



# The 7th International Electronic Conference on Medicinal Chemistry (ECMC 2021)

01-30 NOVEMBER 2021 | ONLINE

## Polymorphisms in estrogenic metabolic pathway genes: clinical significance and lifetime outcome.

**Micaela Almeida<sup>1,\*</sup>, Mafalda Soares<sup>1</sup>, José Fonseca Moutinho<sup>1,2</sup>, António Polónia<sup>3,4,5</sup>, Ana Cristina Ramalinho<sup>1,2</sup>, and Luiza Breitenfeld<sup>1</sup>**

<sup>1</sup> Centro de Investigação em Ciências da Saúde (CICS), Faculdade de Ciências da Saúde, Universidade da Beira Interior (UBI), Avenida Infante D. Henrique 6200-506 Covilhã, Portugal

<sup>2</sup> Centro Hospitalar Universitário da Cova da Beira (CHUCB), Quinta do Alvito 6200-251 Covilhã, Portugal.

<sup>3</sup> Laboratório de Anatomia Patológica, Ipatimup Diagnósticos, Rua Júlio Amaral de Carvalho, Porto 45, 4200-135, Portugal

<sup>4</sup> Faculdade de Medicina, Universidade do Porto, Alameda Prof Hernâni Monteiro, Porto 4200-319, Portugal

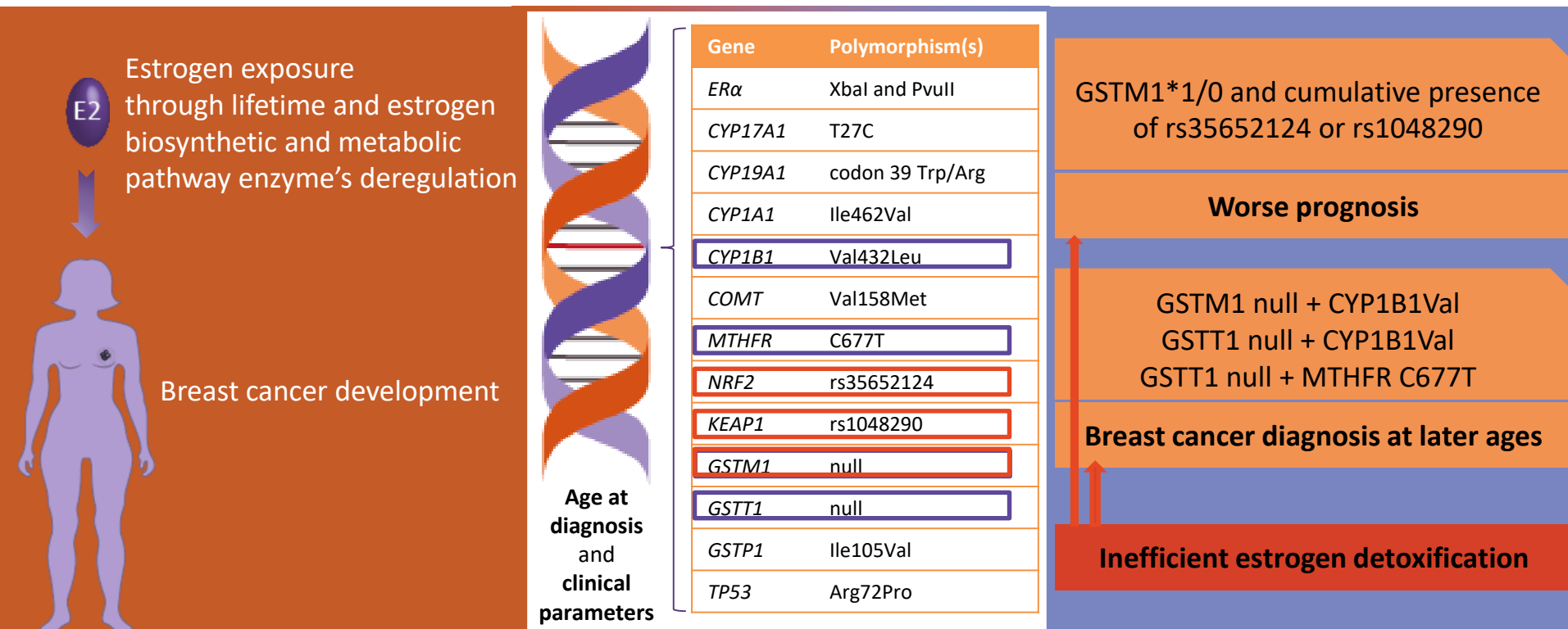
<sup>5</sup> Instituto de Investigação e Inovação em Saúde (i3S), Universidade do Porto, Rua Alfredo Allen, 208, Porto 4200-135, Portugal

