

***MTHFR* C677T Polymorphism and Recurrent Early Pregnancy Loss Risk in North Indian Population**

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Abstract

Recurrent early pregnancy loss (REPL) is a multifactorial disorder as both genetic and environmental factors contribute to the development of disease. Folate metabolism is an important mechanism to ensure proper fetal growth. Hyperhomocysteinemia leads to a number of disorders and REPL is one of them. In a case-control study DNA from 106 cases with the history of 3 or more REPL and 140 healthy fertile controls with successful pregnancy outcomes were genotyped for C677T single-nucleotide polymorphism (SNP) of the *MTHFR* (methylenetetrahydrofolate reductase) gene through polymerase chain reaction-restriction fragment length polymorphism (PCR-RFLP), which was further confirmed by sequencing. Allele frequencies of REPL cases were compared with healthy controls and a statistically significant association was found between REPL and the mutant T allele ($\chi^2 = 8.786$, odds ratio [OR] = 2.20, 95% confidence interval [CI] = 1.323-3.9658, $P = .003$). The genotype frequencies of SNP C677T also differ significantly between these 2 groups ($\chi^2 = 8.237$, $P = .016$). The OR for heterozygous CT in the REPL versus controls is 1.9591 (95% CI = 1.0285-3.7318, $P = .04$). The OR for TT homozygous is 6.3009 (95% CI = 1.2065, $P = .02$). Combined odds ratio of CT and TT against the control has been calculated as 2.2194 (95% CI = 1.2029-4.0952, $P = .02$) which is also significant. Thus the present study clearly indicates that homozygosity and heterozygosity for the *MTHFR* C677T polymorphism confer a 6.3009- and 1.9591-fold increased risk of idiopathic REPL, respectively.