

# Concordance between human epidermal growth factor receptor 2 status by immunohistochemistry and next-generation sequencing in tissue and blood samples from gastric and gastroesophageal junction cancers

Steven B Maron,<sup>1</sup> Kathleen Burke,<sup>2</sup> Pooja Gupta,<sup>3\*</sup> Kristen Sarlo,<sup>4</sup> Marija Tesic-Schnell,<sup>5</sup> Michele Sue-Ann Woo,<sup>4</sup>

Samuel J Klempner<sup>6</sup>

<sup>1</sup>Memorial Sloan Kettering Cancer Center, New York, NY, US; <sup>2</sup>Tempus AI, Inc., Chicago, IL, US; <sup>3</sup><Department>, <Function>, <Business Unit>, AstraZeneca, Gaithersburg, MD, US; <sup>4</sup>Daiichi Sankyo, Inc., Basking Ridge, NJ, US; <sup>5</sup>US Medical Affairs, Oncology Business Unit, AstraZeneca, Gaithersburg, MD, US; <sup>6</sup>Massachusetts General Hospital Cancer Center, Boston, MA, US

\*Has moved institution since the publication was initiated; institution at time of publication: <TBC>

## Objectives

- Evaluate the concordance of detecting human epidermal growth factor receptor 2 (*HER2* [*ERBB2*]) gene amplification by tissue or plasma next-generation sequencing (NGS) with *HER2* immunohistochemistry (IHC) / in situ hybridization (ISH) status in patients with gastric or gastroesophageal junction cancers (GC/GEJC) using the Tempus<sup>®</sup> database
- Assess the correlation in *HER2* copy number between paired tissue and plasma samples in patients with GC/GEJC

## Conclusions

- This analysis showed moderate concordance of *HER2* amplification detection by NGS in tissue (positive percent agreement [PPA], 69.9%) or plasma (PPA, 57.5%) with *HER2*-positive (*HER2*+; IHC 3+ or IHC 2+ / ISH-positive [ISH+]) GC/GEJC determined by IHC/ISH assessment
- High concordance (>90%) per negative percent agreement (NPA) was observed for *HER2* amplification determined by tissue/plasma NGS with *HER2*-negative (*HER2*-; IHC 0, IHC 1+, or IHC 2+ / ISH-negative [ISH-]) GC/GEJC determined by IHC/ISH
- As NGS was not sufficient to detect all patients with *HER2*+ GC/GEJC, these data support IHC/ISH as the standard testing method for identifying patients with *HER2*+ GC/GEJC
  - Further study is needed to determine if NGS testing could complement IHC/ISH in identifying patients who may benefit from *HER2*-directed therapy
- Limitations of the population characteristics and statistical analyses should be considered when interpreting the results of this analysis

## Plain language summary

**Why did we perform this research?** Human epidermal growth factor receptor 2 (*HER2*) is a protein found at higher-than-normal levels on the cell surface of some tumors.<sup>1</sup> The levels of *HER2* can be measured, which can help identify people who may benefit from treatments that specifically target *HER2*.<sup>2</sup> The standard method to identify people with tumors that have high levels of *HER2* (known as *HER2*-positive [*HER2*+]) tumors is by testing tissue samples using two techniques called immunohistochemistry (IHC) and in situ hybridization (ISH). Next-generation sequencing (NGS) is a technique already used in clinical practice to identify a range of genetic variations linked to disease, including whether there are higher-than-normal levels of the *HER2* gene (known as *HER2* amplification). NGS may offer an alternative method to identify *HER2*+ tumors; however, it is currently unclear how well NGS detection of *HER2* amplification can capture people with *HER2*+ cancers. This study assessed whether *HER2* amplification (determined by NGS testing) could identify people with *HER2*+ tumors (identified by IHC/ISH) of the stomach (gastric cancers [GC]) or where the stomach meets the esophagus (gastroesophageal junction cancers [GEJC]).

