

CYP2C19 polymorphisms impacts citalopram metabolism

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(HealthDay)—For patients with major depressive disorder, certain *CYP2C19* polymorphisms contribute to citalopram (CIT) metabolism, according to research published in the December issue of the *Journal of Clinical Pharmacy and Therapeutics*.

Z. Uckun, Ph.D., from Mersin University in Turkey, and colleagues determined *CYP2C19* genetic polymorphisms and their impact on the metabolism of CIT. Genotypes were determined by polymerase chain reaction-restriction fragment length polymorphism method in 209 healthy individuals and 50 patients with <u>major depressive disorder</u>.

The researchers found that the *CYP2C19*1* and *CYP2C19*17* allele frequencies were 71.0 and 18.0 percent, respectively, for the healthy group and 81.1 and 18.9 percent, respectively, for the patient group (P >



0.05). Patients with the *CYP2C19*1/*1* genotype had significantly higher mean plasma concentration and the mean dose-corrected plasma levels of demethylcitalopram (DCIT) compared to patients with *CYP2C19*1/*2* or *CYP2C19*2/*2* genotypes (P CYP2C19*1/*2 and *CYP2C19*2/*2* genotypes, the mean metabolic ratio (MR, CIT/DCIT) was significantly higher (P CYP2C19*1/*1 and *CYP2C19*1/*17* genotypes (P > 0.05).

"Our data suggest that *CYP2C19*17* polymorphism does not have a significant effect on CIT metabolism," the authors write. "In contrast *CYP2C19*2* polymorphism has a prominent role and is likely to contribute to interindividual variability in CIT metabolism in vivo at therapeutic doses."

More information: Abstract

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