\* Corresponding author: [micaelacpalmeida@gmail.com](mailto:micaelacpalmeida@gmail.com)



# Polymorphisms in estrogenic metabolic pathway genes: clinical significance and lifetime outcome.

## Graphical Abstract



**Abstract:** Polymorphisms in estrogenic metabolic pathway genes: clinical significance and lifetime outcome.

Lifetime exposure to high estrogens levels and deregulation of enzymes of estrogen biosynthetic and metabolic pathway are considered breast cancer risk factors. Our research team has previously evaluated, in women with hormone-dependent breast cancer, *NRF2* and *KEAP1* polymorphisms (alone or in association), with breast cancer prognosis, in cases with *GSTM1*-present genotype. We identified that *GSTM1*\*1/0 genotype and cumulative presence of at least one allele mutated in *KEAP1* and/or *NRF2* polymorphisms might be associated with worse prognosis. Moreover, older women were carriers of both *GSTM1* null and *CYP1B1*Val genotypes, similar results were found for the cumulative presence of *GSTT1* null and *CYP1B1*Val genotypes and for *MTHFR* C677T and *GSTT1* null polymorphism. These results lead us to genotype other enzymes involved in estrogen biosynthetic/metabolic pathways, namely: *CYP17A1* T27C, *CYP19A1* codon 39 Trp/Arg, *COMT* Val158Met, *CYP1A1* Ile462Val, *GSTP1* Ile105Val, *ER $\alpha$*  PvuII and XbaI polymorphisms and TP53 Arg72Pro (DNA damage signalling and repair pathway) polymorphism. The genotyping analysis was evaluated, alone or in association, with breast cancer prognosis and with age at breast cancer diagnosis, we found no association ( $p$ -value>0.05). These results emphasize that polymorphisms in *NRF2/KEAP1* complex and the null polymorphisms of *GSTM1/GSTT1* should be further investigated as targets for breast cancer therapy. In addition, polymorphisms related to inefficient estrogen detoxification might be a trigger to hormone-dependent breast cancer development at later ages, and should be considered in breast cancer risk assessment models.

**Keywords:** Breast Cancer, estrogens, polymorphisms, risk model, targeted therapy



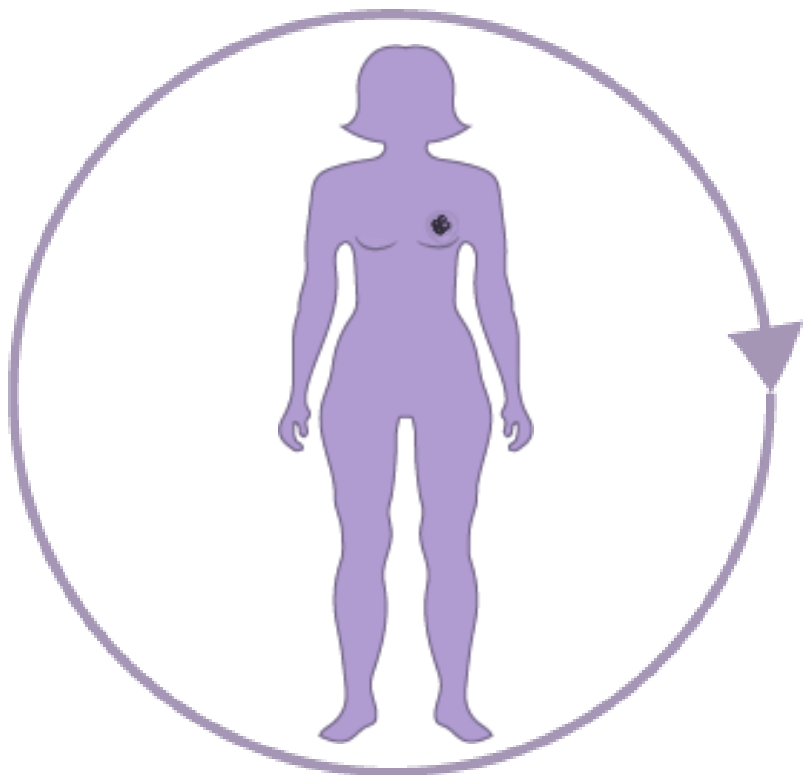
The 7th International Electronic Conference on Medicinal Chemistry

