TITLE: TRAIL, DR4, AND DR5 GENE EXPRESSION: APOPTOSIS DYSREGULATION AND GASTRIC TUMORIGENESIS

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ABSTRACT:

Helicobacter pylori (H. pylori) is one of the main causes of gastric diseases including Gastric Cancer (GC), which is one of the leading causes of cancer deaths worldwide. In addition to *H. pylori* infection, deregulation of apoptosis process can be a risk factor for cancer. TNF-related apoptosis-inducing ligand (TRAIL) is responsible for promoting apoptosis by binding to its Death Receptors TRAIL-R1 (DR4) and TRAIL-R2 (DR5). TRAIL is selective to neoplastic cells; however, these cells can present resistance to this pathway, especially GC cells. We aimed to evaluate the expression level of TRAIL, DR4, and DR5 along with the presence of H. pylori in patients with dyspeptic symptoms and GC, to identify the influence of these pro-apoptotic genes in the gastric tumorigenesis. H. pylori was detected using PCR, and gene expression analysis was performed by realtime quantitative PCR. The patients were divided in groups according to histopathological analysis (Control, Gastritis, and Cancer groups). Of the 244 gastric biopsy samples analyzed, 103 tested positive for the bacterium, which was most prevalent in the Cancer group (73.1%). DR4 and DR5 genes were overexpressed in the Cancer group compared to the Control (p = 0.0007) and Gastritis (p < 0.0001) groups, whereas the TRAIL gene was less expressed when comparing the same groups (p <0.0001). Our results indicate that GC cells express high levels of DR4 and DR5 in attempt to stimulate apoptosis while expression of TRAIL is decreased in the same type of cancer, confirming the resistance of this cancer to apoptosis via TRAIL.

Keywords: Helicobacter pylori, gastric cancer, TRAIL, DR4, DR5

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