

## Evaluating the role of CHEK2 p.(Asp438Tyr) allele in inherited breast cancer predisposition

### *Familial Cancer*

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**Supplementary Table 1.** *CHEK2* primers used in the study

<b>Id</b>	<b>Expected product size (bp)</b>	<b>Start Position (Human GRCh37/hg19)</b>	<b>Sequence (5' - 3')</b>
<i>CHEK2</i> p.(Asp438Tyr) Forward	162	22:29091122	TGAGACTTCTGCCAGACTTC
<i>CHEK2</i> p.(Asp438Tyr) Reverse		22:29091264	CCACTGAGAATGCCACTTGA
<i>CHEK2</i> c.1100delC Forward	132	22:29091770	AGAACTTCAGGCAGCAAGTA
<i>CHEK2</i> c.1100delC Reverse		22:29091881	TGGCAAGTTAACATTATTCC

**Supplementary Table 2.** Tumor parameters of *CHEK2* p.(Asp438Tyr) carriers compared with the tumors of non-carriers

Category	Mut	%	WT	%	p <sup>a</sup>	OR	95 % CI
<b>T class</b>							
1	5	45.5	1143	64.0		0.47	0.14-1.54
2	5	45.5	566	31.7	(1 vs. 2. 3. 4)		
3	0	0.0	56	3.1			
4	1	9.1	20	1.1			
<b>M class</b>							
Neg	12	100	2002	97.8	1	-	-
Pos	0	0	45	2.2			
<b>Node status</b>							
Neg	6	50.0	1312	64.4		0.55	0.18-1.72
Pos	6	50.0	725	35.6			
<b>ER status</b>							
Neg	2	16.7	276	14.2		1.21	0.26-5.54
Pos	10	83.3	1666	85.8			
<b>PR status</b>							
Neg	2	16.7	439	22.8	1	0.68	0.15-3.10
Pos	10	83.3	1485	77.2			
<b>HER2 status</b>							
Neg	12	100	1588	85.4		-	-
Pos	0	0	271	14.6			
<b>Grade</b>							
1 (well diff.)	5	41.7	371	18.8		1.58	0.43-5.87
2 (mod. diff.)	4	33.3	921	46.7	(1 and 2 vs. 3)		
3 (poorly diff.)	3	25.0	682	34.5			
<b>Morphology</b>							
Ductal	8	66.7	1588	77.9		2.82	0.84-9.42
Lobular	4	33.3	307	15.1	(Lobular vs. all other)		
Medullary	0	0	10	0.5			
Other	0	0	133	6.5			
<b>Type</b>							
LumA	10	83.3	1414	76.4		2.04	0.44-9.40
LumB	0	0	193	10.4	(Triple negative vs. all other)		
HER2	0	0	78	4.2			
Triple negative	2	16.7	165	8.9			
<b>KI-67 index</b>							
0 (negative)	1	8.3	146	7.8		5.26	1.15-24.06
1 (weak)	9	75.0	765	40.9	(0 and 1 vs. 2 and 3)		
2 (moderate)	1	8.3	502	26.9			
3 (strong)	1	8.3	456	24.4			

*CI* confidence interval, *ER* estrogen receptor, *LumA* luminalA, *LumB* luminalB, *M* primary metastasis, *Mut* variant carrier, *Neg* negative, *OR* odds ratio, *Pos* positive, *PR* progesterone receptor, *T*tumor size, *WT* wild-type.

<sup>a</sup> Fisher's exact test.