

Abstract

Esophageal cancer is one of the most invasive type of gastrointestinal cancer, the most common cause of cancer-related deaths in the world. Most esophageal cancers occur in the age group 70-50 years. The frequency of this disease in men more than women. *Astrocyte elevated gene -1*(*AEG-1*, also known as *MTDH*, and *Lyric*) has been demonstrated to play a potential role in several significant aspects of tumor progression. Expression or genomic amplification This gene can serve as a biological marker to identify subgroups of patients who require more advanced treatments targeting *AEG-1* and benefits are used. In the present study, the expression of *AEG-1* in formalin fixed paraffin-embedded (FFPE) Esophageal cancer tissue samples in the Iran were studied. Fifteen FFPE Esophageal cancer tissue samples from medical centers were analyzed for *AEG-1* gene expression level by real-time quantitative reverse transcriptase polymerase chain reaction. All PCR reactions were performed in triplicate for both target gene (*AEG-1*) and internal control (*B-actin*) with the $2^{-\Delta\Delta C_t}$ method. Differences in target gene expression levels in patients and controls were evaluated with t-test. $P \leq 0.05$ was considered to be significant. All analyses were performed using the SPSS 13 software (SPSS, Inc., Chicago, IL). The results showed *AEG-1* gene over expression. There was a statistically significant difference in the prevalence of *AEG-1* expression between patients and control ($p < 0.05$). Our results demonstrated *AEG-1* gene over expression in formalin fixed paraffin-embedded (FFPE) esophageal cancer tissue samples compared with controls.

Key words: Esophageal cancer, *AEG-1*, , Reverse transcriptase real-time polymerase chain reaction



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**Evaluation of *Astrocyte elevated gene -1* expression in
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