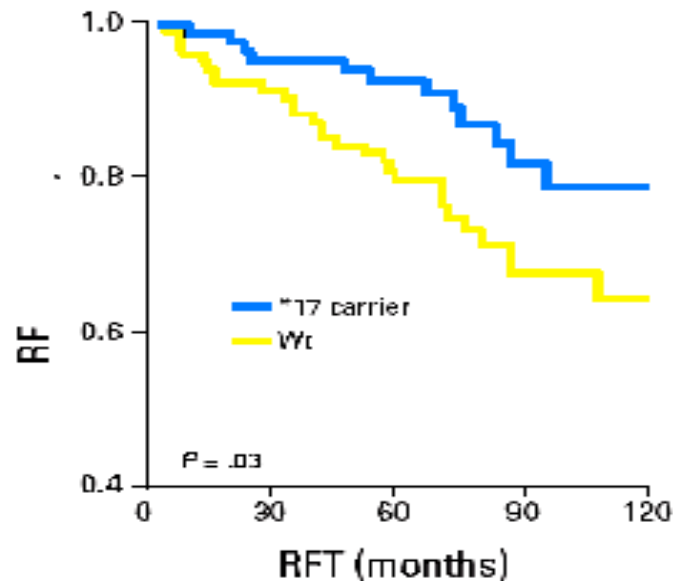
<sup>b</sup> 1661G>C, 2850C>T, and 4180G>C are found in a wide range of alleles



# Impact of *CYP2C19*\*17 Allele

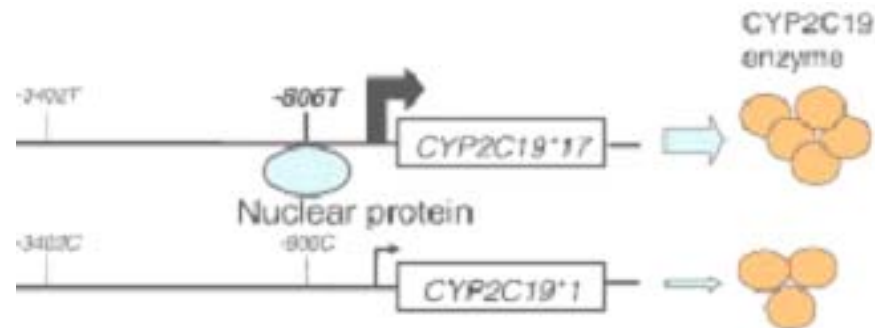
Carriers of *CYP2C19*\*17 allele showed (Schroth et al. 2007)

- Decreased risk of relapse  $p=0.02$  OR=0.43
- Favorable RFT  $p=0.03$  HR=0.45



# *CYP2C19\*17*

- Associated with increased CYP2C19 activity
- UM phenotype
- Carriers of this allele include 18% of Swedes and Ethiopians, but only 4% Chinese



