

CYP2C19 Pharmacogenomics–based *Helicobacter pylori* eradication therapy by proton pump inhibitors

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Two cytochromes P450 (CYPs), CYP2C19 and CYP3A4, are involved in the metabolism of proton pump inhibitors (PPIs) such as omeprazole, lansoprazole, rabeprazole and pantoprazole. Among these two CYPs, CYP2C19 is a genetically determined polymorphic enzyme, showing an interethnic difference in the incidence of poor metabolizer (PM) phenotypes/genotypes: the frequency of PMs is about 13 to 24% in four Asian (Chinese, Korean, Indonesian and Japanese) populations, being much greater than that (2 to 4%) in American- and European-Caucasian populations. Thus, if the CYP2C19 pharmacogenetic (or pharmacogenomic) entity would have any clinical implication, this entity should be of a more therapeutic concern among the Asian than among the Caucasian Patients.

For the hypothetical reasoning as mentioned above, I and my colleagues have accomplished several studies on PPIs in *H. pylori*-positive Japanese patients as follows: in a study on CYP2C19 genotype status and dual therapy with omeprazole and amoxicillin for cure rates of *H. pylori* and peptic ulcer healing, cure rates in patients with wt/wt, wt/m1 or wt/m2 and m1/m1 or m1/m2 genotype status were 28.6, 60.0 and 100%, respectively, and ulcer healing rates depended on success or failure of *H. pylori* as well as on the CYP2C19 genotypic status.

This trend was also confirmed in *H. pylori*- positive patients with gastritis who underwent dual therapy with rabeprazole and amoxicillin. In another study with a PPI (omeprazole or lansoprazole), amoxicillin and clarithromycin (a triple therapy) in 261 Japanese patients with *H. pylori*- positive peptic ulcer, we have observed that the majority of patients without initial eradication of *H. pylori* had the homozygous extensive metabolizer (hom EM) CYP2C19 genotype, but these hom EM patients were successfully re-treated with a high-dose lansoprazole and an antibiotic to which *H. pylori* was sensitive, such as amoxicillin, even when the patients were infected with clarithromycin-resistant strains of *H. pylori*. In addition, we have recently observed that cure rates of gastroesophageal reflux disease (GERD) depended

not only on the *CYP2C19* genotype status but also on the pre-treatment grades of GERD during the treatment with lansoprazole.

Based upon our findings as presented above. I wish to propose a pharmacogenetic- (or pharmacogenomic-) based tailor-made therapeutic strategy for peptic ulcer or GERD in our future clinical practice. Lastly, because the lowest therapeutic success for eradicating *H. pylori* infection was observed in patients with hom EM genotype of *CYP2C19* who more often showed a nocturnal acid breakthrough, a histamin 2 (H₂-) receptor antagonist like famotidine, instead of a PPI, should be re-evaluated for the hom EM patients with peptic ulcer or GERD.