

Abstract

Background: Gastric adenocarcinoma (GAC) is a serious disease with dismal outcome. Discovering novel molecular targeted therapies is a recent point of research to improve prognosis. One of the newly discovered targets is the receptor tyrosine kinases (RTKs); it is a member of trans-membrane receptors which had important roles in proliferation and apoptosis. RTKs were found to have different expression patterns in several malignancies. HER2 neu which is a member of HER family is a proto-oncogene that is formed of four receptor tyrosine kinases. Erythropoietin-producing hepatocellular (Eph) molecules are major RTKs members and one of those molecules is EphA2 which has many different functions in cancer as tumor initiation, progression, angiogenesis and spread. We aimed to explore the expression patterns of both HER2 neu and EphA2 in GAC patients using immunohistochemistry, and to correlate their expressions with clinico-pathological factors and prognosis of our patients

Methods: 100 GAC patients were diagnosed as GAC. Then we analyzed the correlations between their expressions and disease outcome of GAC patients.

Results: HER2 neu and EphA2 positive expressions in GAC were positively correlated with tumor grade and stage ($p < 0.001$ and $p = 0.002$ respectively), inadequate response to therapy ($p < 0.001$ and $p = 0.002$ respectively), increase recurrence rate of GAC ($p = 0.002$), and with poor survival ($p < 0.001$).

Conclusion: GAC patients with high expressions of both HER2 neu and EphA2 had unfavorable prognosis.

E) No significant correlation was found between patient's age or sex, histopathological subtype, initial site, or size of the tumor with markers expression.

Treatment response and survival analysis (Table 3 and Figure 3)

therapy response

Patients with advanced disease (stage III) were assessed for response. Of the 29 patients, 21 patients (72.4%) had CR, 2 patients (6.9%) had PR, and 3 patients (10.3%) had SD and PD (Table 4).

Table 3: Correlation between immunohistochemical markers and disease outcome of our patients.

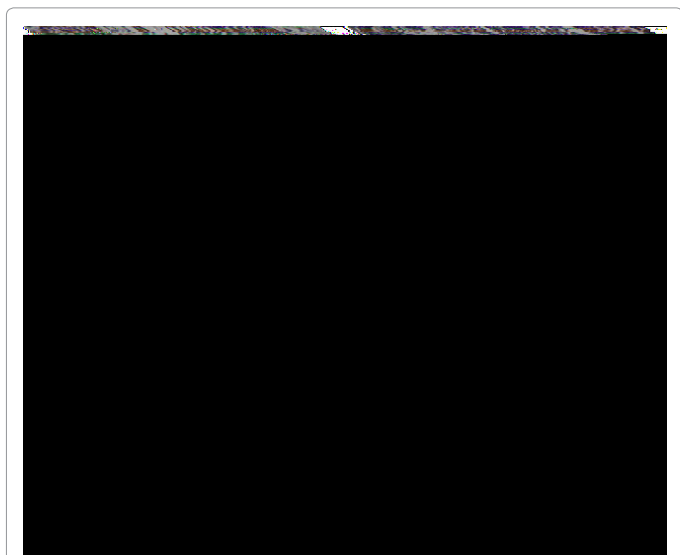


Figure 1: Immunohistochemical staining of HER2 neu in gastric adenocarcinoma (GAC): (A) High membranous expression of poorly differentiated GAC X400 (B) High membranous expression of moderately differentiated GAC X400. (C) Low membranous expression of moderately differentiated GAC X400. (D) Low membranous expression of well differentiated GAC 400. A, B, C and D the