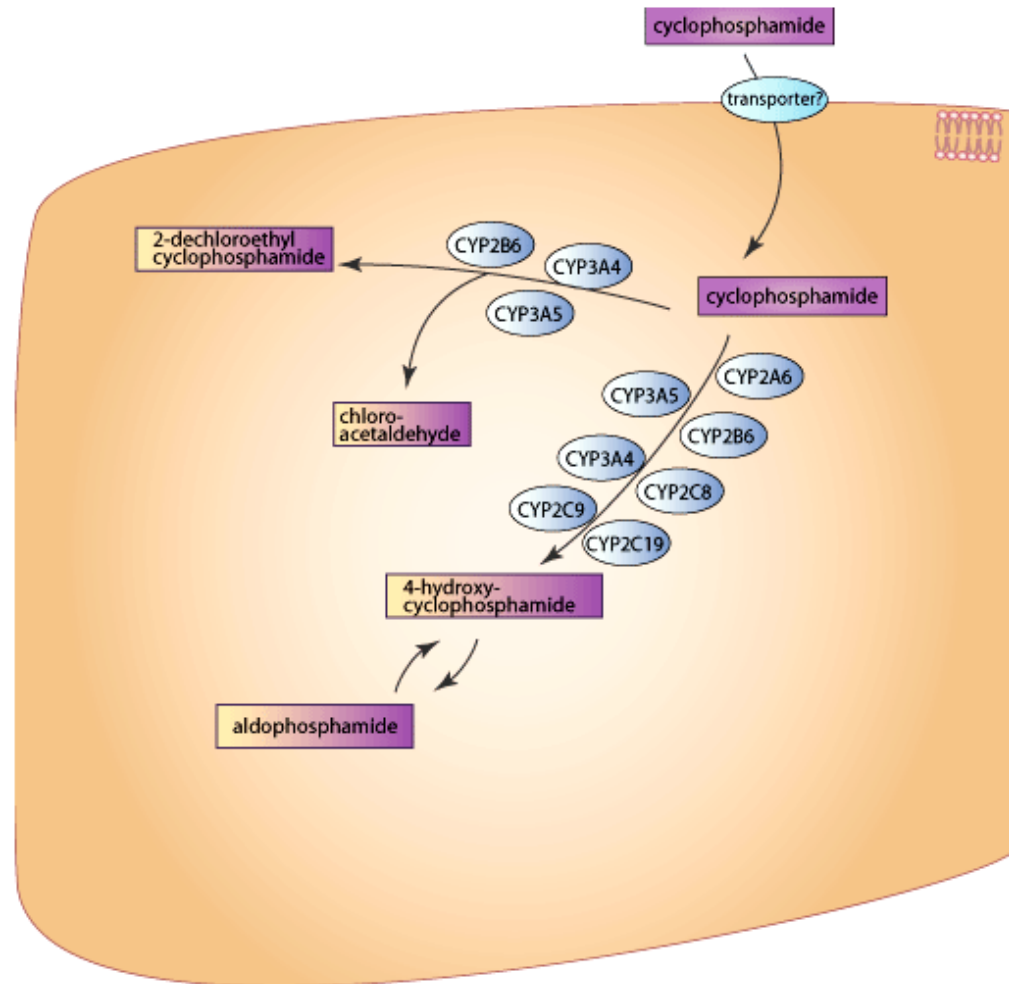
## **II. Pharmacogenetics of Cytotoxic Agents**

# Chemotherapy

- Cyclophosphamide-methotrexate-fluorouracil (CMF)- “golden standard”
- Other agents i.e. taxanes, anthracyclines (FEC, FAC).
- Adjuvant chemotherapy is considered for more than 70% of BC, but is not beneficial for all of them
- Response rates
- Toxicity

# Cyclophosphamide

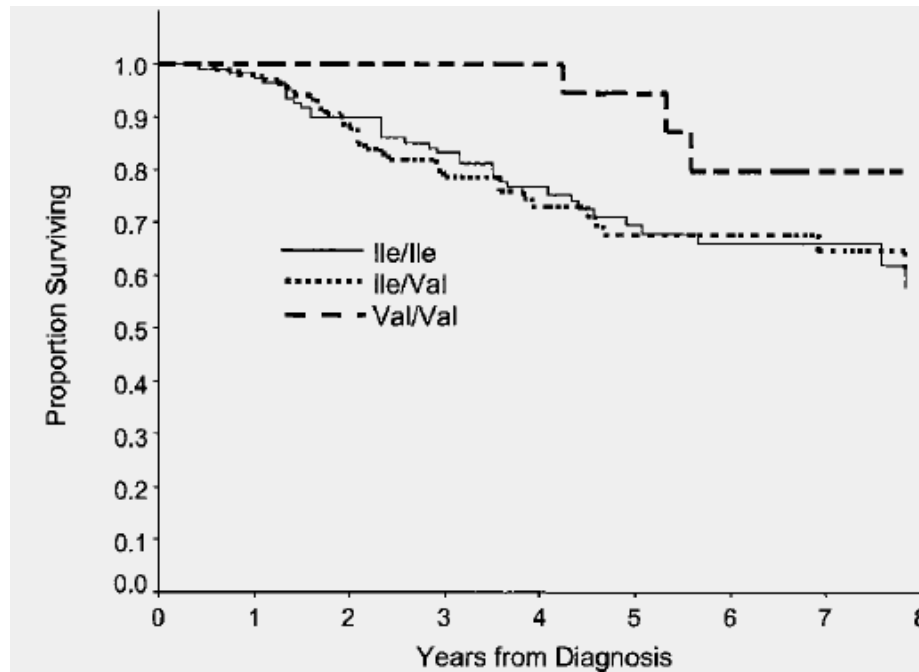
- Alkylating agents
- Prodrug- requires activation by P-450 enzymatic complex
- Eliminated by glutathione S-transferases (GST) A1 and P1