**How did we perform this research?** Clinical records of people with GC/GEJC were sourced from the Tempus<sup>®</sup> database. The primary analysis assessed whether *HER2* gene amplification determined by tissue or blood NGS testing could identify people with *HER2*+ tumors confirmed by IHC/ISH. Only people who had tissue or blood samples taken for NGS testing within 30 days of their samples taken for IHC/ISH assessment were included in the primary analysis. The secondary analysis evaluated samples from people with GC/GEJC who had both tissue and blood samples (taken within 30 days of each other) that underwent NGS testing and assessed whether the NGS results from the tissue sample were similar to those from the blood sample in the same person.

**What were the findings of this research?** Approximately 70% of people with *HER2*+ GC/GEJC determined by IHC/ISH had *HER2* amplification detected by NGS testing of cancer tissue and approximately 58% of people with *HER2*+ GC/GEJC determined by IHC/ISH had *HER2* amplification detected by NGS testing of their blood. Of the people without *HER2*+ GC/GEJC (known as *HER2*-negative [*HER2*-] GC/GEJC), NGS testing of tissue or blood showed that the majority (>90%) of them did not have *HER2* amplification.

**What are the implications of this research?** Results from NGS testing for *HER2* amplification could identify almost all people with *HER2*- GC/GEJC and some people with *HER2*+ GC/GEJC. These findings support IHC/ISH testing as the gold standard for determining *HER2*+ cancers. Further study is needed to determine if NGS testing could complement standard IHC/ISH testing in identifying people with GC/GEJC who may benefit from *HER2*-directed therapy.

**Where can I access more information?** For more information about this study, please reach out to the presenter, Dr. Maron at [marons@mskcc.org](mailto:marons@mskcc.org), or the corresponding author, Dr Klempner at [sklempner@mgb.org](mailto:sklempner@mgb.org).

1. Uzunpamk B, et al. *Ann Oncol*. 2023;34:1035–1046; 2. Zhang H, et al. *Histopathology*. 2024;85:3–19



Poster



Supplementary material

Please scan this quick response (QR) code with your smartphone camera or app to obtain a copy of these materials. Alternatively, please click on the following link: [\[hyperlink to the QR code landing page\]](#)

Copies of this poster obtained through this QR code are for personal use only and may not be reproduced without permission from ASCO GI 2025 and the authors of this poster.

This study was sponsored by AstraZeneca. In March 2019, AstraZeneca entered into a global development and commercialization collaboration agreement with Daiichi Sankyo for trastuzumab deruxtecan (T-DXd; DS-8201).

Poster presented at ASCO GI 2025, January 23–25, San Francisco, CA, US and online, by Dr Steven B Maron. Corresponding author: Dr Samuel J Klempner ([sklempner@mgb.org](mailto:sklempner@mgb.org)).

## Introduction

- HER2* expression is seen in a wide range of solid tumor types and is often associated with poor prognosis in various cancers<sup>1–5</sup>
  - An estimated 18% of patients with GC/GEJC have *HER2*+ (IHC 3+ or IHC 2+ / ISH+) tumors<sup>6</sup>
- HER2*-directed therapy has demonstrated clinically meaningful benefit in patients with *HER2*+ advanced GC/GEJC,<sup>7,8</sup> with current guidelines recommending *HER2* testing to identify those with metastatic disease who may benefit from *HER2*-directed therapy<sup>9</sup>
- IHC/ISH are the standard testing methods for determining *HER2* positivity in clinical practice
- Although NGS is a testing method commonly used by clinical oncology specialists to identify a panel of disease biomarkers, including *HER2* (*ERBB2*) amplification, there are limited data on whether NGS can identify patients with *HER2*+ GC/GEJC
- This study assessed the concordance of detecting *HER2* amplification by tissue or plasma NGS with *HER2* status determined by IHC/ISH in advanced GC/GEJC

