Review Article

HER-2 status and its clinicopathologic significance in breast cancer in patients from southwest China: re-evaluation of correlation between results from FISH and IHC

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Abstract: This study aimed to re-evaluate the concordance between FISH and IHC for HER-2 status, and evaluate its clinicopathologic significance in breast cancer by the new guideline (2014) in China. We determined the HER-2 status in 589 cases of invasive breast cancer, the concordance and correlation between these two results and their relationship to cliniopathological characteristics were evaluated. The rate of HER-2 gene amplification identified by FISH was 31.07% (183/589). Concordance was detected in 93.67% with IHC 0/+, 32.26% in IHC 2+, and 61.16% in IHC 3+. The concordance of 54.67% was observed with a Kappa coefficient of 0.189 (P = 0.000), and a positive correlation was found between IHC and FISH (r = 0.427, P = 0.000). Moreover, the expression of ER was negatively correlated with HER-2 gene amplification and the expression of Her-2/neu protein (r = -0.419, P < 0.001; r = -0.144, P < 0.001; respectively), and the PR expression and ER/PR status was negatively correlated with HER-2 gene amplification (r = -0.226, P < 0.001; r = -0.258, P < 0.001), but not significantly with Her-2/neu expression. Interestingly, the tumor size and histological grade were positive correlation with HER-2/neu protein expression. The coincidently positive rate between FISH and IHC in breast cancer is very high, IHC score 0/+ is strongly consistent with HER-2 gene amplification, IHC score 2+~3+ is a preferred method to detect the HER-2 gene status, FISH still remains a standard method of evaluation of the HER-2 gene amplification.

Keywords: Breast neoplasms, human epidermal growth factor receptor 2 (HER-2), clinical pathological features, fluorescence in situ hybridization (FISH), immunohistochemistry (IHC)

Introduction

The human epidermal growth factor receptor-2 (HER-2) proto-oncogene, which plays a crucial role in the proliferation and differentiation of cells, is located on chromosome 17q11.2-q12, and encodes Her-2/neu protein. Amplification of this gene is associated with rapid progression of the disease, metastatic potential and resistance to tamoxifen [1]. Patients who with HER-2 gene amplification and Her-2/neu protein overexpression may indicate a worse situation, and are less responsive to hormonal therapies, whereas the use of anti-Her-2/neu monoclonal antibody, trastuzumab (Herceptin), especially with the combination with chemotherapy, is effective for those populations [2].

Therefore, to evaluate HER-2 status has become critical for treatment of breast cancer patients. Currently, the most commonly used method is immunohistochemistry (IHC) and fluorescence in situ hybridization (FISH), whereas the correlation between the results from these two methods, as well as the significance of the correlation in clinicopathology of breast cancer, has not been well established yet, on the basis of a new guideline in China.

Materials and methods

Specimens

A total of 589 paraffin-embedded samples from patients of invasive breast cancer were

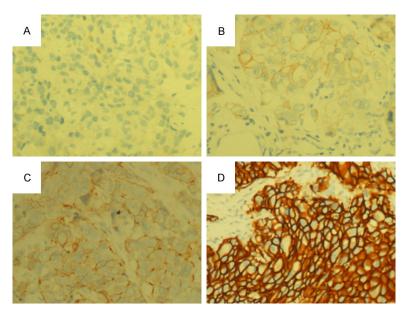


Figure 1. IHC scored of Her-2/neu protein. A. IHC 0: Negative membrane staining, ×200. B. IHC 1+: Weak membrane staining, ×200. C. IHC 2+ Moderate membrane staining, ×200. D. IHC 3+: Strong membrane staining, ×200.

The sections were then incubated with the primary antibody (erbB-2, 1:100, clone 4B5, Roche, Germany) at 37°C for 30 min. Thereafter, sections were rinsed with Trisbuffered saline (TBS) three times, and then incubated with the secondary antibody (DAKO Rabbit/Mouse Chengdu China) at room temperature for 30 min, and washed with TBS. Finally, the DAB was used to illuminate the positive staining signals, and the sections were counterstained with hematoxylin. The positive staining signals were sorted into 4 grades, according to the standard procedures [3].

