

## Human epidermal growth factor receptor-2 gene amplification in gastric cancer using tissue microarray technology

Dimitrios Tsapralis, Ioannis Panayiotides, George Peros, Theodore Liakakos, Eva Karamitopoulou

Dimitrios Tsapralis, Theodore Liakakos, 3rd Department of Surgery, Attikon University Hospital, University of Athens, Rimini 1, 12464 Athens, Greece

Ioannis Panayiotides, 2nd Department of Pathology, University of Athens, Rimini 1, 12464 Athens, Greece

George Peros, 4th Department of Surgery, Attikon University Hospital, University of Athens, Rimini 1, 12464 Athens, Greece

Eva Karamitopoulou, Department of Pathology, Aretaieion University Hospital, University of Athens, Vass. Sophias 76, 11528 Athens, Greece

**Author contributions:** Tsapralis D performed the research; Panayiotides I provided paraffin blocks from gastric cancer cases; Peros G contributed patients and data; Liakakos T contributed patients and designed the research; Karamitopoulou E designed the research, analyzed the data and wrote the paper.

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Correspondence to: Eva Karamitopoulou, Assistant Professor, Department of Pathology, Aretaieion University Hospital, Vass. Sophias 76, 11528 Athens, Greece. [ekaramit@med.uoa.gr](mailto:ekaramit@med.uoa.gr)  
Telephone: +30-210-7286427 Fax: +30-210-6018459

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### Abstract

**AIM:** To assess human epidermal growth factor receptor-2 (HER2)-status in gastric cancer and matched lymph node metastases by immunohistochemistry (IHC) and chromogenic *in situ* hybridization (CISH).

**METHODS:** 120 cases of primary gastric carcinomas and 45 matched lymph node metastases from patients with full clinicopathological features were mounted onto multiple-punch and single-punch tissue microarrays, respectively, and examined for HER2 overexpression and gene amplification by IHC and CISH.

**RESULTS:** Twenty-four tumors (20%) expressed HER2 immunohistochemically. An IHC score of  $\geq 2+$  was observed in 20 tumors (16.6%). HER2 amplification was detected by CISH in 19 tumors (15.8%) and in their matched lymph node metastases. A high concordance

rate was found between HER2 positivity (as detected by IHC) and *HER2* gene amplification (as detected by CISH), since 19 of the 20 IHC positive cases were amplified (95%). All amplified cases had 2+ or 3+ IHC results. Amplification was associated with intestinal phenotype ( $P < 0.05$ ). No association with grading, staging or survival was found.

**CONCLUSION:** In gastric cancer, HER2 amplification is the main mechanism for HER2 protein overexpression and is preserved in lymph node metastases.

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**Key words:** Human epidermal growth factor receptor-2; Immunohistochemistry; Chromogenic *in situ* hybridization; Gastric cancer

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### INTRODUCTION

Alterations of the human epidermal growth factor receptor-2 (*HER2*) gene are implicated in the development and progression of many tumors<sup>[1-4]</sup>. In breast cancer, HER2 amplification has been found in about 20% of cases and was linked to poor prognosis<sup>[5,6]</sup>. Breast cancer patients with HER2 amplification have been effectively treated with the monoclonal antibody trastuzumab, a HER2 inhibitor<sup>[7-10]</sup>. Recently, a number of studies have suggested a response to trastuzumab therapy for other

cancers with HER2 amplification, including germ cell, endometrium and salivary duct carcinoma<sup>[11-13]</sup>.

In gastric cancer, HER2 amplification has been found in 7% to 27% of tumors<sup>[14-19]</sup>. Reports of trastuzumab therapy in metastatic gastric cancer showed complete tumor regression and disappearance of the metastases in two cases<sup>[20,21]</sup>. A phase III randomized study (Trastuzumab for HER2-positive metastatic gastric cancer) in patients with inoperable, metastasizing and/or recurring gastric cancer with HER2 overexpression or *HER2* gene amplification, documented that 47.3% of the patients who received trastuzumab, along with their chemotherapy, showed a significant regression of the primary tumor and/or the metastases. Moreover, trastuzumab caused a prolongation of the median survival time by 2.4 mo in all patients<sup>[22]</sup>. Based on these reports, gastric cancer patients with HER2 overexpression and/or amplification could be good candidates for trastuzumab therapy.