## Results

- A total of 1578 records of patients with GC/GEJC were identified from the Tempus<sup>®</sup> database for the primary analysis
  - Tissue NGS testing was performed in 1507/1578 patients and plasma NGS testing in 305/1578 patients; demographics and clinical characteristics of these patients are shown in **Table 1**
  - In patients with tissue and blood NGS results, 973 (64.6%) and 204 (66.9%) had Stage 4 disease and 189 (12.5%) and 21 (6.9%) had received prior chemotherapy, respectively
- According to IHC/ISH status, 229 (15.1%) and 40 (13.1%) patients were *HER2*+ among those with tissue and plasma NGS testing, respectively

**Table 1. Demographics and clinical characteristics of patients identified for the primary analysis**

n (%)	Patients with a tissue NGS result (n=1507)*	Patients with a plasma NGS result (n=305)†
<b>Age, years†</b>		
<50	261 (17.3)	57 (18.7)
50–60	304 (20.2)	74 (24.3)
60–70	450 (29.9)	76 (24.9)
70–80	359 (23.8)	74 (24.3)
≥80	132 (8.8)	24 (7.9)
Not available	1 (0.1)	0 (0)
<b>Sex, male</b>	1005 (66.7)	201 (65.9)
<b>Primary disease site</b>		
Cardia	605 (40.1)	104 (34.1)
Stomach	902 (59.9)	201 (65.9)
<b>Disease stage†</b>		
1–3	191 (12.7)	31 (10.2)
4	973 (64.6)	204 (66.9)
Not available	343 (22.8)	70 (23.0)
<b>HER2 IHC/ISH status</b>		
IHC 0/1+	987 (65.5)	217 (71.1)
IHC 2+ / ISH unknown‡	158 (10.5)	32 (10.5)
IHC 2+ / ISH–	133 (8.8)	16 (5.2)
IHC 2+ / ISH+	47 (3.1)	10 (3.3)
IHC 3+	182 (12.1)	30 (9.8)
<b>Received prior chemotherapy</b>	189 (12.5)	21 (6.9)

\*Patients with results from a tissue NGS xT panel (xT.v1, xT.v2, xT.v3, xT.v4); †patients with results from a plasma NGS xF.v2 panel; ‡patients with *HER2* IHC 2+ tumors without an ISH result were excluded from concordance analysis. *HER2*, human epidermal growth factor receptor 2; ISH, in situ hybridization; IHC, immunohistochemistry; NGS, next-generation sequencing

## Acknowledgments

The authors acknowledge Arthur Lambert (employee of AstraZeneca) and Angelica Nunes (employee of Daiichi Sankyo) for interpretation and scientific review of data, and feedback, and Yoshinobu Shiose (employee of Daiichi Sankyo) for scientific review of data and feedback. Under the guidance of the authors and in accordance with Good Publications Practice (GPP), medical writing and editorial support was provided by Samuel Shields, PhD, of Helios Medical Communications, part of the Helios Global Group, and was funded by AstraZeneca

## Methods

### Patient populations

- De-identified clinical and molecular data from patients with cancer in the US were retrospectively identified in the Tempus<sup>®</sup> database
- For the primary analysis, patients with GC/GEJC, a *HER2* IHC score (IHC 0–3+) curated by clinicians, and tissue or plasma samples that underwent NGS testing were included
  - Only patients who had samples for NGS collected within 30 days of the reported IHC sample collection date were evaluated
- For the secondary analysis, patients with GC/GEJC and a curated *HER2* IHC score who underwent tissue and plasma NGS testing (NGS samples collected within 30 days of each other; variable allele frequency >5%) were included
  - As this analysis exclusively assessed NGS results, patients were not excluded based on IHC sample collection date

### Concordance of *HER2* amplification status, determined by NGS, with *HER2* IHC/ISH status

- Of the patients identified as *HER2* amplified by tissue and plasma NGS, 95.2% and 92.0% were *HER2*+ by IHC/ISH, respectively
- PPA of *HER2* amplification by tissue NGS with *HER2*+ tumors was 69.9% and *HER2* amplification by plasma NGS with *HER2*+ tumors was 57.5% (**Table 2**)
- NPA of *HER2* amplification by tissue NGS with *HER2*- tumors was 99.3% and *HER2* amplification by plasma NGS with *HER2*- tumors was 99.1% (**Table 3**)
- Correlation between IHC/ISH status with tissue *HER2* major copy number and plasma *HER2* log<sub>2</sub>(copy number) is shown in **Figures 1 and 2**, respectively