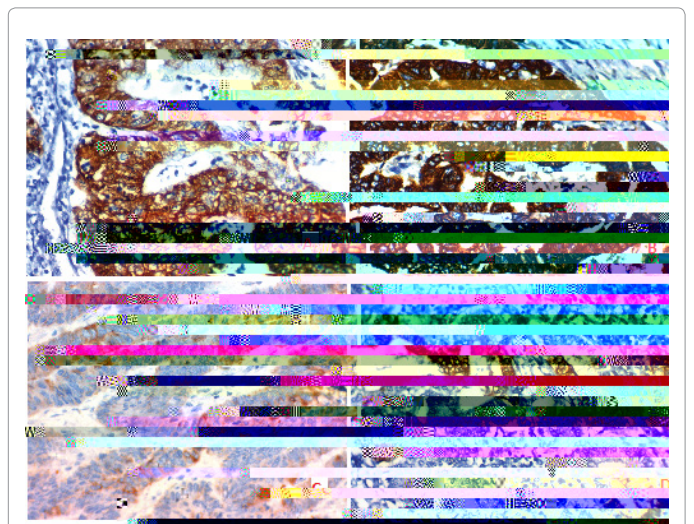


Figure 2: Immunohistochemical staining of EphA2 in gastric adenocarcinoma (GAC) : (A) High expression in the cytoplasm of poorly differentiated GAC X400; (B) High expression in the cytoplasm of moderately differentiated GAC X400; (C) Low expression in the cytoplasm of moderately differentiated GAC X400; (D) Low expression in the cytoplasm of well differentiated GAC X400. A, B, C and D the

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but no significant relation was found between expression pattern of HER2 neu and tumor recurrence.

Survival analysis: After a median follow up of 40 (Range: 6-58) months, Positive expressions of EphA2& HER2 neu were significantly associated with shortened disease free survival (DFS) and overall survival (OS) ($p < 0.001$ for both, Table 4, Figure 3).

Discussion

Many researchers had investigated the prognostic role of HER2 expression in cancers of many organs [11]. Previous studies have reported that the frequency of IHC detection of HER2 overexpression in GC varies from 10% to 22.1% [17], while positive expression of HER2 neu was detected in 50% of our cases and this high percentage of positive expression may be due to small patients number or due to inclusion of early as well as advanced stages in our study.

In our research we found that positive expression of HER2 neu was correlated with higher tumor grade, high incidence of L.N metastases and advanced stage of the cancer. Our results were close to Leni et al. [18] who reported that HER2 overexpression was more frequent in advanced GAC with high grade and advanced stage, and that was significantly associated with high disease recurrence and poor prognosis. Our results were consistent with Jia et al. who reported that HER2 over-expression was correlated with increased depth of invasion ($P=0.045$), lymph-nodes metastasis ($P=0.026$), and elevated clinical stage ($P=0.026$) but was not significantly associated with patient age, gender or cancer location [17]. Our results may be explained by that HER2 overexpression leads to increase cellular proliferation and inhibits apoptosis resulting in uncontrolled and excessive growth and spread of cancer.

Other different results were detected by Park et al. and Oh et al. [19,20] who stated that gastric tumor with HER2 neu amplification was only associated with old age and tumor size but it had no relation to prognosis. This discrepancy may be due the use of different immunohistochemical clones, the number of examined cases or the selection criteria that implied further study on a larger scale. Tessa and Raghuvver [21] assessed the expression HER-2 in cervical cancer and proved that it was positively associated with increasing the grade of cancer, presence of lymph node metastases and parametrial spread which was in agree with our results. Our results were also compatible with Park et al. [22] who studied Her2 amplification in colon cancer and reported that it was associated with higher rates of nodal metastasis and decreased patient survival. Hence, HER2 neu overexpression was found to be a prognostic factor for GAC and was negatively correlated with survival rates that were similar to results of Zhang et al. [23] and Park et al. [19] however, Jeung et al. [24] found no significant relation between HER2 neu expression and grade or stage of GAC; such difference that may be related to the nature of studied group and their number.

Also we found that positive expression of EphA2 was positively correlated with tumor grade, L.N metastases and tumor stage ($p=0.005$, $p=0.002$ and $p < 0.001$ respectively). The results were similar to Huang et al. [7] and may be explained by that EphA2 stimulates proliferation, migration and spread of GAC cells mainly by increasing the expression of the epithelial mesenchymal transition markers like snail, N-cadherin, b-catenin, stimulating the Wnt/b-catenin pathway and by inhibition of E-cadherin in GAC cells. EPHA2 is overexpressed in a wide range of cancers and is associated with poor prognosis [25]. Many recent studies investigated the RTKs such as EphA2 and reported them as targets for molecular therapy for GAC [26-29], and also proved that

EphA2 overexpression was positively correlated with factors that controlled angiogenesis and invasion in cancer cells because EphA2 receptor activation allowed vascular endothelial growth factor (VEGF)-dependent endothelial cell transport, sprouting, survival and expression of metalloproteinase, and these may be the causes of the poor clinical outcome of cancer patients with EphA2 overexpression, moreover the EphA2-EphrinA1 signaling axis regulates many steps that are essential for carcinogenesis and stimulation of downstream