# Cyclophosphamide and GSTP1

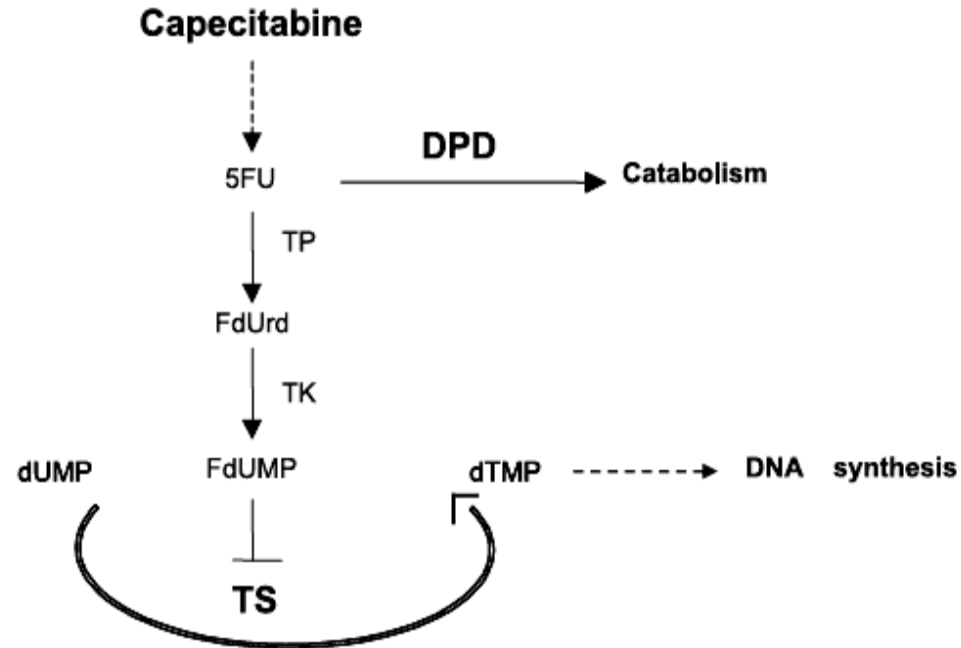
Study of 240 patients treated with cyclophosphamide  
(*Sweeney et al. 2000*)

- Low- activity Val/Val GSTP1 genotype showed better survival      HR=0.3



# 5-Fluorouracil

- Uracil analogue
- Prodrug- active form FdUMP
- TS is a main target
- 85% inactivated by DPD
- Toxicity caused by DPYD polymorphisms