**Table 2. Comparison of tissue *HER2* amplification status, determined by NGS, with *HER2* IHC/ISH status\***

Tissue <i>HER2</i> amplification status, n	HER2 IHC/ISH status, n		Total
	HER2+ (IHC 3+ or IHC 2+ / ISH+)	HER2- (IHC 0, IHC 1+, or IHC 2+ / ISH-)	
<b>Detected</b>	160	8	168
<b>Not detected</b>	69	1112	1181
<b>Total</b>	229	1120	
	<b>PPA 69.9%</b>	<b>NPA 99.3%</b>	

\*Patients with *HER2* IHC 2+ tumors with an unknown ISH result were excluded. *HER2*, human epidermal growth factor receptor 2; ISH, in situ hybridization; IHC, immunohistochemistry; NGS, next-generation sequencing; NPA, negative percent agreement; PPA, positive percent agreement

**Table 3. Comparison of plasma *HER2* amplification status, determined by NGS, with *HER2* IHC/ISH status\***

Plasma <i>HER2</i> amplification status, n	HER2 IHC/ISH status, n		Total
	HER2+ (IHC 3+ or IHC 2+ / ISH+)	HER2- (IHC 0, IHC 1+, or IHC 2+ / ISH-)	
<b>Detected</b>	23	2	25
<b>Not detected</b>	17	231	248
<b>Total</b>	40	233	
	<b>PPA 57.5%</b>	<b>NPA 99.1%</b>	

\*Patients with *HER2* IHC 2+ tumors with an unknown ISH result were excluded. *HER2*, human epidermal growth factor receptor 2; ISH, in situ hybridization; IHC, immunohistochemistry; NGS, next-generation sequencing; NPA, negative percent agreement; PPA, positive percent agreement

## Disclosures – Dr Steven B Maron

Received honoraria from American Physician CME, Clarion Healthcare, Clinical Care Options., Medpage, MJH Healthcare Holdings LLC, OncoViews, Physicians' Education Resource, Practicing Clinician Exchange, and Vindico Medical Education; had a consulting or advisory role with Amgen, Calcium Company, Daiichi Sankyo, Elevation Oncology, Novartis, Pinetree Therapeutics, and Purple Biotech; received research funding from Epic Sciences and Guardant Health; and received travel, accommodation, or expenses from AstraZeneca

### Testing methods

- Concordance was assessed according to PPA and NPA
  - PPA was defined as the percentage of patients with *HER2* amplification detected in the subgroup with *HER2*+ (IHC 3+ or IHC 2+ / ISH+) tumors
  - NPA was defined as the percentage of patients with *HER2* amplification not detected in the subgroup with *HER2*- (IHC 0, 1+, or 2+ / ISH-) tumors
- The thresholds to determine tissue and plasma *HER2* amplification were defined as *HER2* copy number ≥8 or log<sub>2</sub>(copy number) ≥0.5, respectively
  - Additional amplifications were captured by manual inspection
- Statistical analyses are reported in a descriptive manner; no confirmatory hypothesis testing was planned

### Primary analysis

- Evaluate the concordance of tissue *HER2* amplification status, determined by NGS, with *HER2* IHC/ISH status
- Evaluate the concordance of plasma *HER2* amplification status, determined by NGS, with *HER2* IHC/ISH status