FISH

studied, of which 487 samples were obtained from Department of Pathology, the Affiliated Hospital of Southwest Medical University from November 2010 to July 2015, and 102 samples were provided by the Department of Pathology, People's Hospital of Deyan, Sichuan, China, from October 2010 to March 2013. Patients came from southwest China including parts area of Sichuan, Yunnan, Guizhou and Chongqing province, and the samples had IHC markers (ER, PR and Ki67) and were confirmed by clinicopathological information including age, tumor size, lymph node metastasis, and histological grade. The tissues were sectioned (3 µm thickness) for both FISH and IHC. This experimental protocol was pre-approved by the Medical Ethics Committee of Southwest Medical University (No. 20130051).

Immunohistochemistry

Her-2/neu protein expression was detected by using EnVision technology. Briefly, the paraffin embedded tissue were sectioned with 3 μ m thick and placed on poly-L-Lysine coated slides, every case was both with positive and negative control. The slides were dried overnight at 60°C. After deparaffinization and blocking of endogenous peroxidase, sections were transferred in retrieval solution (Tris-EDTA pH: 9.0) for 3 min, followed by washing with tap water.

FISH was performed using the HER-2/neu probes kit (Jingpujia Medical Technology Co, Ltd, Beijing, China). Firstly, 3 µm thick sections were mounted on poly-L-Lysine coated slides and baked overnight at 60°C. After deparaffinizing in xylene for 10 min two times and dehydrated in 100%, 85% and 70% ethanol for 2 min each and air-dry. Secondly, immersed the slides in distilled water in Pressure-cooker (121°C) for 3 min and for 4 min after disconnect the power supply, after washed in sodium saline citrate (2×SSC, pH 7.0) for 5 min twice at 37°C, the side were then digest for 18 min in protease solution (200 µg/ml) at 37°C, and then washed in 2×SSC for 5 min twice again. After that, 10 µl hybridization buffer containing the probe was applied to the target tissue on the slides. The sections were covered with a coverslip, and then the coverslip was sealed using rubber cement. The hybridization was conducted in a hybridization oven (ThermoBrite Statspin) under the following conditions: denature at 83°C for 8 min, incubation at 42°C in a humidified chamber for 16 hrs. Finally, slides were washed in 2×SSC for 10 min at 46°C, followed by 0.1% NP40 at 46°C for 5 min, and rinsed in 70% ethanol. After air-dry, 10 µl DAPI was applied and coverslip was gently placed. The hybridization signals were analyzed using a Nikon 80i fluorescence microscope (Nikon, Tokyo, Japan). The results were re-analyzed

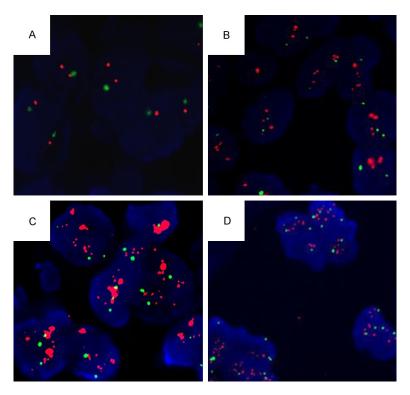


Figure 2. FISH results (CEP17 stained green and HER-2 gene stained red). A. Negative, a ration of HER-2 to CEP17 signals less than 2.0 and with an average HER2 copy number < 4.0 signals/cell. B. Equivocal: a ratio of HER-2 to CEP17 signals < 2.0 with an average HER2 copy number \geq 4.0 and < 6.0 signals/cell. C. HER-2 gene amplification: red signal were clustered obvious. D. HER-2 gene amplification: a ratio of HER-2 to CEP17 signals < 2.0 but the average HER2 copy number \geq 6.0 signals/cell.

Table 1. Correlation between IHC and FISH in HER-2 status

IHC Score		FISH Status	Total	Concordance		
	Negative	Equivocal	Positive	TOLAT	Concordance	
-/+	148	1	9	158	93.67% (148/158)	
++	198	12	100	310	32.26% (100/310)	
+++	42	5	74	121	61.16% (74/121)	
Total	388	18	183	589		

IHC, immunohistochemistry; FISH, fluorescence in situ hybridization. Kappa = 0.189, P = 0.000; r = 0.427, P = 0.000.

based on the methods of the latest ASCO-CAP HER2 Test Guideline Recommendations (Wolff AC et al., 2013), e.g. if HER-2 signal (red) were clustered obvious, it was determined that clustered amplification, otherwise, 30 invasive tumor nuclei ratio of HER-2 to CEP17 of \geq 2.0 or < 2.0 but the average HER2 copy number \geq 6.0 signals/cell was considered as HER2 amplification whereas \leq 2.0 and with an average HER2 copy number < 4.0 signals/cell was reported as non-amplification, while when HER2/CEP17

ratio < 2.0 with an average HER2 copy number ≥ 4.0 and < 6.0 signals/cell was taken as an equivocal result that required additional 70 nuclei or repeat of the test.

