Genetic Screening

가 가 가 가 가 가 가 . 가 BRCA1(breast cancer susceptibility gene 1) BRCA2 . BRCA1 BRCA2 17q21 13q12.3 100,000bp 70,000bp 가 . BRCA 가 (integrity) DNA (penetrance) . BRCA1/BRCA2 가 55 - 75% TP53, PTEN, STKIII/LKB1, ATM, 가 MLH1/MSH2 (Table 1). BRCA1/BRCA2 1. 5 - 10%

Table 1. Genetic Predisposition to Breast Cancer

Syndrome	Gene	Inheritance	Cancers
Breast/ovarian cancer syndrome	BRCA1	AD	Breast, ovary
	BRCA2	AD	Breast, ovary, prostate, pancreas
Li-Fraumeni syndrome	TP53	AD	Breast, brain, sarcoma, leukemia, adrenocortical carcinoma
Cowden disease	PTEN	AD	Breast, ovary, thyroid, colon
Peutz-Jegher syndrome	STKIII/LKB1	AD	GIT, breast
Ataxia-telangiectasia	ATM	AD	Breast
Site-specific breast cancer	CHEK2	AD	Breast
Muir-Torre syndrome	MLH1/MSH2	AD	Colorectal, breast

	(2004) 40 60
2. BRCA1/BRCA2	15% BRCA1/BRCA2
	(founder routetien) PDCA 7
71	(founder mutation) BRCA 가
가	가 가
BRCA1/BRCA2	BRCA
. BRCA 가 ,	. (2004)
, 가 ,	, 2.8%
가	(BRCA1 1.8%, BRCA2 1.0%) BRCA
. Ashkenazi 2%	, 12.7%
BRCA1/BRCA2 가,	(BRCA1 8.7%, BRCA2 4%)가
0.1% BRCA1 가 .	· 가 .
	, 가
	first - degree relatives 가 ,
BRCA1 0~5.7%,	-
BRCA2가 0.2~8.5% , 40	
JNOA22 0.2~0.370 , 40	BRCA 가
DDCA4 07 0 60/ DDCA27 4.2 6 60/	BNCA /
BRCA1 0.7~8.6%, BRCA2가 1.3~6.6%	DDCA4
(Table 2).	BRCA1 가
BRCA	84%, 44%
가 55% ,	(Table 3) 가
가 75% .	가
(2002)	, BRCA2 BRCA1
가 21 가 9가	•
(43%) BRCA1/BRCA2	BRCA
5 Breast Cancer Information Core (BIC)	,

Table 2. BRCA Mutations Among Cases of Female Breast Cancer Unselected for Family History (Modified Data from Table 1 in Hum Mutat 20: 413-424, 2002)

Population	Groups	Estimated prevalence			
		BRCA1 (%)	BRCA2 (%)	Total (%)	
Europe	Unselected	0.4 - 5.7	0.2 - 8.5	1.8 - 5.9	
	Selected*	0.7 - 6.8	1.3 - 2.4	2.0 - 9.0	
North America	Unselected	0.0 - 2.6	3.1	3.1	
	Selected	5.9 - 7.5	3.4 - 6.6	9.4 - 13.1	
Australia	Unselected	-	-	-	
	Selected	2.3 - 3.8	2.3	4.6	
Asia	Unselected	0.8 - 4.4	2.3 - 4.1	5.1 - 6.7	
	Selected	3.6 - 8.6	2.4	6.0	
Korea	Unselected				
	()	1.4	1.2		
	()	1.8	1.0	2.8	
	Unselected				
	()	8.7	4.0	12.7	
	()	10	8.7	15	

^{*}Selected groups were collected by age, family history, and bilaterality etc.

가 가 3. 가 National Cancer Institute(NCI) 가 BRCA1/BRCA2) Johns Hopkins 가 가 가 , 가 가 가 4. Ashkenazi 가 BRCA1 BRCA2 가 가 가 , 35 BRCA1/BRCA2 가 가 가 가 가 가 100% , F-CSGE(flu-BRCA1/BRCA2 penetrance 가 orescent conformation - sensitive gel electrophoresis) BRCA1 가 DHPLC(denaturing high performance liquid chro-84%, matography) 가 44% 가 BRCA1/BRCA2 18 가

Table 3. Cumulative Risk by Age of Breast Cancer in Women from Families with BRCA1 and BRCA2 Mutations (Easton et al, 1995, 1997)

100/100

297,000

(\$2700)

가

Age (yrs)	Cumulative risk (%)		
_	BRCA1	BRCA2	
30	3.2	4.6	
40	19.1	12.0	
50	50.8	46.0	
60	54.2	61.0	
70	85.0	86.0	

Personal Characteristics

Breast cancer diagnosed at an early age

가

Bilateral breast cancer

, 25

A history of both breast and ovarian cancer

The presence of breast cancer in 1 or more male family members

CA - 125

가

가

BRCA1

Family Characteristics1

Multiple cases of brest cancer in the family
Both breast and ovarian cancer in the family
One or more family members with 2 primary cancers
Ashkenazi Jewish background

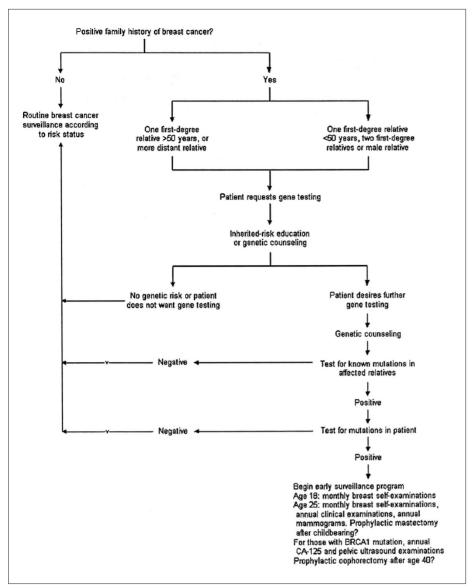


Fig. 1. Suggested algorithm for genetic screening in women with a family history of breast cancer (based on recommendations of NIH and National Human Genome Research Institute)

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- 1. : . 10
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