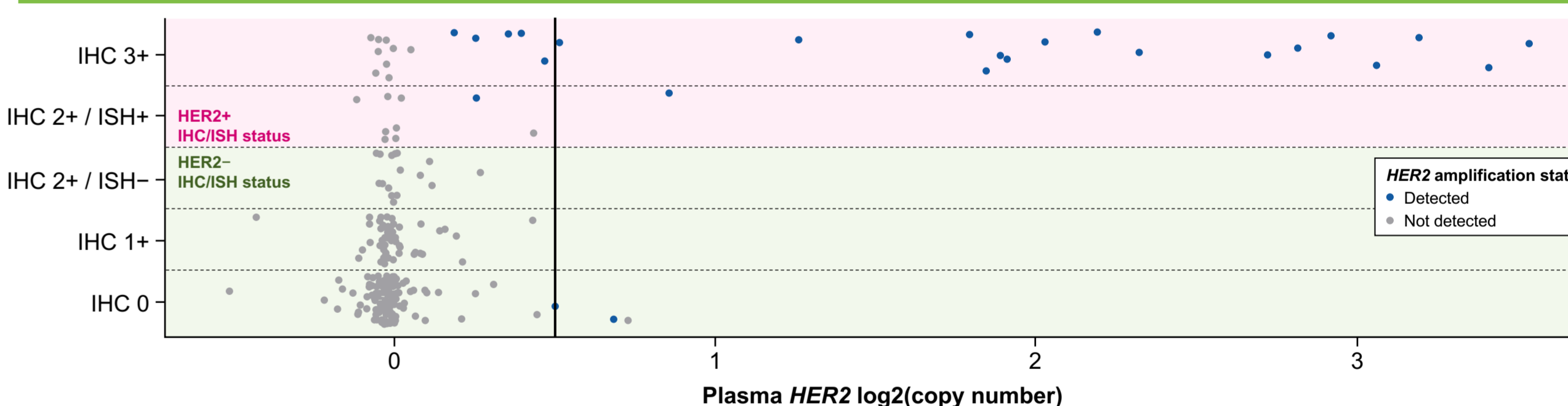
### Secondary analysis

- Assess the correlation between *HER2* log<sub>2</sub>(copy number) calling in paired tissue and plasma NGS

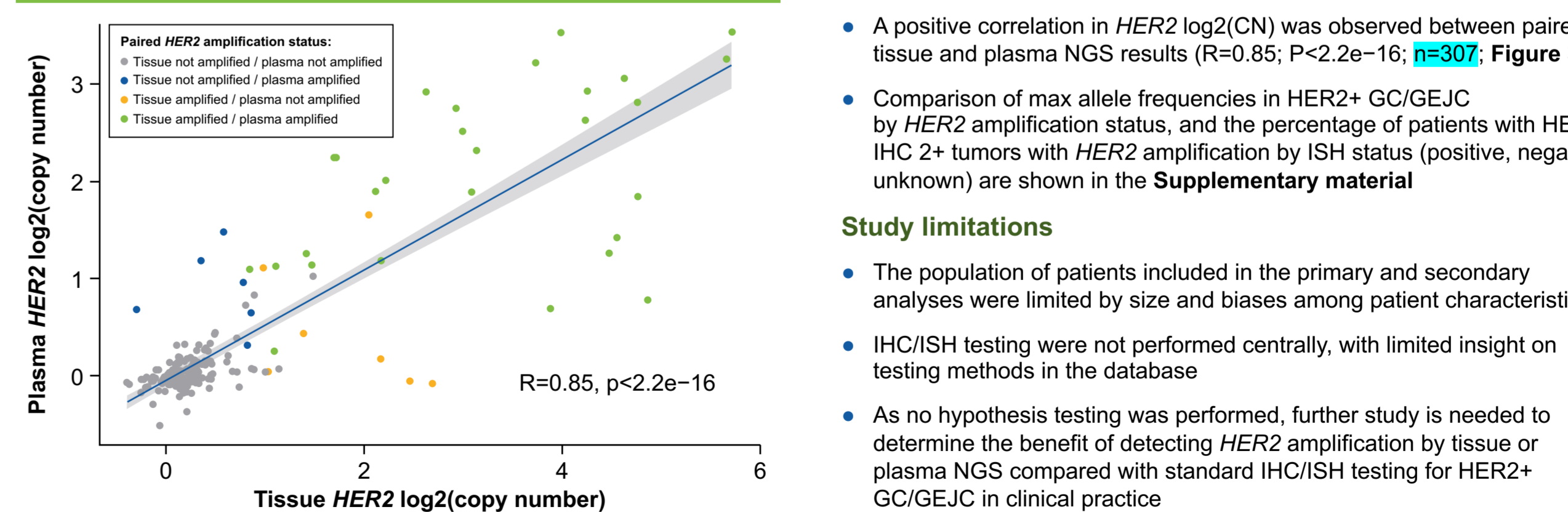
**Figure 1. Correlation of tissue NGS with *HER2* IHC/ISH status**



