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VacA and *cagA* genotypes of *H. pylori* and *IFN-* γ expression in chronic gastritis and gastric cancer patients

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Helicobacter pylori represents the major cause of chronic gastritis, duodenal and gastric ulcer and gastric cancer. Clinical outcome of *H. pylori* infection is determined by the virulence factors of bacteria, environment and immune response of the host. A differential expression of cytokines between *H. pylori* positive and *H. pylori* negative patients or between gastritis and gastric cancer patients was reported. The aim of present study was to compare the IFN- γ expression between *vacA* and *cagA* genotypes of *H. pylori* in chronic gastritis and gastric cancer patients. Ninety-six patients with chronic gastritis and twenty with gastric cancer were recruited. *H. pylori* infection, *vacA* and *cagA* genotyping were accomplished via PCR from total DNA of gastric biopsies, the IFN- γ expression was determined by immunohistochemistry. Of the patients overall, 66 (56.9%) were infected with H. pylori, among patients with chronic gastritis 54 (56.3%) and 12 (60%) in gastric cancer were *H. pylori* positive. The predominant *vacA* and *cagA* genotypes of *H. pylori* were vacAs1m1 (87%) and cagA positive (77.8%) in chronic gastritis and 83.3% and 58.3% in gastric cancer. We observed a variation of expression of IFN- γ in patients with chronic gastritis and 83.2% and 58.3%) compared with *vacAs1m1* (82%) and s2m2 (72%) genotypes and in patients infects with *H. pylori* cagA positive (82%) respect to *H. pylori* could be important factors that increase the damage in gastric cancer.

Biography

Martínez-Carrillo Dinorah Nashely has completed her master's degree at the age of 29 years from Universidad Autónoma de Guerrero. She has published 4 papers in reputed journals. She is currently pursuing her Ph.D.

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