

Table 11. Risk's level according to genotypes combination

	Cases	Controls			
Genotype	n	n	OR	95% CI	P
Reference genotype	254	633	1.0		
Intermediate risk	643	776	2.32	1.79-2.54	< 0.001
High risk	1291	767	4.22	3.55-5.02	< 0.001

^aAdjusted for age, age at menarche, age at FFTP, number of full term pregnancies, first-degree family history of breast cancer, history of benign breast disease, use of oral contraceptives, WHR, alcohol consumption, smoking habits, level of education and postmenopausal use of estrogen. ORs are calculated relative to subjects with the reference genotype.

Reference genotype GSTP1ValVal-PON1/192RR-PON1/55LL-GLO1AA.

Intermediate risk: GSTP1IleVal-PON1/192QR-PON1/55LM-GLO1AE.

High risk: GSTP1IleIle-PON1/192QQ-PON1/55MM-GLO1EE.