01-30 NOVEMBER 2021 | ONLINE



# Introduction

Estrogens have been identified as carcinogens, and most of the risk factors for breast cancer are related to an increased or prolonged exposure to estrogens.



Biosynthetic pathway

Metabolic pathway

Estrogen signalling  
Estrogen Receptor (ER)

Apoptosis inhibition



The 7th International Electronic Conference on Medicinal Chemistry

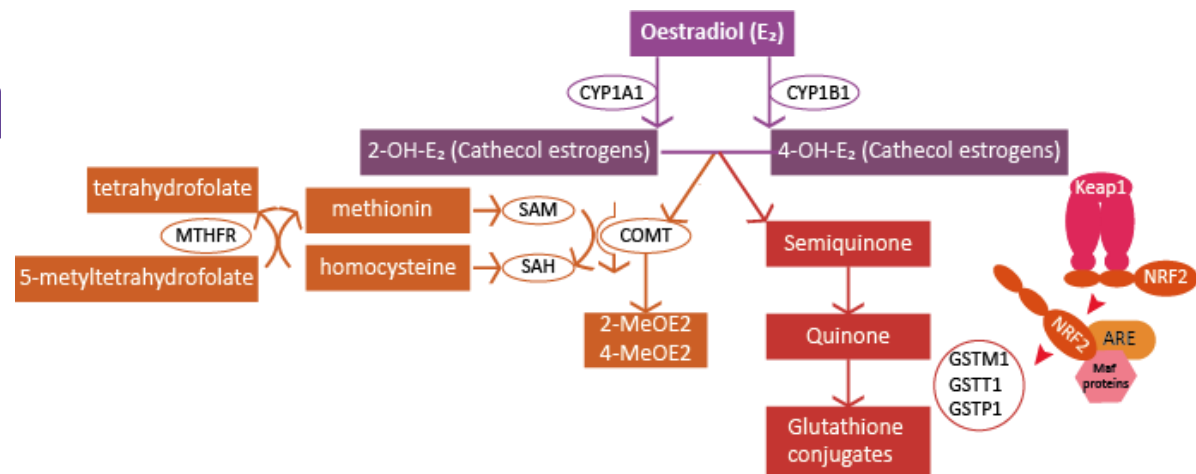
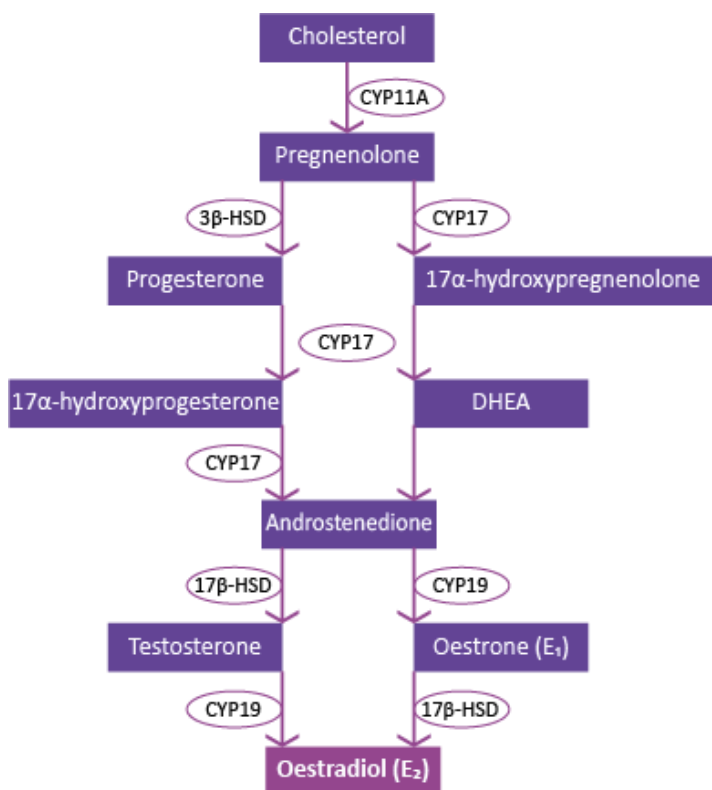
01-30 NOVEMBER 2021 | ONLINE