HER2 testing can be performed either by immunohistochemical evaluation of protein expression or by evaluating the gene copy number by *in situ* hybridization, most commonly using fluorescence *in situ* hybridization (FISH). However, while immunohistochemistry (IHC) is a relatively inexpensive, easy to perform method for most pathology laboratories, FISH is technically demanding, expensive and requires special equipment<sup>[23-25]</sup>. An alternative method, chromogenic *in situ* hybridization (CISH), is a combination of *in situ* hybridization with a detection system using a chromogen similar to IHC. Slides are visible under a light microscope and show correlation with morphology. A number of studies compared HER2 testing with IHC, FISH and CISH in breast carcinoma and have shown good correlation between CISH and FISH results<sup>[25-30]</sup>.

We evaluated HER2 overexpression and gene amplification by IHC and CISH, respectively, in 120 cases of gastric carcinoma patients and 45 matched lymph node metastases mounted onto multiple-punch and single-punch tissue microarrays respectively. Our data suggests that, in gastric cancer, HER2 amplification is the main mechanism for HER2 protein overexpression and is preserved in lymph node metastases.

## MATERIALS AND METHODS

### Patients

The current study involved 120 non-consecutive patients with gastric carcinoma, surgically treated at the 3rd and 4th Departments of Surgery, University of Athens, between 2004 and 2007. Histomorphological data were reviewed from the corresponding hematoxylin and eosin stained slides. Clinical data were obtained from corresponding reports. Clinicopathological information included: gender, age, tumor diameter, histological subtype, tumor location, pT stage, pN stage, pM stage, vascular and lymphatic invasion, survival time, and information on post-operative therapy. Characteristics of patients are summarized in Table 1.

Table 1 Characteristics of patients with gastric cancer

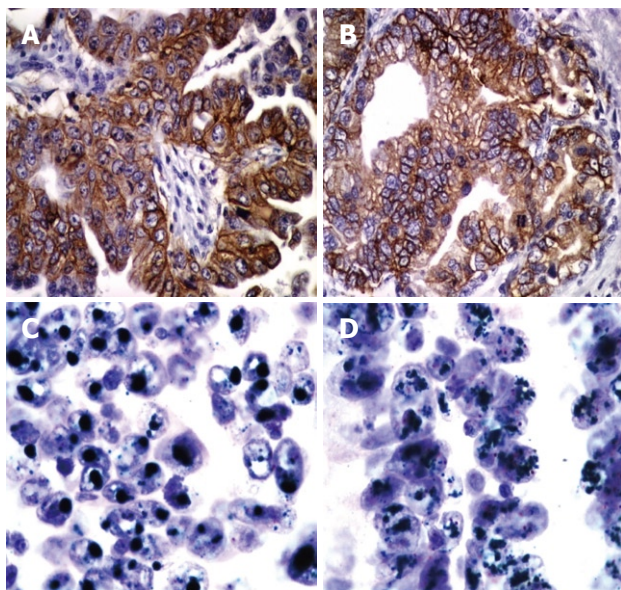
Clinicopathological feature	Frequency n (%)
Patient age at diagnosis (yr)	Mean 69.6, min-max 27-96
Tumor diameter (cm)	Mean 4.6, min-max 1.3-12
Gender	Male 84 (70) Female 36 (30)
Histological type	Intestinal 80 (66.66) Diffuse 24 (20) Mixed 16 (13.33)
Tumor location	Cardia 37 (30.8) Corpus 39 (32.5) Antrum 44 (36.66)
pT stage	pT1 15 (12.5) pT2 65 (54.16) pT3-4 40 (33.33)
pN stage	pN0 36 (30) pN1 43 (35.8) pN2+3 41 (34.2)
pM stage	pM0 105 (87.5) pM1 15 (12.5)
Tumor grade	G1-2 42 (35) G3 78 (65)
Venous invasion	Present 37 (30.8) Absent 83 (69.2)
Lymphatic invasion	Present 84 (70) Absent 36 (30)
Adjuvant therapy	None 32 (26.6) Treated 88 (73.4) Chemotherapy 55 (62.5) Chemo/Radiotherapy 33 (37.5)
5-year survival (%)	(95% CI) 38.9 (25-52)