Statistical evaluation

Statistical analysis was carried out using SPSS version 13.0. IHC and FISH results were done using "Kappa" as measure of concordance. Groups were compared using Wilcoxon test, and spearman rank correlation analysis and chi-square test. *P* < 0.05 was considered as statistically significant.

Results

In the present study, 589 specimens of invasive breast cancer paraffin sections of southwest China women were detected. The IHC slide was scanned for immunostaining evaluation scored IHC 0, 1+, 2+, 3+ (Figure 1A-D). Of the 589 samples detected by FISH, 65.87% (388/589) showed HER-2 negative (Figure 2A), 3.06% (18/589) were detected report equivocal (Figure 2B), and the rate of HER-2 gene amplification identified was 31.07% (183/ 589; Figure 2C, 2D). Concordance was detected in 93.67% with IHC 0/+, 32.26% in IHC 2+, 61.16% overall in

IHC 3+. The concordance of 54.67% was observed with a Kappa coefficient of 0.189 (P = 0.000) and a positive correlation was found between IHC score and FISH results (r = 0.427 P = 0.000) (**Table 1**).

The relationship between FISH and IHC, and the relevance to cliniopathological characteristics, was also investigated in this study. There were significant differences between the expression of ER, PR, ER/PR status and FISH

HER-2 and its significance in breast cancer

Table 2. Association of HER-2/neu protein expression, HER-2 gene status and clinical-pathological characteristics

	FISH results			P value	P value	IHC Score			P value	P value
	Negative	Equivocal	Positive	,	(Correlation test)	-/+	++	+++	(Ranksum test)	(Correlation test)
Age				P > 0.05	P > 0.05				P > 0.05	P > 0.05
≤ 45	200	6	81			82	151	54		
> 45	188	12	102			76	159	67		
ER				P = 0.000	P = 0.000				P = 0.000	P = 0.000
-	84	3	108			39	110	46		
+	81	7	51			27	78	34		
++	107	3	15			54	47	24		
+++	116	5	9			38	75	17		
PR				P = 0.000	P = 0.000				P = 0.035	P > 0.05
-	133	5	104			56	141	45		
+	112	7	49			64	69	35		
++	86	3	18			29	52	26		
+++	57	3	12			9	48	15		
ER/PR				P = 0.000	P = 0.000				P > 0.05	P > 0.05
-/-	71	4	74			36	82	31		
+/-	56	5	32			20	66	7		
-/+	29	6	11			16	24	6		
+/+	232	3	66			86	138	77		
Ki67				P > 0.05	<i>P</i> > 0.05				P > 0.05	P > 0.05
≤ 50	259	11	118			107	199	82		
> 50	129	7	65			51	111	39		
Tumor size				P > 0.05	P > 0.05				P > 0.05	P = 0.000
< 2	101	4	50			52	70	33		
≥ 2, < 5	182	9	85			91	158	27		
≥ 5	105	5	48			15	82	61		
lymph node				P > 0.05	P > 0.05				P > 0.05	P > 0.05
0	145	3	57			53	107	45		
≥ 1, < 4	114	10	58			42	106	34		
≥ 4	129	5	68			63	97	42		
Histological grade				P > 0.05	P > 0.05				P = 0.022	P = 0.003
1	109	6	42			51	87	19		
II	151	9	87			65	127	55		
III	128	3	54			42	96	47		

ER, estrogen receptor; PR, progesterone receptor.

results ($x^2 = 108.171$, P = 0.000; $x^2 = 30.914$, P < 0.001; $x^2 = 39.845$, P < 0.001), Expression of ER, PR also significant difference to HER-2/neu protein expression ($x^2 = 19.430$, P < 0.001; $x^2 = 8.609$, P < 0.05). The expression of ER was negatively correlated with HER-2 gene amplification and the expression of Her-2/neu protein (r = -0.419, P = 0.000; r = -0.144, P = 0.000). The PR expression and ER/PR status was negatively correlated with HER-2 gene amplification (r = -0.226, P = 0.000; r = -0.258, P = 0.000), but not significantly with Her-2/neu protein

expression (P > 0.05), otherwise the tumor size and histological grade were positive correlation with HER-2/neu protein expression, but not with HER-2 amplification (**Table 2**).