# Introduction



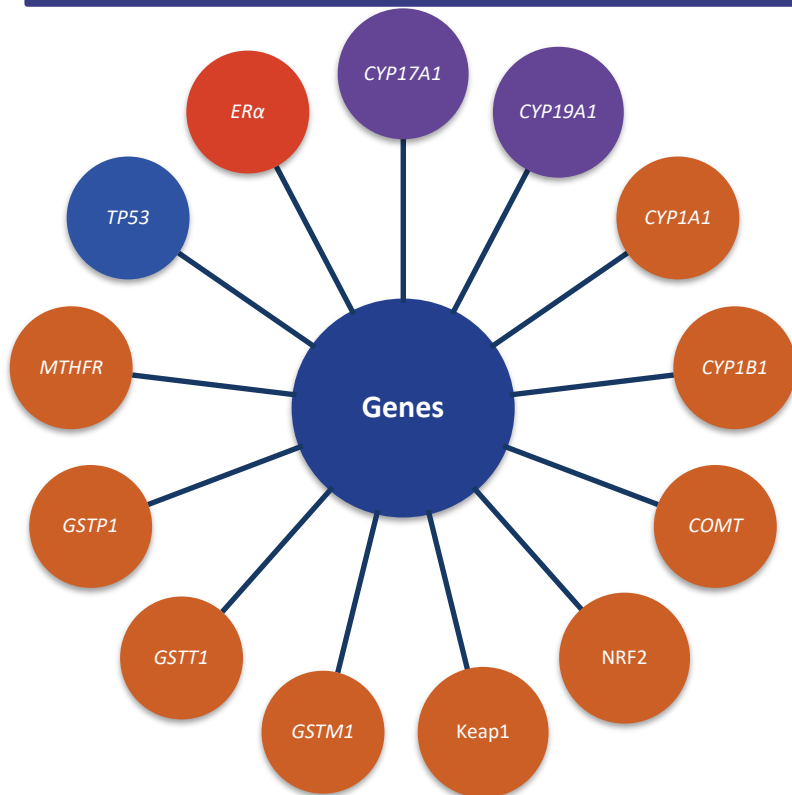
## Biosynthetic and metabolic pathways



# Introduction



## Genes



Biosynthetic pathway

Metabolic pathway

Estrogen signalling  
Estrogen Receptor (ER)

Apoptosis inhibition

Free-radical-mediated DNA damage, single-strand breaks, estrogen–DNA adducts formation, protein oxidation and lipid peroxidation, what triggers genetic instability and cellular damage.



The 7th International Electronic Conference on Medicinal Chemistry

01-30 NOVEMBER 2021 | ONLINE



# Introduction



## Framework

Our research team correlated:

GSTM1 and GSTT1 null

CYP19A1 arginine (C) allele

TP53 proline allele

Increased risk of breast cancer development.

Carriers of xx genotype of  
Er $\alpha$  XbaI

Reduced risk of breast cancer development.



The 7th International Electronic Conference on Medicinal Chemistry

01-30 NOVEMBER 2021 | ONLINE



# Introduction



## Framework

*GSTM1* null polymorphism taking into concern the three genotypes:

- *GSTM1*\*1/1 (present)
- *GSTM1*\*1/0 (heterozygous)
- *GSTM1*\*0/0 (null)

*NRF2* 35652124  
and  
*KEAP1* rs1048290

Clinical parameters in order to evaluate breast cancer prognosis

*GSTM1* and *GSTT1* null polymorphism with *CYP1B1* Val432Leu and *MTHFR* C677T.