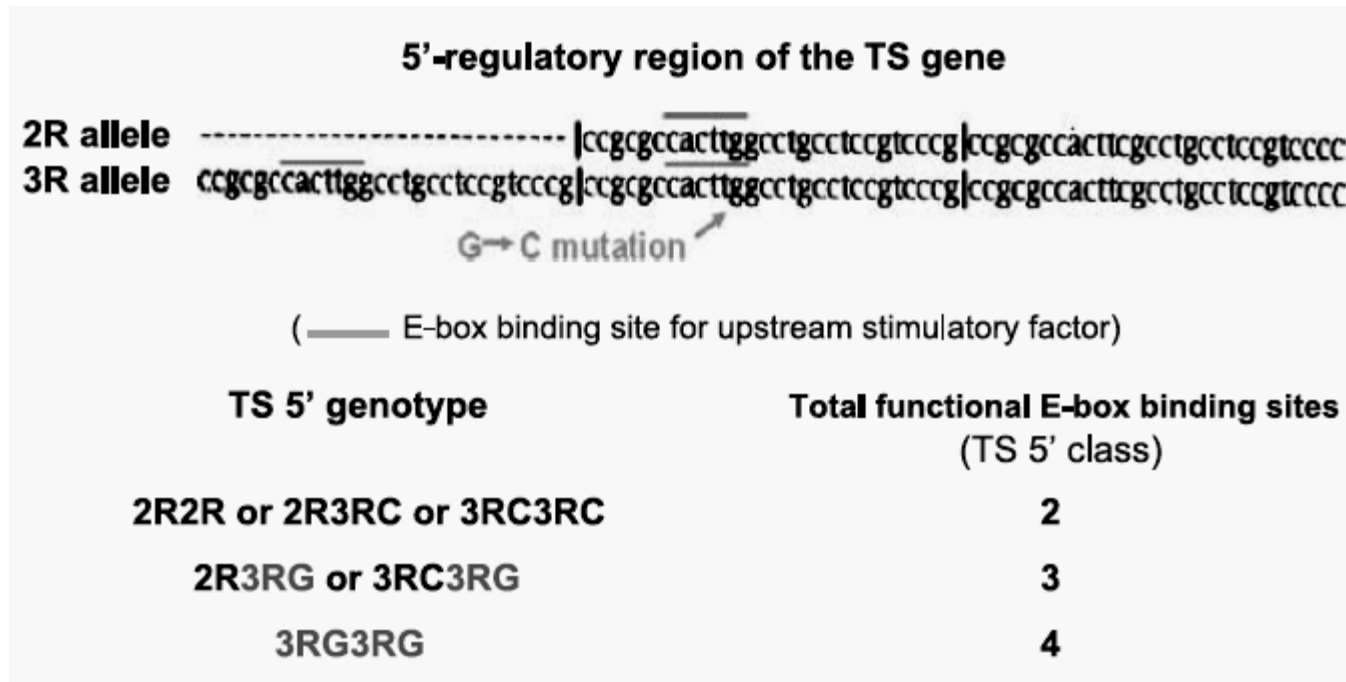


# 5-FU Toxicity and DPYD Polymorphisms

- Wide variation of enzyme activity among individuals
- At least 20 polymorphic loci are associated with reduced DPD activity, and 3-5% of population are heterozygous carriers of deficient allele
- *DPYD\*2A* allele accounts for ~ 50% of known nonfunctional DPYD alleles
- Reduced DPD activity leads to 5-FU toxicity

# TS and 5-FU

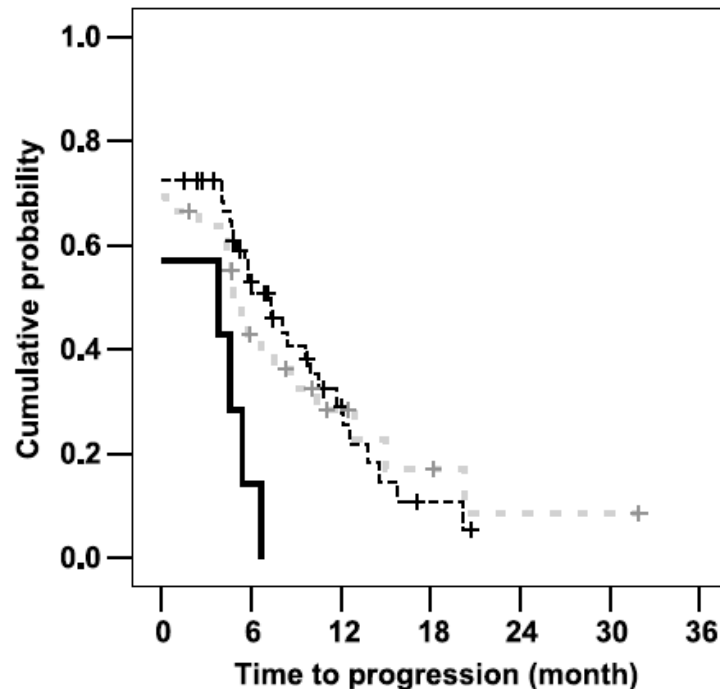
- Inhibition of TS by 5-FU  $\longrightarrow$  cell death
- Polymorphisms G/C in 28bp tandem repeat sequence of TS increases gene expression



# TS and 5-FU

Carriers of TS 5'genotype class 4 (3RG3RG) showed  
(Largillier et al. 2006)

- rapid disease progression ( $p=0.037$ )
- trend toward a higher global toxicity grade 3 and 4



