

Investigation of Soluble HER2 and Transforming Growth Factor Beta-1 Serum Levels in Gestational Trophoblastic Disease

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Abstract HER2/neu and TGF- β 1 are over-expressed in various types of malignancies. It appears that they play an important role in the biologic behavior of tumors and have prognostic value. Gestational trophoblastic diseases (GTDs) comprise of a heterogeneous group characterized by abnormally proliferating trophoblastic tissues, ranging from benign to malignant. The objective of this study was to measure and compare the serum levels of s-HER2 and TGF- β between patients with GTDs and pregnant and non-pregnant controls. Serum levels of s-HER2 and TGF- β 1 were determined by ELISA method in 95 GTD patients (55 complete moles, 32 persistent moles, and 8 choriocarcinoma), 30 normal pregnant controls, and 22 normal non-pregnant controls. Mean serum level of s-HER2 did not differ significantly between patients and controls. TGF- β 1 serum level was significantly higher in GTD patients (20.29 ± 10.68 pg/ml with 95% confidence interval (CI) of 18.10–22.48 pg/ml) compared with pregnant controls (10.26 ± 11.84 pg/ml with 95% CI of 5.75–14.76 pg/ml) and non-pregnant controls (7.27 ± 9.61 pg/ml with 95% CI

of 3.01–11.53 pg/ml) ($P < 0.001$). Our findings suggest that TGF- β 1 serum levels in GTD patients may represent a potential prognostic marker. Further investigations with larger sample size and more frequent sampling are required to elucidate this issue.

Keywords HER2 · TGF- β 1 ·
Gestational trophoblastic disease

Introduction

Gestational trophoblastic diseases (GTDs) consist of a spectrum of disorders characterized by an abnormal proliferation of trophoblastic tissue. GTDs include hydatidiform mole (complete and partial), invasive mole, choriocarcinoma, and two rare variants called placental site trophoblastic tumor (PSTT) and epithelioid trophoblastic tumor (ETT). It is considered today the most curable gynecologic cancer [1, 2].

HER2/neu oncogene (also known as c-erbB-2) is a member of the epidermal growth factor receptor (EGFR) family. It is over-expressed in various types of cancers including breast, ovarian, lung, gastric, and oral cancers; it is one of the biomarkers used as a prognostic and predictive factor in some solid malignancies [3]. Clinical and molecular studies suggest that increased soluble HER2 (s-HER2) expression may contribute to disease progression by mediating angiogenesis, metabolic adaptation, and other aspects of metastasis that define the cancer phenotype [4].

Increased expression of HER2/neu in breast cancer cells results in transactivation of EGFR family [5]. Several investigations have shown that over-expression of HER2

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is present in metastatic breast cancers and is associated with resistance to chemotherapy, resulting in decreased survival of patients [6, 7]. Over-expression of HER2/neu was also found to be a poor prognostic factor for survival from advanced-stage ovarian cancer and endometrial cancer [8].

Transforming growth factor- β (TGF- β) is a cytokine that almost every cell in the body produces and has receptors for it. It acts as a modulatory cytokine released by T regulatory cells during the course of cell-mediated immune response. TGF- β regulates cellular proliferation by inhibition, thus in normal cells it works as a tumor suppressor. Mutations in the TGF- β pathway in cancer cells render these cells resistant to the growth inhibition of TGF- β , causing uncontrolled proliferation of the cells [9, 10].

There are three isoforms of TGF- β , each encoded by a distinct gene. TGF- β 1 messenger RNA is expressed in endothelial, hematopoietic, and connective-tissue cells. The levels of TGF- β 1 in serum can be measured and used as diagnostic or prognostic marker for human diseases [9]. Increased serum level of TGF- β 1 has been detected in some solid malignancies such as gastric and breast cancer [11, 12]. High serum levels of TGF- β 1 protein are also found in patients with invasive prostate cancer [13] and colorectal cancer [14].

TGF- β signaling pathway plays an important role in the pathogenesis and progression of GTDs [15, 16]. Increased expression of TGF- β 1 mRNA has been found in tissues from molar pregnancy [17].

The aim of the present study was to determine the serum level of s-HER2 and TGF- β 1 in GTD patients and to compare the values with those of pregnant and non-pregnant healthy women. No studies investigating serum level of s-HER2 and TGF- β 1 in GTD patients have been published up to date.

