



Predictive potential role of glutathione S-transferases polymorphisms in response to chemotherapy and breast cancer prognosis

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ABSTRACT. The aim of this study was to evaluate the role of *GSTM1* null/present, *GSTT1* null/present, and *GSTP1* polymorphisms in the clinical response to chemotherapy and treatment outcome of breast cancer. The *GSTM1*, *GSTT1*, and *GSTP1* Ile105Val polymorphism genotypes were analyzed using polymerase chain reaction coupled with restriction fragment length polymorphism. Conditional logistic regression analysis revealed that breast cancer patients carrying the GG genotype of *GSTP1* Ile105Val showed a significantly better response to chemotherapy compared to those expressing the AA genotype [odds ratio = 2.66, 95% confidence interval (CI) = 1.24-5.91, P = 0.007]. The Cox proportional hazards model indicated

that the GG genotype of *GSTP1* Ile105Val in breast cancer patients was correlated with a lower risk of death from all causes than those with AA genotype. The adjusted hazard ratio (95%CI) for the GG genotype of *GSTP1* Ile105Val was 0.44 (0.18-0.99; P = 0.03). In conclusion, the results of our study indicated that the GG genotype of *GSTP1* Ile105Val was significantly associated with better response to chemotherapy and longer overall survival, compared to the wide-type genotype.

Key words: Glutathione S-transferases; Polymorphism; Chemotherapy; Breast cancer