Age at diagnosis of breast cancer



The 7th International Electronic Conference on Medicinal Chemistry

01-30 NOVEMBER 2021 | ONLINE

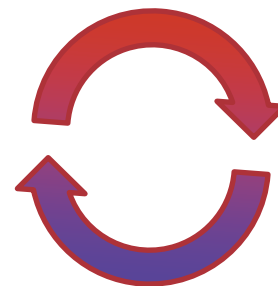
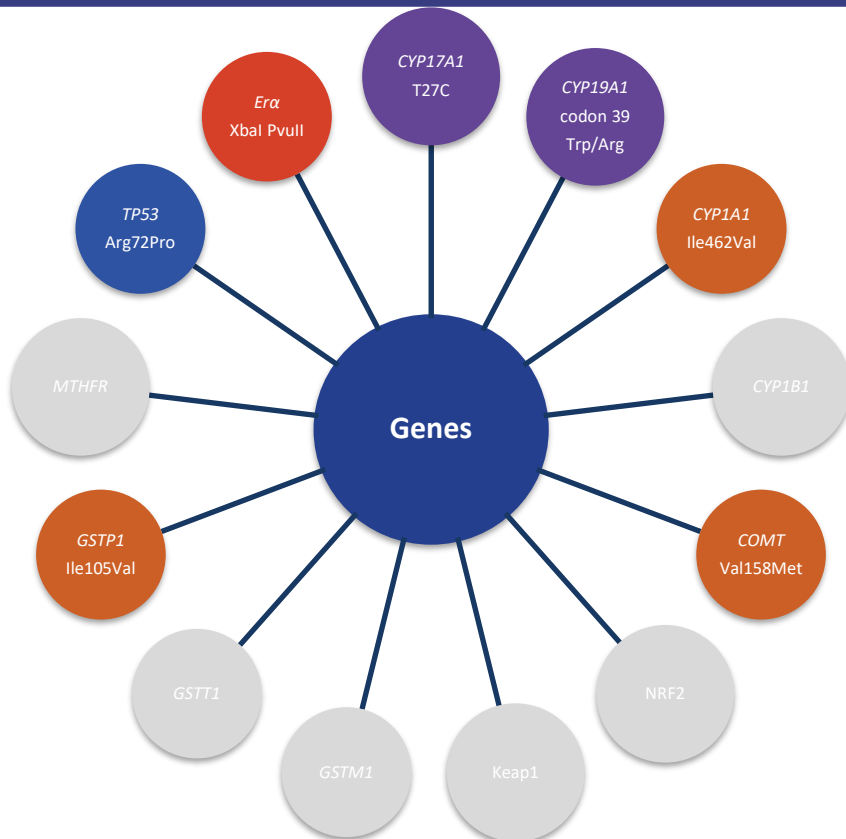