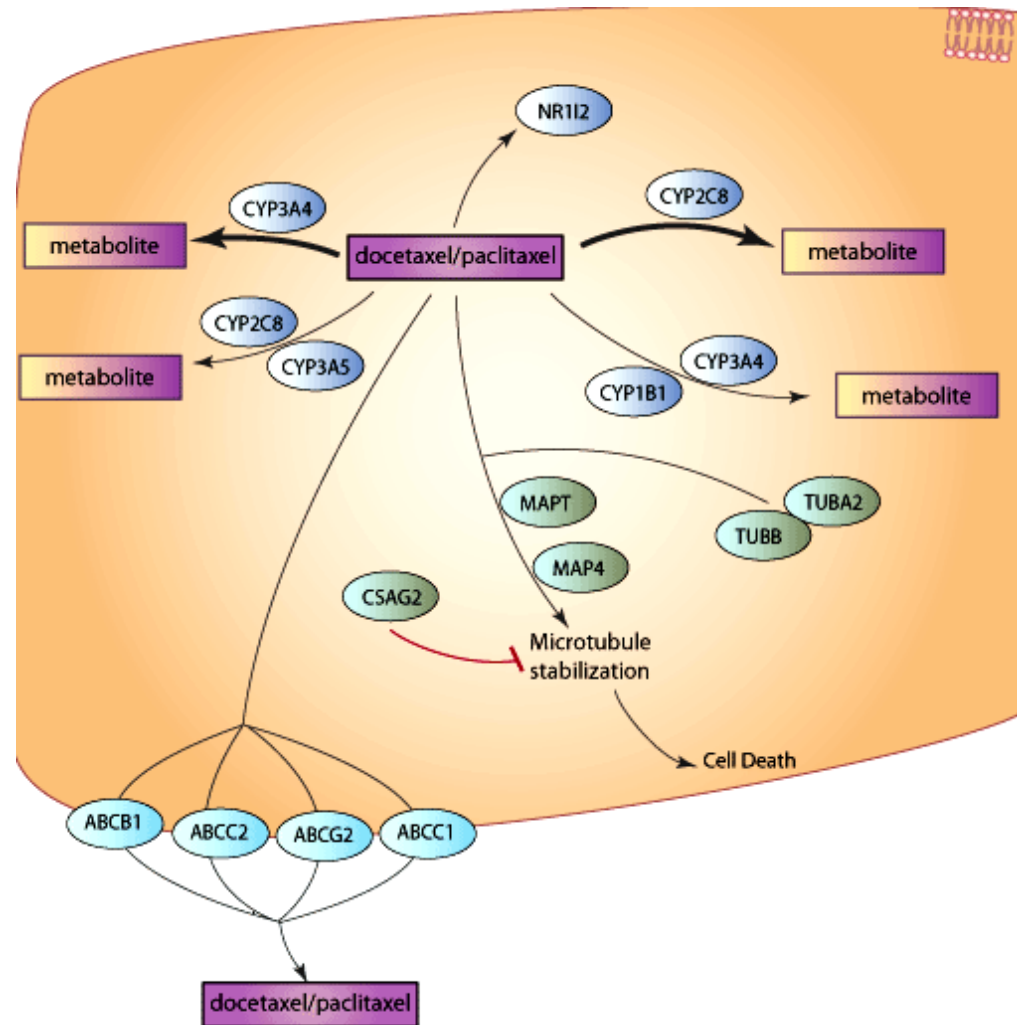
# Commercial Testing



## TherapID™: 5-FU test

- is a simple test that analyzes the genetic variations in the **DPYD** gene in order to predict the risk for serious side effects in patient treated by 5-FU.

# Paclitaxel



- Block cell cycle progression
- Broad spectrum of activity against solid tumors
- Dose dependent induced toxicity

# Paclitaxel Resistance

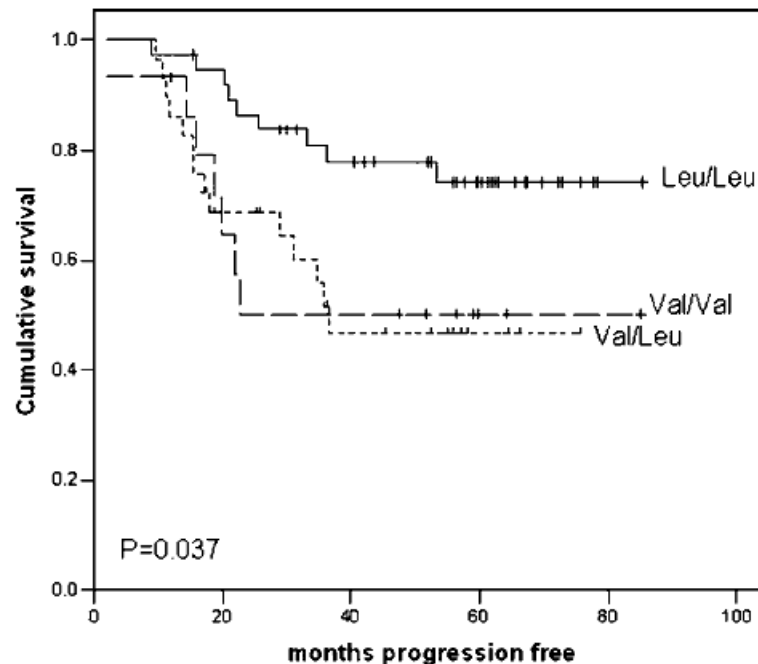
Molecular Mechanism	Implicated in paclitaxel resistance
P-glycoprotein [17–19]	yes
Tubulin mutations and isoforms [24, 28, 1]	yes
Impaired apoptotic response (bcl-2, bax) [49, 50]	yes
Alterations in microtubule-binding proteins (stathmin, tau) [39, 42, 51]	yes