### Specimen characteristics

Paraffin-embedded tissue blocks of primary tumors and matched positive lymph nodes were retrieved from the Department of Pathology, University of Athens. The use of this material was approved by the local Ethics committee. Two tissue microarrays (TMAs) were constructed. The first included punches from primary tumors. In order to exclude bias due to possible tumor heterogeneity, each patient had multiple tumor punches taken from formalin-fixed, paraffin-embedded blocks using a tissue cylinder with a diameter of 1 mm, which were subsequently transferred into one recipient paraffin block (3 cm × 2.5 cm) using a semiautomated tissue arrayer. Each patient had on average 5.1 tissue punches included on this array, including at least 4 tumor punches. The second TMA included single punches from matched metastatic lymph nodes in 45 patients.

### Assay methods

**IHC:** Five µm TMA sections were dewaxed and rehydrated in distilled water. Endogenous peroxidase was blocked using 0.5% H<sub>2</sub>O<sub>2</sub>. To determine the HER2 expression immunohistochemically, the HercepTest™ (Dako, Glostrup, Denmark) was used according to the manufacturer's protocol. Following pressure cooker-mediated antigen retrieval sections were incubated with the prediluted primary antibody. Control samples included normal gastric mucosa and breast cancer tissue. Immunostaining was scored by an experienced gastrointestinal pathologist following a 4-step



**Figure 1** Examples of human epidermal growth factor receptor-2 immunohistochemical expression and amplification in primary and metastatic gastric cancer. Immunohistochemistry shows strong membranous staining of human epidermal growth factor receptor-2 (HER2) in intestinal type gastric cancer (A) and (B) ( $\times 400$ ). Chromogenic *in situ* hybridization assay shows amplification of HER2 in primary gastric cancer (C) and the corresponding lymph node metastasis (D). Clustered green signals represent the amplified *HER2* gene, while red signals represent centromere 17. Cell nuclei are counterstained with hematoxylin ( $\times 1000$ ).

score (0, 1+, 2+, 3+), according to the consensus panel recommendations on HER2 scoring for gastric cancer<sup>[31]</sup>.

**CISH:** HER2 CISH was performed using a CISH HER2 probe and Immunodetection Kit (ZytoDot2C SPEC HER2/CEN 17 Probe Kit). TMA sections were deparaffinized and incubated for 5 min in 3% H<sub>2</sub>O<sub>2</sub>, followed by Heat Pretreatment Solution EDTA in a covered staining jar standing in a boiling water bath at 98 °C for 15 min. After washing in distilled water, Pepsin Solution (ES1) was applied and slides were incubated for 5 min at room temperature in a humidity chamber. Sections were then washed in distilled water, dehydrated in increasing ethanol, and air dried. ZytoDot2C SPEC HER2/CEN 17 Probe was applied and sections were covered with a coverslip sealed with a layer of hot glue. Samples were then denatured at 80 °C for 5 min, transferred to a humidity chamber and left to hybridize overnight at 37 °C. On day 2, immunodetection was performed according to the manufacturer's instructions and sections were counterstained with Hematoxylin and mounted.

### Statistical analysis

$\chi^2$  tests and contingency tables were used to analyze the relationship between IHC and CISH, and categorical parameters. Overall survival was estimated by the Kaplan-Meier method and evaluated by log-rank testing. All analysis were carried out using SAS (V9, The SAS Institute, NC, United States).