# Introduction



## Framework



Clinical parameters in order to evaluate the breast cancer prognosis

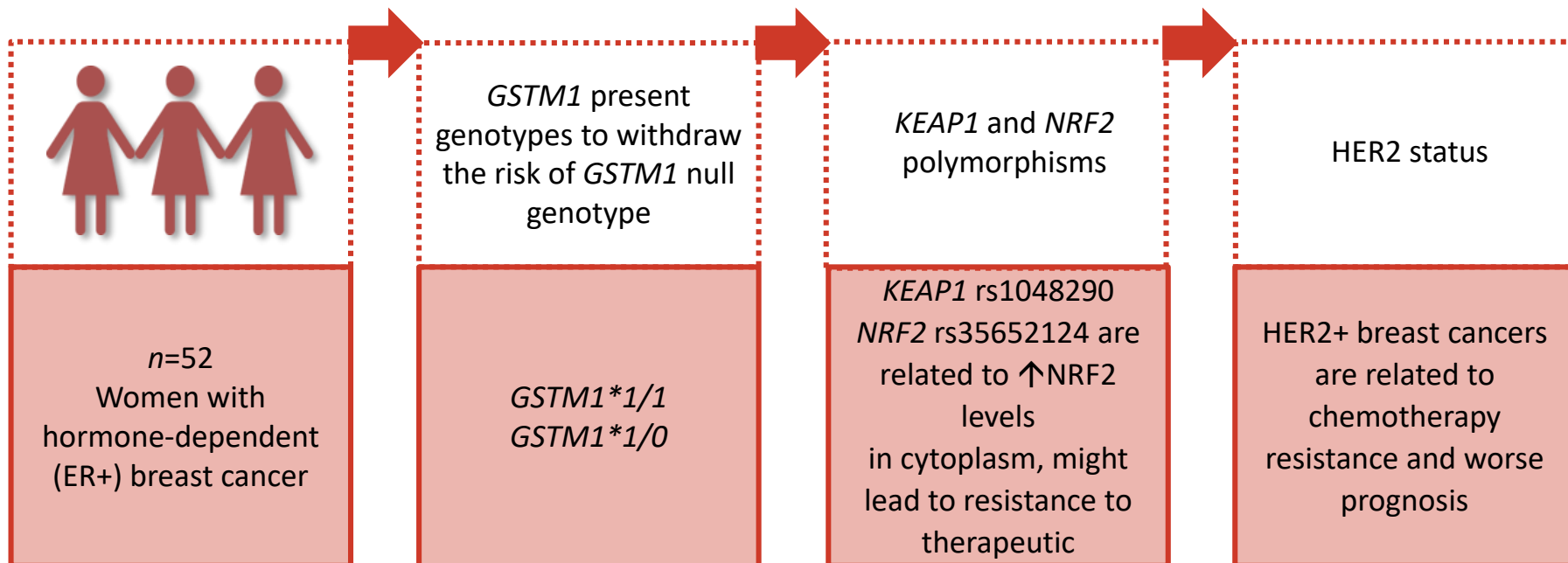
Age at diagnosis



# Results and discussion



Metabolism regulators, phase II enzymes and breast cancer prognosis



The 7th International Electronic Conference on Medicinal Chemistry

01-30 NOVEMBER 2021 | ONLINE



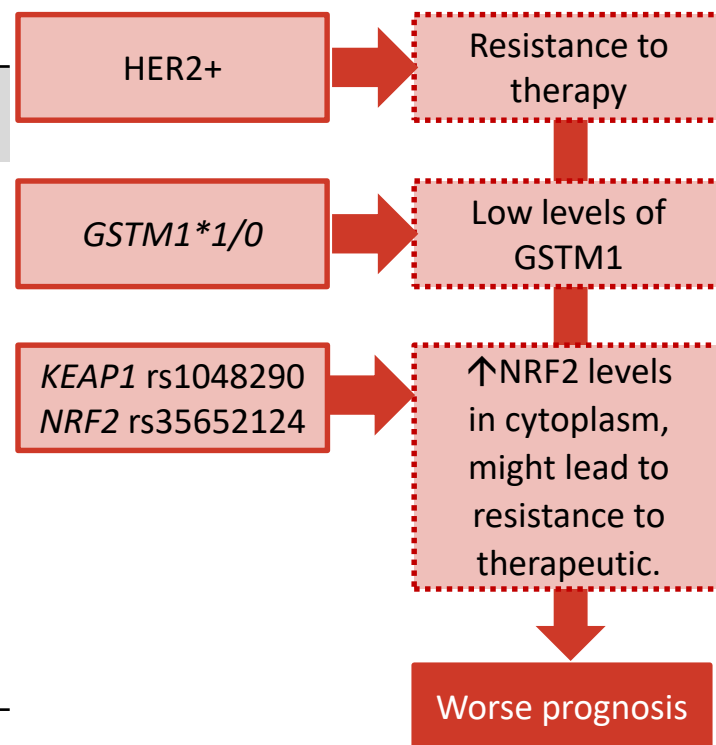
# Results and discussion



Metabolism regulators, phase II enzymes and breast cancer prognosis

Association of *KEAP1* and *NRF2* polymorphisms in cases with *GSTM1* present genotype (\*1/1 or \*1/0) and correlation with HER2 status

Genotype		HER2 status n (%)		P value
		+	-	
<i>KEAP1</i> rs1048290				<b>0.044</b>
<i>GSTM1</i>				
CC	1/1	0	0	
CC	1/0	4 (15.4)	0	
CG/GG	1/1	0	1 (3.8)	
CG/GG	1/0	18 (69.2)	3 (11.5)	
<i>NRF2</i> rs35652124				<b>0.043</b>
<i>GSTM1</i>				
AA	1/1	0	0	
AA	1/0	9 (34.6)	2 (7.7)	
AG/GG	1/1	0	1 (3.8)	
AG/GG	1/0	13 (50)	1 (3.8)	



