CYP2D6 genotype and the association between tamoxifen treatment and risk of contralateral breast cancer in the WECARE Study

UBC Controls

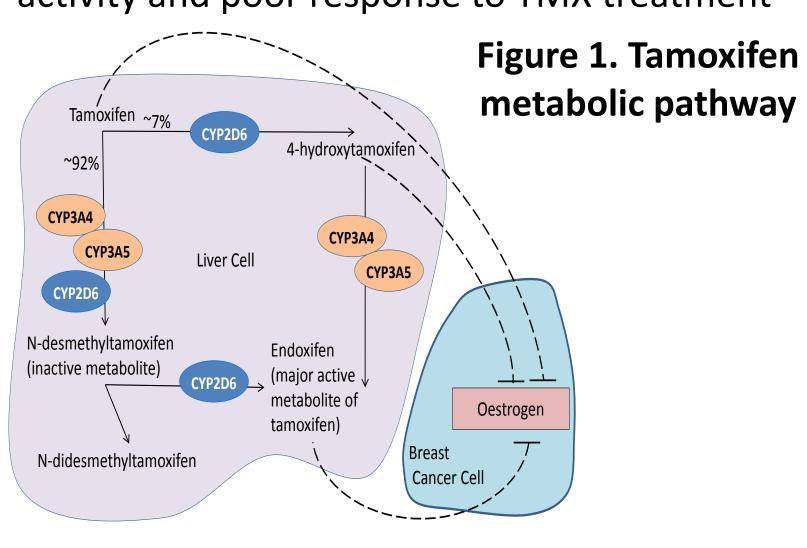
Siok Leong^{1*}, Anne Reiner¹, Xiaolin Liang¹, Elizabeth Comen², Jonine Bernstein¹, Jennifer Brooks³, Irene Orlow¹ and the WECARE Study Collaborative Group

¹Department of Epidemiology & Biostatistics, Memorial Sloan Kettering Cancer Center, New York NY ²Department of Medicine, Memorial Sloan Kettering Cancer Center, New York NY ³Dalla Lana School of Public Health, University of Toronto Canada



BACKGROUND

The most common malignancy in women who survive a first unilateral breast cancer (UBC) is the development of a second primary in the contralateral breast (CBC). UBC patients with estrogen receptor positive tumors are often treated with tamoxifen (TMX), which is metabolized into more active compounds by the Cytochrome P450 2D6 (CYP2D6) (Figure 1). Women who carry certain CYPD26 gene variants have been shown to have reduced enzyme activity and poor response to TMX treatment



Hypothesis: Women with poor metabolizing CYP2D6 phenotypes have a worse response to TMX and a subsequent increased risk of developing CBC compared to women not treated with TMX who are effective metabolizers

<u>Goal</u>: (I) To develop a specific assay for genotyping the *CYP2D6* full gene deletion and polymorphisms that were not previously captured by GWAS; (II) to investigate the risk of developing CBC according to TMX treatment and *CYP2D6* genotypes

Table 1. Patient characteristics

CBC Cases

Age at first dx 46 yr Period at risk* 6.3 yr Menopausal status^ Premenopausal 1,	an (range) Median (range) 46 yr (23-5 r (24-54) 5.5 yrs (1-19 s (1-19.8) N (%) N (%) 1,676 (76	4)
Period at risk* 6.3 yr Menopausal status^ Premenopausal 1,	rs (1-19.8) 5.5 yrs (1-19 N (%) N (%)	•
Menopausal status^ Premenopausal 1,	N (%)	9.8)
Premenopausal 1,		
•	1,676 (76	
Postmenopausal 3	- · · · · · · · · · · · · · · · · · · ·)
	389 (26) 522 (24)	
Unknown	8 (1) 14 (1)	
First-degree family hist	ory of breast cancer	
No 1,	.004 (66) 1,706 (77)
Yes 4	197 (33) 466 (21)	
Unknown	20 (1) 40 (2)	
Stage of first diagnosis		
Local 1,	.061 (70) 1,442 (65)
Regional 4	148 (29)	
Unknown	12 (1) 11 (1)	
Chemotherapy		
No 6	923 (42)	
Yes 8	322 (54) 1,289 (58)
Radiation treatment		
No 6	541 (42) 522 (24)	
Yes 8	380 (58) 1,689 (76)
Unknown	- 1 (0)	
Hormone treatment		
	964 (63) 1,270 (57)
	557 (37) 940 (42)	
Unknown	- 2 (0)	
ER status at first diagno		
	797 (52) 1,254 (57)
	467 (31) 561 (25) 267 (47) 207 (48)	
Other/unknown 2 ER estrogen recentor: *Perio	257 (17) 397 (18)	