# Paclitaxel and *CYP1B1*\*3 Allele

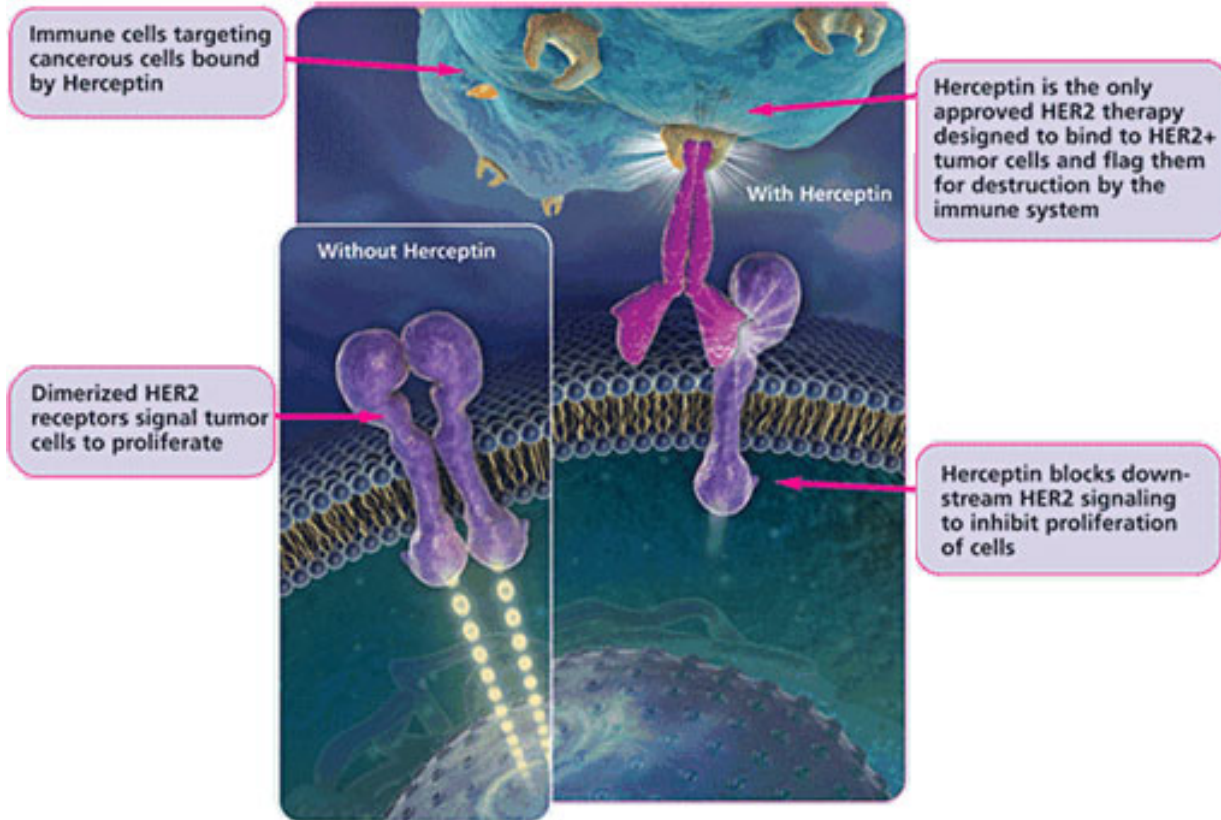
Study of 93 patients with primary breast cancer treated by paclitaxel (*Marsh et al.2007*)