**Figure 2. Correlation of plasma NGS with *HER2* IHC/ISH status**



**Figure 3. Correlation of paired tissue and plasma *HER2* log<sub>2</sub>(copy number)**



### Correlation of paired tissue and plasma *HER2* log<sub>2</sub>(copy number)

- A positive correlation in *HER2* log<sub>2</sub>(CN) was observed between paired tissue and plasma NGS results (R=0.85; P<2.2e-16; n=307; **Figure 3**)
- Comparison of max allele frequencies in *HER2*+ GC/GEJC by *HER2* amplification status, and the percentage of patients with *HER2* IHC 2+ tumors with *HER2* amplification by ISH status (positive, negative, unknown) are shown in the **Supplementary material**

### Study limitations

- The population of patients included in the primary and secondary analyses were limited by size and biases among patient characteristics
- IHC/ISH testing were not performed centrally, with limited insight on testing methods in the database
- As no hypothesis testing was performed, further study is needed to determine the benefit of detecting *HER2* amplification by tissue or plasma NGS compared with standard IHC/ISH testing for *HER2*+ GC/GEJC in clinical practice

## References

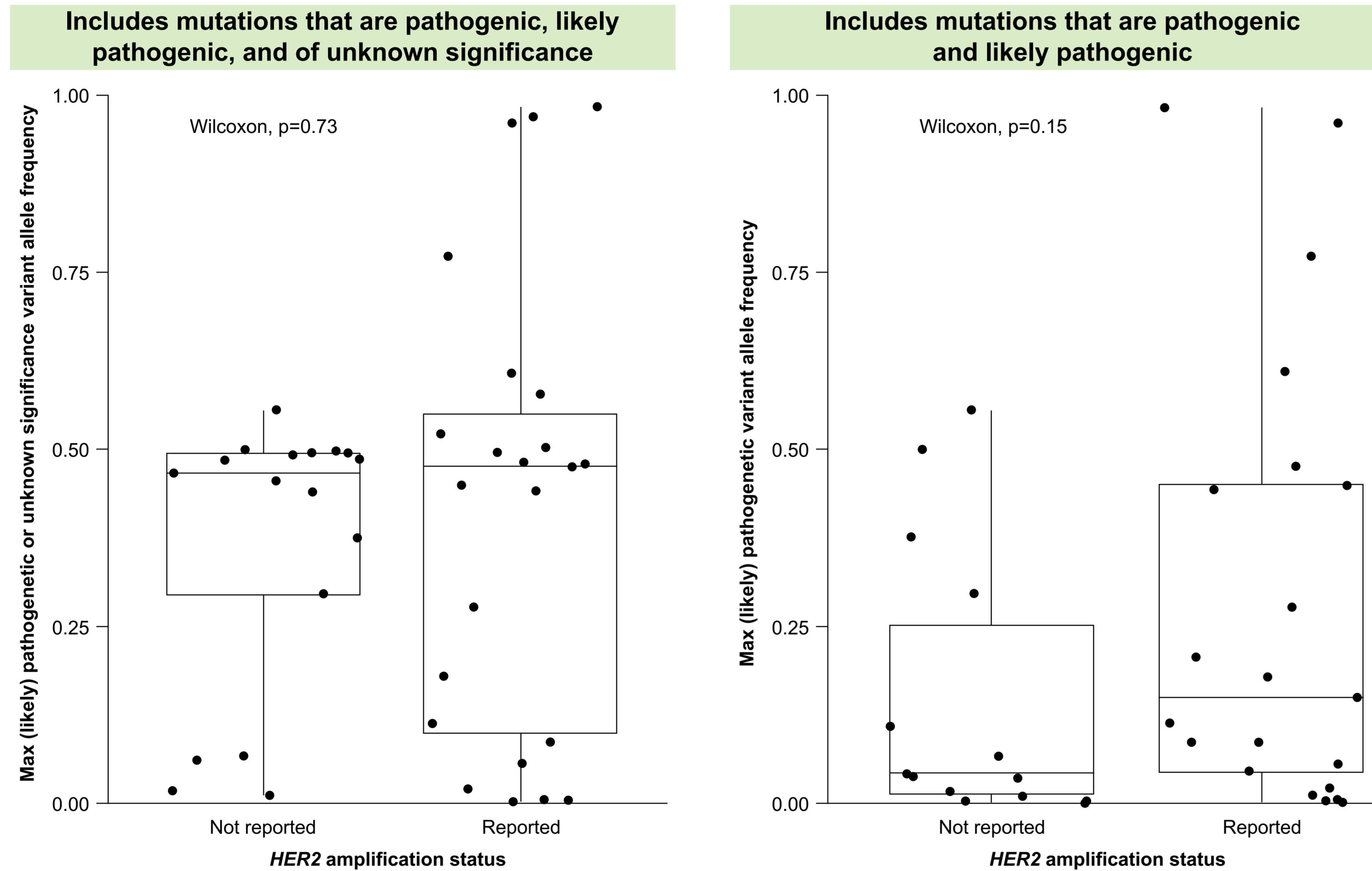
- Yan M, et al. *Cancer Metastasis Rev*. 2015;34:157–164
- Li Z, et al. *eBioMedicine*. 2020;62:103074
- Uzunpamk B, et al. *Ann Oncol*. 2023;34:1035–1046
- King F, et al. *Mol Cancer*. 2023;22:6
- Halle MK, et al. *Br J Cancer*. 2018;118:378–387
- Van Cutsem E, et al. *Gastric Cancer*. 2015;18:476–484
- Shitara K, et al. *N Eng J Med*. 2020;382:2419–2430
- Bang Y-J, et al. *Lancet*. 2010;376:687–697
- Lordick F, et al. *Ann Oncol*. 2022;33:1005–1020