ER, estrogen receptor; *Period refers to CBC diagnosis; ^menses or pregnancy within 2yrs of UBC dx

Table 2. Characteristics of studied CYP2D6 polymorphisms

RefSeq	Polymorphism	Effect	Allele nomenclature^	Metabolizing Phenotype
Reference	wt	wt	*1A	Extensive (EM)
rs16947; rs1135840	2850C>T, 4180G>C	R296C; S486T	*2D	Extensive (EM)
rs35742686	2549delA	FS	*3A	Poor (PM)
rs3892097	1846G>A	Splicing	contributes to *4A, 4B-H, and 4J-P with other SNPs	Poor (PM)
	Full deletion	X	*5	Poor (PM)
rs5030655	1707delT	FS	*6A-B-C-D	Poor (PM)
rs5030656	2615_2617del AAG	K281del	*9	Intermediate (IM)
rs1065852; rs1135840	100C>T; 4180G>C	P34S; S486T	contributes to*10A-B-D with other SNPs	Intermediate (IM)
rs28371725	2988G>A	Splicing	contributes to *41 and *91 with other SNPs	Intermediate (IM)

RefSeq, reference sequence; FS, frameshift; x, no protein/abolished function; ^ Nomenclature as per database http://www.cypalleles.ki.se accessed 02/09/17

RESULTS

Figure 4. Insertion/deletion

deleted (3.5Kb, yellow arrows) fragments

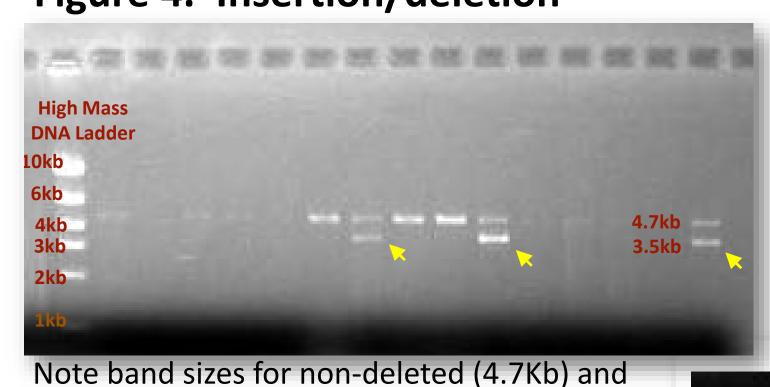
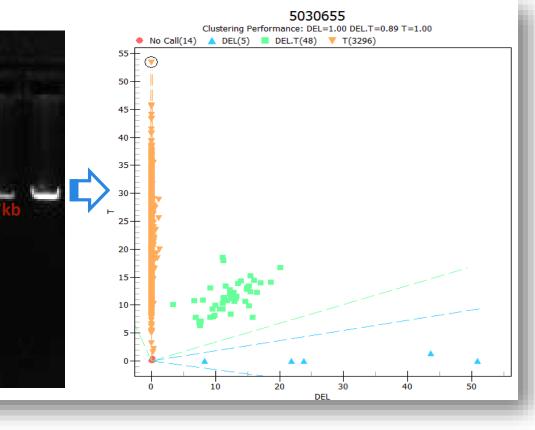


Figure 5. Two-stage genotyping

A specific 6.7Kb amplicon was amplified (Left panel) and used as template in the MassArray genotyping (Right panel)



METHODS

- 1121 CBC 'cases' & 2212 UBC 'controls' from the Women's Environmental Cancer and Radiation Epidemiology (WECARE) Study
- Detailed epidemiologic and clinical data (<u>Table 1</u>)
- CYP2D6, located in a complex locus (<u>Figure 2</u>), was tested for full deletion using 3 primers in a long-range PCR (Figures 3 & 4)
- Smaller variants (<u>Table 2</u>) were typed with a modified MassArray assay (Agena Bioscience), in a two-stage assay (Figure 5)
- Conditional logistic regression was used to estimate risk ratios (RR) and 95% confidence intervals (CI) for the association between TMX and risk of CBC, stratified by CYP2D6 genotype