Materials and Methods

Subjects A total of 95 patients with GTD (mean age of 27.56 years) referred from Obstetrics and Gynecology outpatient clinics or hospital wards of Shiraz University of Medical Sciences were studied, including 55 patients with complete hydatidiform mole, 32 patients with persistent hydatidiform mole, and 8 patients with choriocarcinoma. Diagnosis was made based on histopathologic examination of the tissue obtained from endometrial curettage. Six months after evacuation a follow-up sample was obtained from six patients with complete mole and five patients with persistent mole (selected from the total of 95 patients). Thirty healthy pregnant women regional volunteers with normal pregnancy and gestational ages of 10 to 22 weeks with single alive fetus and 22 healthy non-pregnant women in reproductive age were also recruited as controls in this

case-control study. Controls were those attending Obstetrics and Gynecology outpatient clinics for routine prenatal check-ups or for routine visits. No significant difference was found between case and control groups regarding age, gravidity, parity, and gestational age.

This study was approved by our university ethics committee. Informed consent was obtained from all participants.

s-HER2 and TGF- β 1 Serum Levels Determination Sera from patients (prior to molar evacuation) and controls were collected in sterile clean dry tubes, separated, and kept frozen at -70°C until used. s-HER2 and TGF- β 1 serum levels were determined using commercial ELISA kits (BenderMed System, Austria for sHER2 and Biosource, USA for TGF- β 1), according to manufacturer recommendations.

Statistical Analysis The data were analyzed using SPSS software (version 11.5.0; SPSS Inc., Chicago, IL, USA). T-test, Mann–Whitney test, and one way ANOVA were used to compare groups, when appropriate. Serum levels are expressed as mean $\pm$ standard deviation (SD). A *P*-value of less than 0.05 was considered significant.

Results

Serum levels of s-HER2 and TGF- β 1 were measured in 95 GTD patients, 30 pregnant controls, and 22 non-pregnant healthy women. The mean serum level of s-HER2 was 3.05 ± 1.30 ng/ml in GTD patients, 3.19 ± 1.62 ng/ml in healthy pregnant controls, and 2.77 ± 1.67 ng/ml in non-pregnant controls, as depicted in Table 1, the difference was not statistically significant ($P > 0.05$).

As presented in Table 2, mean serum level of TGF- β 1 was significantly higher in GTD patients (20.29 ± 10.68 pg/ml with 95% confidence interval (CI) of 18.10–22.48 pg/ml) compared with pregnant controls (10.26 ± 11.84 pg/ml with

Table 1 Mean serum level of s-HER2 in GTD patients, healthy pregnant controls, healthy non-pregnant controls, and follow-up group*

	Number	Mean s-HER2 serum level (ng/ml)
Total GTD patients	95	3.05 ± 1.30
Normal pregnant controls	30	3.19 ± 1.62
Normal non-pregnant controls	22	2.77 ± 1.67
Follow-up group	11	3.12 ± 0.68

*No statistically significant difference was found between patients and controls

Table 2 Mean serum level of TGF- β 1 in GTD patients, healthy pregnant controls, healthy non-pregnant controls, and follow-up group

	Mean TGF- β 1 serum level (pg/ml)	95% CI	<i>P</i> -value*
Total GTD patients	20.29 $\pm$ 10.68	18.10–22.48	N/A
Normal pregnant controls	10.26 $\pm$ 11.84	5.75–14.76	<0.001
Normal non-pregnant controls	7.27 $\pm$ 9.61	3.01–11.53	<0.001
Follow-up group	11.12 $\pm$ 10.46	5.08–17.16	<0.001

**P*-value comparing patients with other groups

CI: Confidence interval; Mean $\pm$ SD

95% CI of 5.75–14.76 pg/ml) and non-pregnant controls (7.27 $\pm$ 9.61 pg/ml with 95% CI of 3.01–11.53 pg/ml) (P <0.001). Statistically significant differences existed (P <0.001) when comparing patients with the follow-up group. In addition, the difference between TGF- β 1 serum level of patients 6 months following evacuation and serum level of control groups (pregnant or non-pregnant) did not prove to be statistically significant (P >0.05).

Comparing serum levels of s-HER2 and TGF- β 1 between subgroups of patients (55 complete hydatidiform mole, 32 persistent hydatidiform mole, and eight choriocarcinoma patients) revealed no statistically significant difference (data not given).