- *CYP1B1*\*3 (L432V) allele significantly associated with progression- free survival

p= 0.037



# Herceptin

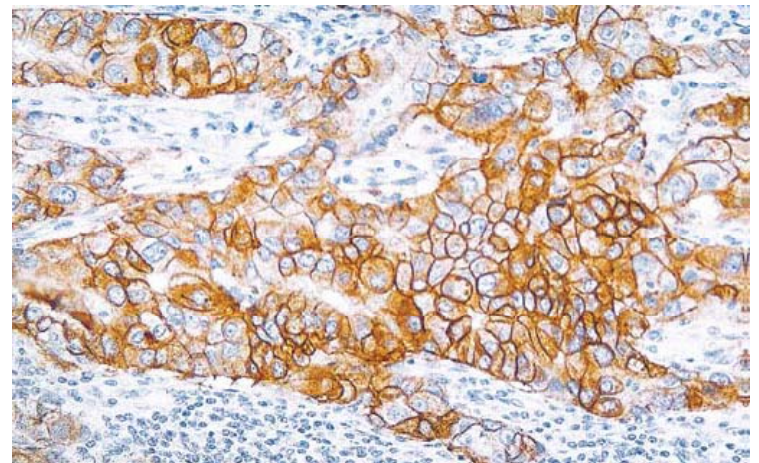
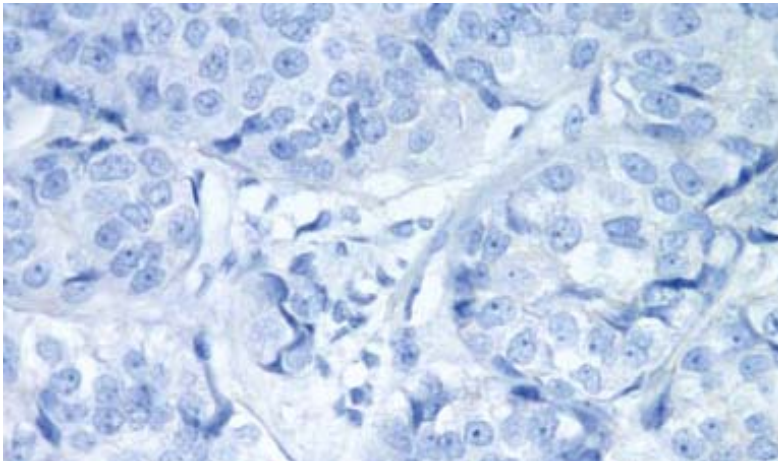


Monoclonal antibody targets an extracellular domain of the HER-2 receptor

# HercepTest™



- The only FDA-approved diagnostic method to aid in the selection of patients for Herceptin therapy is immunohistochemistry (IHC).

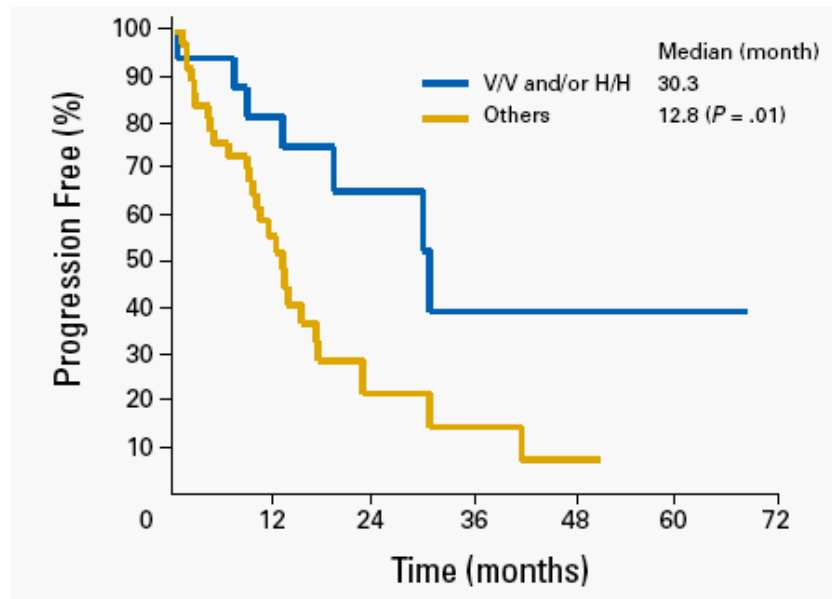


# Trastuzumab

FcγR polymorphisms associated with PFS

(Musolino et al. 2008)

- FcRIIIa 158 V/V and FcRIIa 131 H/H genotype
- p=0.01



# Strategies to Overcome Therapy Resistance