**Table 2** Lauren phenotype, human epidermal growth factor receptor-2 immunohistochemistry and chromogenic *in situ* hybridization in gastric carcinoma

		Diffuse ( <i>n</i> = 24)	Mixed ( <i>n</i> = 16)	Intestinal ( <i>n</i> = 80)
HER2 IHC	0	23	14	59
	1+	0	0	4
	2+	1	0	5
	3+	0	2	12
HER2 CISH	Non amplified	24	14	63
	Amplified	0	2	17

HER2: Human epidermal growth factor receptor-2; IHC: Immunohistochemistry; CISH: Chromogenic *in situ* hybridization.

## RESULTS

### HER2 immunohistochemistry

HER2 protein expression was observed in 24 of the 120 gastric carcinomas (20%). In more detail, one of the 24 diffuse type carcinomas (4.16%) and 23 of the 96 intestinal and mixed type carcinomas (23.95%) showed HER2 protein expression. Immunostaining was always membrane bound and showed basolateral predominance (Figure 1A and B). Immunostaining in mixed type carcinomas was restricted in the intestinal type component. Quantitative analysis of the immunostaining, according to the consensus panel recommendations on HER2 scoring for gastric cancer, resulted in fourteen 3+ cases (11.66%), six 2+ cases (5%) and four 1+ cases (3.33%)<sup>[31]</sup> (Table 2). IHC was interpretable in 652 of the 660 spots (98.8%). Reasons for non-interpretable results were missing tissue spots or absence of tumor tissue.

### HER2 CISH

Tissue spots were scanned for possible intratumoral heterogeneity by using a 10 $\times$  objective lens. CISH hybridization signals of the *HER2* gene appeared as dark green-colored dot-shaped signals. The chromosome 17 centromeric regions appeared as bright red-colored dot-shaped signals. Areas of necrosis and overlapping nuclei were avoided. Signal enumeration was performed using the 40 $\times$  objective lens of a light microscope. HER2 amplification was observed in 19 of the 120 primary gastric carcinomas (15.8%). Amplified cases showed intratumoral heterogeneity with areas of low amplification where HER2 signals appeared as multiple dots or small clusters, and areas of high amplification with presence of large, green *HER2* gene signal clusters (Figure 1C and D). All amplified cases had 2+ or 3+ IHC results (Table 2). HER2 amplification showed significant association with histologic tumor type. Seventeen (21.25%) of the 80 intestinal and two (12.5%) of the 16 mixed type cancers, but 0 (0%) of the diffuse type cancers, were amplified ( $P < 0.002$ , Table 2). No association between *HER2* gene amplification and tumor grade, size, stage or localization was found.

**Table 3** Comparison of human epidermal growth factor receptor-2 immunohistochemistry and chromogenic *in situ* hybridization in primary gastric carcinomas and matched lymph node metastases

	HER2 IHC			HER2 CISH		
	PT	LNM	Concordance	PT	LNM	Concordance
Positive	7	6	85.7%	6	6	100%
Negative	38	39	97.4%	39	39	100%

HER2: Human epidermal growth factor receptor-2; IHC: Immunohistochemistry; CISH: Chromogenic *in situ* hybridization; PT: Primary gastric cancer; LNM: Lymph node metastases.

### HER2 alterations in lymph node metastases

Comparative analysis of primary tumors and corresponding lymph node metastases, performed in 45 cases, showed a high concordance in the presence of HER2 overexpression or amplification ( $P < 0.0001$ , Table 3).

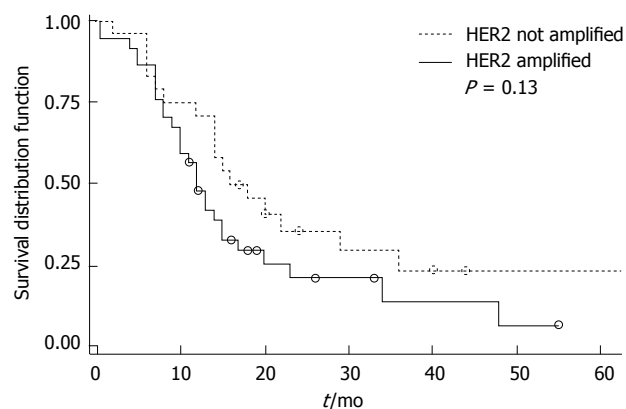
### Clinicopathological characteristics

Increasing tumor grade and stage were associated with reduced patient survival ( $P < 0.0001$  each). No correlation was observed between patient survival and HER2 overexpression or amplification, even after including postoperative therapy of the patients and location of the tumors in the analysis (Figure 2).