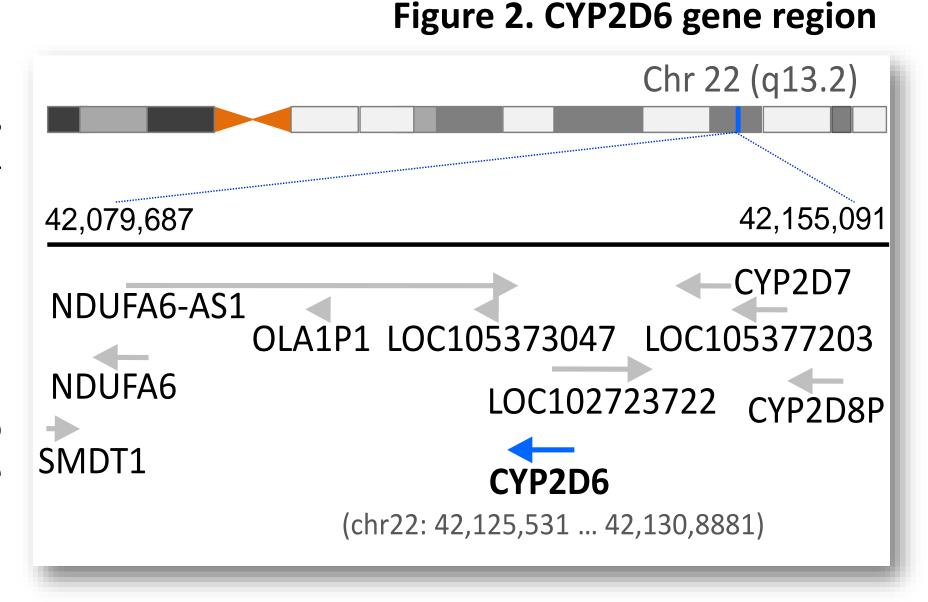
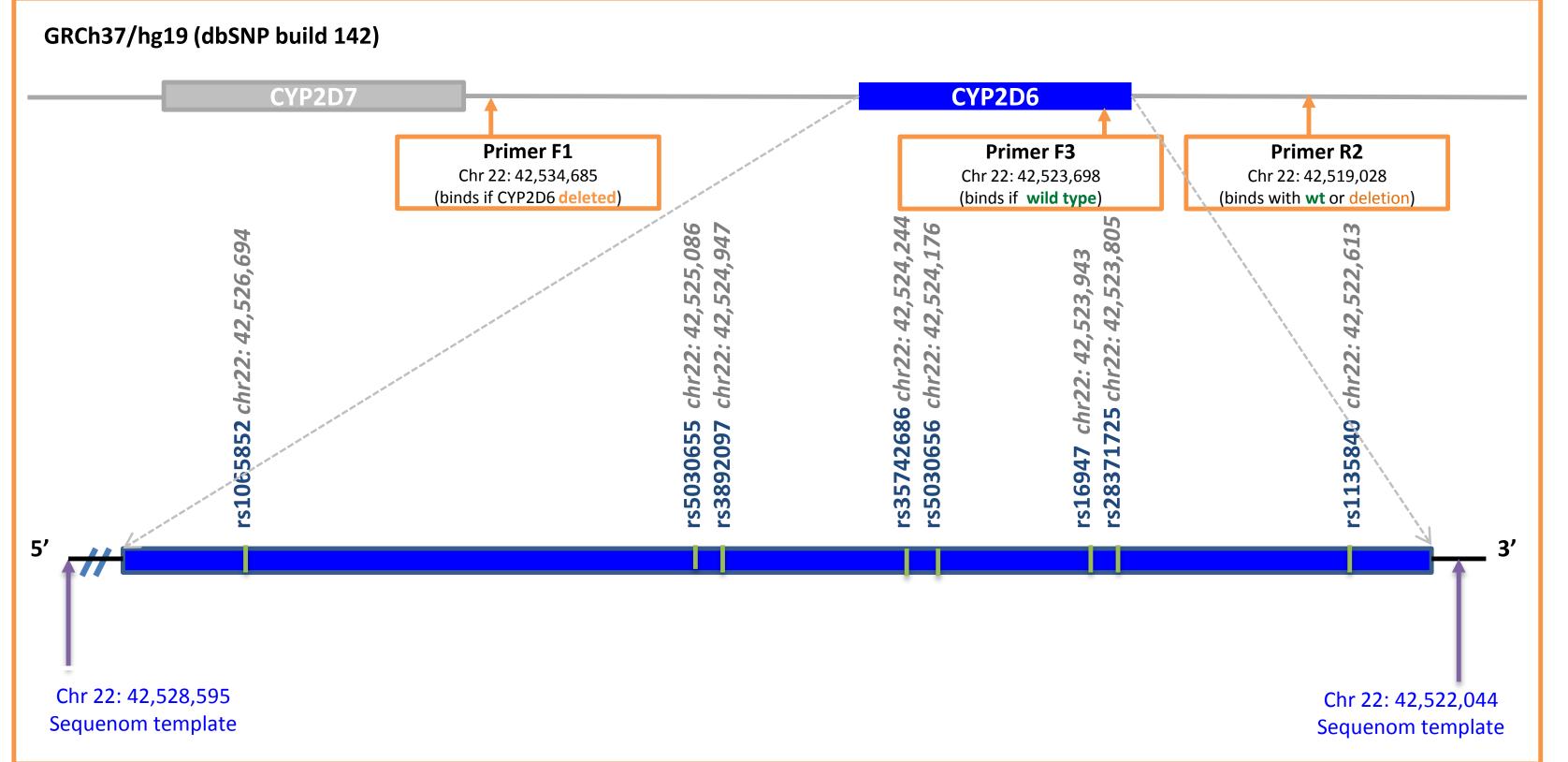