The 7th International Electronic Conference on Medicinal Chemistry

01-30 NOVEMBER 2021 | ONLINE



# Results and discussion



## Polymorphisms in estrogen metabolic pathway and age at diagnosis

n=157

<i>GSTM1</i>	<i>CYP1B1</i> Val432Leu	Age, n (%)		p-value
		<50	≥50	
Present	Leu/Leu (WT)	8 (5.1)	10 (6.4)	1
Present	Leu/Val + Val/Val	9 (5.7)	38 (24.2)	0.038*
Null	Leu/Leu (WT)	2 (1.3)	19 (12.1)	0.013*
Null	Leu/Val + Val/Val	12 (7.6)	59 (37.6)	0.012*

<i>GSTT1</i>	<i>CYP1B1</i> Val432Leu	Age, n (%)		p-value
		<50	≥50	
Present	Leu/Leu (WT)	10 (6.4)	21 (13.4)	1
Present	Leu/Val + Val/Val	17 (10.8)	62 (39.5)	0.239
Null	Leu/Leu (WT)	0	8 (5.1)	0.062
Null	Leu/Val + Val/Val	4 (2.5)	35 (22.3)	0.022*

<i>GSTT1</i>	<i>MTHFR</i> C677T	Age, n (%)		p-value
		<50	≥50	
Present	CC (WT)	10 (6.4)	30 (19.1)	1
Present	CT+TT	17 (10.8)	53 (33.8)	0.933
Null	CC (WT)	2 (1.3)	13 (8.3)	0.351
Null	CT+TT	2 (1.3)	30 (19.1)	0.034*

*GSTM1* null  
+  
*CYP1B1*Val

*GSTT1* null  
+  
*CYP1B1*Val

*GSTT1* null  
+  
*MTHFR* T allele

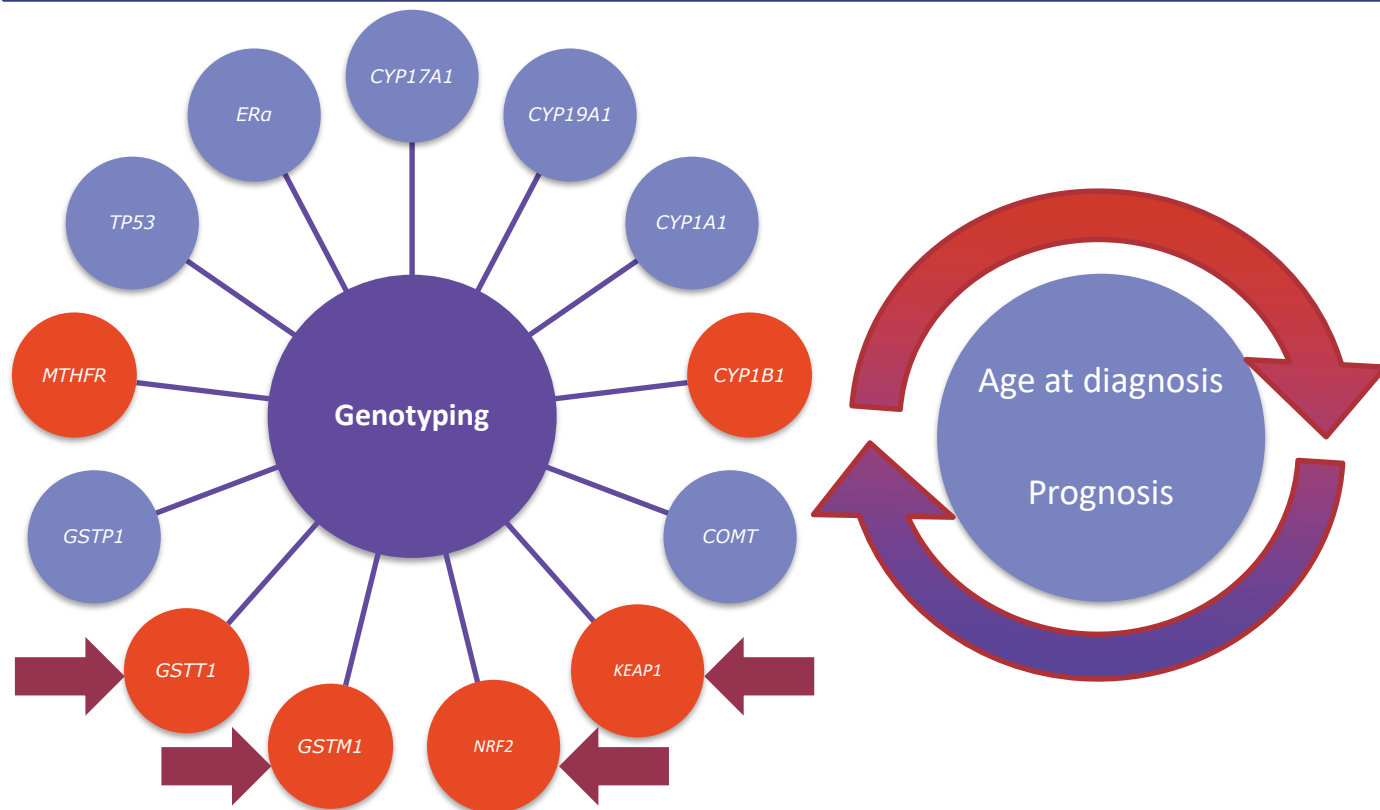
Breast cancer  
diagnosis at  
≥ 50 years old



