

Genetic polymorphisms of CYP1A1, GSTM1, GSTT1, and GSTP1 genes in B-cellular chronic lymphocytic leukemia (B-CLL)

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Abstract

Aim: The aim of this study was to examine the relationship between genetic polymorphisms in GSTM1, GSTT1, GSTP1, and CYP1A1 genes and their susceptibility to B-CLL.

Methods: The DNA for our research was isolated from the leukocytes of venous peripheral blood taken from 146 patients with B-cell CLL and 221 healthy individuals, using a standard phenol-chloroform extraction method. GSTM1 and GSTT1 genotypes were determined by a multiplex PCR technique that detects homozygous deletions of the mentioned genes and the β -globin housekeeping gene as an internal control. CYP1A1 mutation A4889G (Ile462Val) and GSTP1 polymorphisms A1578G (Ile105Val) and T2293C (Ala114Val) were identified by the PCR-RFLP approach.

Results: For the first time it was shown that the mutant genotype (GSTP1-105IleVal+GSTP1-105ValVal) occurs significantly more often in B-CLL patients than that of healthy individuals (65.1% ps. 53.4%, $\chi^2 = 4.92$, $p < 0.05$; OR=1.63, 95% CI=1.06÷2.50), whereas the "wild" genotype GSTP1-105IleIle was significantly underrepresented in patients with CLL compared with those of healthy individuals (34.9% ps. 46.6%, $\chi^2 = 4.92$, $p < 0.05$; OR=0.62, 95% CI=0.40÷0.95).

At the same time the difference in distribution of homozygous deletion of the GSTM1 and GSTT1 genes and also of the CYP1A1-Ile462Val and GSTP1-Ala114Val genotypes between B-CLL patients and healthy individuals was statistically not significant ($p > 0.05$).

Conclusions: The results of the current study suggested that the GSTP1 polymorphism may influence the risk of B-CLL developing.

Keywords: B-CLL, genetic susceptibility, CYP1A1 polymorphism, GSTM1 polymorphism, GSTT1 polymorphism, GSTP1 polymorphism