## DISCUSSION

The HER2 protein is a frequently analyzed gene product, especially in breast cancer. Recent studies have examined HER2 expression in other tumor types, including gastric cancer<sup>[18,32,33]</sup>. Immunohistochemical HER2 expression and protein positivity ( $\geq 2+$ ) was observed in 20% and 14.58%, respectively, of the 120 gastric carcinomas in our study. These results are in keeping with previous reports demonstrating similar frequencies of HER2 overexpression in gastric cancer<sup>[18,32,33]</sup>. HER2 positivity was observed in 23% of the cases in the study by Yano *et al.*<sup>[18]</sup> and in 22.6% of the cases in the study by Kim *et al.*<sup>[33]</sup>. More recently, a TMA study of 166 gastric carcinomas by Marx *et al.*<sup>[32]</sup> found a HER2 positivity rate of 17% and a strong correlation between IHC and HER2 gene amplification detected by FISH. In our cases, comparable to others, HER2 overexpression and/or amplification were almost exclusively found in gastric cancers of the intestinal type. This finding supports the presence of different molecular characteristics between the main histologic tumor types that seem to develop through different molecular pathways.

No correlation between HER2 overexpression and/or amplification and tumor localization could be demonstrated in our study. This might be contradictory to previous studies where a high frequency of HER2 expression was reported in cardia carcinomas<sup>[34]</sup>. However, adenocarcinomas of the gastroesophageal junction, many of which are Barrett carcinomas, are known to have a high rate of HER2 amplification and cannot always be differentiated from cardia carcinomas. This may lead to



**Figure 2** Kaplan-Meier curve for disease-specific survival and human epidermal growth factor receptor-2 amplification in gastric carcinomas. HER2: Human epidermal growth factor receptor-2.

an artificial increase in the rate of HER2 expression reported in cardia carcinomas.

A high concordance rate was found between HER2 positivity, as detected by IHC, and HER2 gene amplification, as detected by CISH, since 19 of the 20 IHC positive cases (i.e.,  $\geq 2+$ ) were amplified (95%). Many of them (73.6%) had high HER2 gene amplification. However, one HER2 2+ case, detected in a diffuse carcinoma, was not found to be amplified, which may be attributed to a technical error of IHC associated with formalin fixation of the tissue. Alternatively, other mechanisms of HER2 protein overexpression can contribute to inconsistencies between IHC and ISH results.

In the present study we found a high concordance rate of HER2 status between primary tumors and their corresponding lymph node metastases. This finding is in keeping with previous published results, where HER2 amplification status was found to be almost identical in the primary gastric carcinomas and their corresponding lymph node metastases<sup>[32]</sup> and provides further evidence for the role of HER2 amplification in gastric cancer.

Our findings support gastric cancer HER2 amplification as being the main mechanism that leads to HER2 protein overexpression. Similar findings have previously been reported for esophageal cancer<sup>[35]</sup>. Moreover, Tapia *et al.*<sup>[19]</sup>, in a large-scale TMA study, examined more than 4000 samples from 120 different tumors and could not find any tumors with HER2 overexpression in the absence of gene amplification.

HER2 testing has developed over a number of years, and many retrospective studies using formalin-fixed, paraffin-embedded material and different methodologies have provided inconsistent results<sup>[36,37]</sup>. Correlation between technical methods can be used to obtain a high concordance rate and to better define the assays with the best ability to identify patient groups that would benefit from a targeted therapy. Although in breast cancer HER2 IHC is an acceptable method of predicting HER2 status, previous studies have shown marked variation in different diagnostic laboratories regarding the interpretation of positive staining<sup>[25]</sup>. On the other hand, FISH is

technically demanding, expensive and does not produce a permanent archival slide. In our study, CISH testing showed a good correlation to IHC, and therefore seems to represent a reliable methodology for HER2 testing. Moreover, the slides are visible under a light microscope and show good correlation with tumor morphology.