Figure 3. Summary of Assay Design to test deletion and small polymorphisms



Deletion analysis accomplished with 3 primers that amplify a 4.7Kb (wt) or a 3.5Kb (deletion) fragment

Table 3. Association between tamoxifen and risk of CBC

RefSeq	Genotype & Metabolizing —		TMX(-) N (%)		TMX (+) N (%)		RR* (95% CI)*	p-het*
	Phenoty	•	Cases	Controls	Cases	Controls	KK (55% CI)	p-net
	CC		462(44)	655(47)	227(50)	353(45)	0.74 (0.58-0.95)	
rs16947	СТ		441(42)	567(40)	163(35)	317(41)	0.56 (0.43-0.73)	
	TT	EM	140(13)	180(13)	67(15)	106(14)	0.67 (0.44-1.03)	0.28
	GG		357(34)	477(34)	172(37)	265(34)	0.72 (0.55-0.94)	
rs1135840	GC		494(47)	650(46)	187(40)	365(47)	0.55 (0.43-0.70)	
	CC	EM	195(19)	278(20)	105(23)	153(20)	0.82 (0.58-1.17)	0.11
	CC		639(61)	844(60)	278(61)	480(62)	0.65 (0.53-0.81)	
rs1065852	CT		343(33)	460(33)	139(30)	260(33)	0.56 (0.42-0.74)	
	TT	IM	61(6)	104(7)	39(9)	39(5)	1.54 (0.84-2.85)	0.01
rs3892097	GG		683(66)	893(63)	310(67)	525(67)	0.66 (0.54-0.81)	
	GA		316(30)	451(32)	135(29)	234(30)	0.61 (0.46-0.83)	
	AA	PM	42(4)	67(5)	17(4)	21(3)	1.30 (0.58-2.92)	0.24
rs5030655	Т	1	1034(99)	1394(99)	459(99)	769(98)	0.67 (0.56-0.79)	
	deleted	IM	15(1)	20(1)	3(1)	15(2)	0.17 (0.04-0.77)	0.06
rs5030656	AAG		988(94)	1344(95)	444(95)	750(96)	0.67 (0.56-0.79)	
133030030	deleted	IM	61(6)	73(5)	21(5)	35(4)	0.58 (0.28-1.18)	0.70
	G		868(83)	1211(85)	393(85)	651(83)	0.69 (0.57-0.82)	
rs28371725	GA		161(15)	188(13)	61(13)	115(15)	0.56 (0.37-0.85)	
	AA	IM	20(2)	18(1)	10(2)	19(2)	0.33 (0.11-0.97)	0.29
rs35742686	Α		1020(97)	1365(96)	454(98)	763(97)	0.66 (0.55-0.78)	
	deleted		28(3)	50(4)	9(2)	20(3)	0.86 (0.31-2.39)	0.61
full gene	present		903(94)	1221(93)	391(93)	661(94)	0.63 (0.52-0.77)	
	deleted	PM	55(6)	85(7)	31(7)	43(6)	1.43 (0.73-2.80) r age at first prima	0.02

* RR, relative risk among treated compared to untreated group. Adjusted for age at first primary, age at menopause two years prior to first primary, histology & stage of first primary, family hx of breast cancer, chemotherapy at first primary, radiation at first primary, number of full term pregnancies at first primary, and age at menarche. Deleted includes heterozygotes and homozygotes

SUMMARY

- Patients carriers of 2 CYP2D6 copies treated with TMX were at lower risk for CBC, while those carrying 0-1 copies were at higher risk for CBC compared to untreated individuals(pheterogeneity=0.02)
- Women carriers of rs1065852-C/C (reference) treated with TMX were at lower CBC risk, while those with 2 variant alleles (T/T) were at increased risk for CBC (p-heterogeneity=0.01)

CONCLUSION

- CYP2D6 polymorphisms modify response to TMX in relation to risk of CBC
- Further analyses in the context of other CYP2D6 polymorphisms (previously tested by GWAS) are warranted in order to categorize women into distinct TMX-response groups
- Knowledge on genetic modifiers such as CYP2D6 may contribute towards personalized breast cancer treatment