Discussion

GTDs comprise a spectrum of diseases with different clinical presentations that continue to be a significant reproductive health problem worldwide. The rate of metastasis in this neoplastic disease is high due to increased vascularization [2].

Tumor markers and certain immune regulatory mediators have previously been investigated in molar pregnancy and choriocarcinoma [18, 19]. The potential role of s-HER2 and TGF- β 1 as tumor markers in patients with GTDs was investigated in the present study.

HER2/neu is a member of EGFR family that has been investigated in various neoplasms; release of soluble form of HER2/neu and its alterations in the serum during malignancy course remains as a potential biomarker in certain tumors particularly in the case of breast cancer [6, 7].

In the current study, the serum level of s-HER2 was determined in GTD patients and was compared with normal healthy pregnant and non-pregnant controls. Mean concentration of s-HER2 in GTD patients and controls was not statistically different.

The role of Her2/neu in GTDs has been studied previously [15, 16]; however, no data on the serum level of HER2 in GTDs was available in the literature. In a study conducted by Tuncer et al. [15], expression of c-erbB-2 gene product was investigated immunohistochemically (IHC) on sections obtained from GTD patients; they reported the importance of EGFR-related family of oncogenes in the pathogenesis of GTDs and proposed that the increased expression of EGFR in complete mole may influence the development of persistent gestational trophoblastic tumor. Utilizing IHC, Yang et al. [16], found the overexpression of c-erbB-2 protein as a strong predictor for the malignant transformation of complete mole.

Although previous studies [15, 16] found an increased serum level of s-HER2 in GTD patients, we failed to find an association between s-HER2 determination by ELISA in serum and GTDs. This could be explained based on the fact that characteristics of antibodies used in ELISA and IHC and also sensitivity of these tests are different. As IHC is done on formalin fixed tissue compared with sera in our assay, evacuation of tissue and handling of sera should also be considered. On the other hand, differences in the proportion of GTD subtypes in our study and others [15, 16], status of treatment, and even age and ethnicity of the studied populations should be considered. Performing reverse transcription polymerase chain reaction (RT-PCR) to analyse mRNA on tissue to determine the level of HER2 gene expression could be suggested to elaborate more on this issue.

Here serum level of TGF- β 1 was also determined in the studied population. In normal cells TGF- β inhibits cellular proliferation. After tumor cells become resistant to growth inhibition by TGF- β , the tumor cells often increase their production of TGF- β . In response to increased production of TGF- β , the tumor cells, whose growth can no longer be inhibited by TGF- β , become more invasive and metastasize to distant organs due to TGF- β mediated stimulation of angiogenesis and cell motility, TGF- β also enhances the production and influences the adhesive characteristics of the extracellular matrix and suppresses the activities of infiltrating immune cells, helping the tumor escape host immunosurveillance [20–22].

Changes in serum level of TGF- β and variation in its genes have been studied in most reproductive related disorders; in various cancers elevated levels of TGF- β have been reported including gastric carcinoma [23], esophageal cancer [24], and skin malignancy [25]. Measurement of TGF- β protein level in breast cancer patients might help to identify the high-risk population early in tumor progression [11].

In the current study, TGF- β 1 serum levels were significantly higher in GTD patients compared with normal pregnant and non-pregnant controls. In addition to the

notion that TGF- β 1 is released by replicating tumor cells and facilitates tumor growth and invasion by promoting angiogenesis, affecting extracellular matrix and immunosuppression, it could also be postulated that production of self-antigens during cellular proliferation in GTD will trigger T regulatory cells and consequently more TGF- β 1 will be synthesized. TGF- β 1 released by T regulatory cells functionally suppresses the immune effector mechanism [10], giving more chance to neoplastic cells to grow and to migrate to other organs and tissues.

Our findings raise the possibility that TGF- β 1 serum level determination in GTD patients may have prognostic significance. Failing to detect a statistically significant difference in TGF- β 1 serum level between complete mole, persistent mole, or choriocarcinoma in the current study restricts its value in prediction of tumor behavior; however, the limited number of patients in the present study should be considered while interpreting the results.

Further investigations with larger sample sizes and more frequent sampling are required to elucidate the role of TGF- β 1 serum level as a predictive and prognostic factor in the management of GTDs, to nominate it as a serum marker that can be measured to have a clue of persistence or remission of GTD following diagnosis, or to follow the response to treatment.

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