In this study, we used the tissue microarray technique using multiple tissue punches per case to account for possible heterogeneity in terms of protein expression or gene amplification in the primary tumor. Each patient had an average of 4 tumor punches taken. This is particularly important for HER2 testing in gastric cancer since considerable heterogeneity concerning gene amplification has been reported in many studies<sup>[38,39]</sup>. Such heterogeneity was also noted in our study, since in many amplified cases, areas with low and high amplification were found within the same tumor. Multiple sampling thus helped to minimize possible biases in evaluation, as suggested by Goethals *et al*<sup>[40]</sup>, who recommended that at least four punches of primary tumor are required to account for possible heterogeneity. However, a single tissue punch was sampled in the case of lymph nodes since the issue of heterogeneity may be substantially less important. Our study also benefits from complete clinicopathological and follow-up characterization of patients. In contrast to previous studies<sup>[34,41]</sup> we could not demonstrate any association between HER2 positivity and clinical outcome.

Our study provides evidence supporting gastric cancer HER2 amplification as being the main mechanism for HER2 protein overexpression and that HER2 amplification is preserved in the lymph node metastases. Therefore, gastric cancer patients with HER2 overexpression and/or amplification seem to be good candidates for anti-HER2 therapy. Moreover, CISH is a reliable and inexpensive method that can be used for HER2 testing of gastric cancer.

## COMMENTS

### Background

Alterations of the human epidermal growth factor receptor-2 (HER2) gene are implicated in the development and progression of many tumors. Breast cancer patients with HER2 amplification have been effectively treated with the monoclonal antibody trastuzumab, a HER2 inhibitor. Recently, a number of studies have suggested a response to trastuzumab therapy for other cancers with HER2 amplification, including gastric cancer, where HER2 amplification has been found in 7% to 27% of the tumors. Reports of trastuzumab therapy in metastatic gastric cancer showed complete tumor regression and disappearance of the metastases in two cases. Based on these reports, gastric cancer patients with HER2 overexpression and/or amplification could be good candidates for trastuzumab therapy.

### Research frontiers

The HER2 protein is a frequently analyzed gene product, especially in breast cancer. Recent studies have examined HER2 expression in other tumor types, including gastric cancer. The frequency and significance of HER-2/neu amplification in gastric carcinoma are investigated. In this study, immunohistochemical HER2 expression and protein positivity ( $\geq 2+$ ) was observed in 20% and 14.58%, respectively, of the 120 gastric carcinomas.

### Innovations and breakthroughs

The authors evaluated HER2 overexpression and gene amplification by immunohistochemistry and chromogenic *in situ* hybridization (CISH) respectively, in

120 cases of gastric carcinoma patients and 45 matched lymph node metastases mounted onto multiple-punch and single-punch tissue microarrays respectively. The data suggest that, in gastric cancer, HER2 amplification is the main mechanism for HER2 protein overexpression and is preserved in the lymph node metastases.

### Applications

This study provides evidence supporting gastric cancer HER2 amplification as being the main mechanism for HER2 protein overexpression and that HER2 amplification is preserved in the lymph node metastases. Therefore, gastric cancer patients with HER2 overexpression and/or amplification seem to be good candidates for anti-HER2 therapy. Moreover, CISH is a reliable and inexpensive method that can be used for HER2 testing of gastric cancer.

### Peer review

The authors studied HER2/neu expression and gene amplification in a cohort of Greek patients with gastric cancer. The manuscript includes 2 aspects. Firstly, they describe the expression/amplification of HER2-neu in the study group. Secondly, they highlight the potential applicability of CISH as a routine method. Data are mostly well documented and conclusions drawn are appropriate.

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