Please note that these data are placed on this slide to allow for easy review. Should these data be included as a supplement, they will be uploaded as images to the QR code.

## Supplemental results

Figure S1. Comparison of max allele frequencies in HER2+\* GC/GEJC by HER2 amplification status



Data are shown as supplementary material to the poster (abstract number: 478), titled 'Concordance between human epidermal growth factor receptor 2 status by immunohistochemistry and next-generation sequencing in tissue and blood samples from gastric and gastroesophageal junction cancers', presented at ASCO GI 2025, January 23–25, San Francisco, CA, US and online. \*HER2+ was defined as IHC 3+ or IHC 2+ / ISH+ tumors. GC, gastric cancer; GEJC, gastroesophageal junction cancer; HER2, human epidermal growth-factor receptor 2; IHC, immunohistochemistry; ISH, in situ hybridization

Table S1. HER2 amplification status in patients with HER2 IHC 2+ GC/GEJC by ISH status

ISH status (IHC 2+ patients)	HER2 copy number variant amplified	HER2 copy number variant not amplified	Proportion amplified, %
Positive	9	21	30.0
Negative	0	16	0
Unknown	5	27	15.6

Data are shown as supplementary material to the poster (abstract number: 478), titled 'Concordance between human epidermal growth factor receptor 2 status by immunohistochemistry and next-generation sequencing in tissue and blood samples from gastric and gastroesophageal junction cancers', presented at ASCO GI 2025, January 23–25, San Francisco, CA, US and online. \*HER2+ was defined as IHC 3+ or IHC 2+ / ISH+ tumors. GC, gastric cancer; GEJC, gastroesophageal junction cancer; HER2, human epidermal growth-factor receptor 2; IHC, immunohistochemistry; ISH, in situ hybridization