# Results and discussion



Polymorphisms in estrogen synthetic and metabolic pathway: age at diagnosis and prognosis



Significant results allows to emphasize

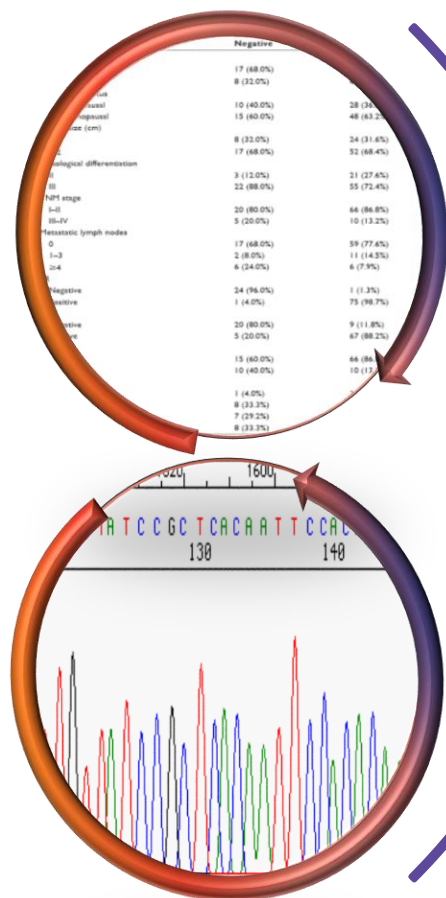
That polymorphisms in *NRF2/KEAP1* complex and the null polymorphisms of *GSTM1/GSTT1* should be consider prognostic markers for prognosis of the disease and of cancer therapy.



The 7th International Electronic Conference on Medicinal Chemistry  
01-30 NOVEMBER 2021 | ONLINE



# Conclusions



Cumulative presence of polymorphisms in *NRF2/KEAP1* and *GSTM1\*1/0* genotype might be associated with worse prognosis in breast cancer patients and with resistance to therapy.

Cumulative presence of polymorphisms in genes related to estrogen metabolism and lifetime exposure to estrogens might lead to breast cancer development at later ages.

Polymorphisms in *NRF2* and *KEAP1*, also as the null polymorphisms of *GSTM1* and *GSTT1* should be further investigated as possible targets for breast cancer therapy.



# Acknowledgments

- All the participants.
- All my supervisors and team members.
- Health Sciences Research Centre – UBI and Cova da Beira Hospital Centre.
- Ipatimup - Institute of Molecular Pathology and Immunology of the University of Porto and i3S – Institute for Research & Innovation in Health.
- “Validation of risk assessment model for breast cancer based on genetic polymorphisms of low penetrance” (ref. PTDC/DTPPIC/4743/2014).
- FCT fellowship - (ref. SFRH/BD146395/2019).



The 7th International Electronic Conference on Medicinal Chemistry

01-30 NOVEMBER 2021 | ONLINE