Discussion

As the accuracy and importance of HER2 testing and its status evaluation, The ASCO/CAP (American Society of Clinical Oncology and the College of American Pathologists) have continuous updated the recommendations in HER-2

testing not only technical guidelines but also interpretation guidelines. We use the recently updated guidelines, the new ASCO/CAP 2013 criteria, in our experiment. The new guideline in order to avoid false negative results contrast to the former one, and the first time combined the HER-2/CEP17 ratio and the HER-2 gene copy number to define the negative and equivocal status. We analyzed the concordance between FISH and IHC for HER-2 status, and re-evaluated its clinicopathologic significance to got results as following using the new guidelines make sure inferred more reliable evaluation results.

Her-2 gene amplification and its receptor protein over expression are known to be associated with poor prognosis. It is significant that correct evaluation of HER-2 status of breast cancer patients to their treatment and prognosis [4]. Generally, IHC assay is used traditionally and relatively cheap and convenient to operate. When IHC 2+, indicating there is nondeterminacy in the Her-2/neu protein expression, at this point, FISH, considering as a gold standard, is a technology of evaluation HER-2 gene amplification at the levels of DNA [1, 5]. Thus, both IHC and FISH are widely used to detect HER-2 status, the concordance between the two methods is extensively investigated, we used the most recent guidelines re-evaluated its concordance there to study further. Recent studies reported that strong correlation between IHC 0/1+ and FISH [1, 5, 6]. In our present study, the concordance between FISH and IHC 0/1+ was 93.67%, which is consistency with the researches, and increased compared with our previous reported 87.5% [7], Otherwise, 388 cases were detected negative in 589 samples which compared 88 cases negative in the 159 patients studied in the previous study, the ratio has increased than old criteria, not exactly the same as Stoss et al. [8] reported. Because the analyzed results do not came from same batch of data and also related to the small cases in previous research. Then we reported the concordance between FISH and IHC 2+ was 32.26%. It is consistency with Sudha SM et al. reported about 16-29% [9, 10], and lower than 50.50% we calculated before [7]. It seems the most recent ASCO/CAP guidelines, which combine the HER2/CEP17 ratio and the mean HER2 gene copy number to define the equivocal status, can effectively reduce the redetection [8].

But showed IHC 2+ should be routinely performed FISH test because its lower concordance rate, in order to accurately make a clinical treatment plan for the patients. Otherwise, several reporters have published their researches to show high concordance rate about between FISH and IHC 3+ [1, 3, 6, 11, 12], by contrast, our study illustrated that numeric only 61.16%, just similar as the previous reported 58.8% [7]. The reasons might be attribute to false positive results of IHC, such as specimens fixation, method of antigen retrieval, differentt antibodies and subjective interpretations, as well as impact of chromosome 17. Overall, our study demonstrated consistency between two technologies by consistency test (Kappa coefficient of 0.189) and a positive correlation was found between FISH and IHC assay, which is consistent with Kovacs' conclusion, but the consistency was not as the above, it's not difficult to see the difference is mainly in IHC 3+ which group still have a great percentage of HER-2 gene without amplification, but patients who with IHC 3+ can directly use targeted drug therapy in currently therapy guideline in the world. Therefore, IHC 3+ may another detected HER-2 gene status by FISH and HER-2 gene amplification patients taking targeted therapy have better prognosis just as the research of Mass RD et al. [13]. A lot of researches were also supported the FISH test results to be the basic standards to determine whether taking the targeted drug [14]. In summary, we recommend that the breast cancer patients should test the HER-2 gene by FISH whatever the IHC assay results, and thus to improve the diagnosis, guide treatment and determine the prognosis.

In the published literatures, there is an inverse association between HER-2 gene amplification and ER expression [15-17], in comparison, our research also indicated the similar results. Furthermore, ER also showed a converse relationship with HER-2/neu protein expression, such as Mirtavoos-Mahyari et al. [18] reported that ER-negative breast cancer tumors were significantly more likely to be HER-2 positive (3+ by immunohistochemistry or positive by fluorescent in situ hybridization) than were ER-positive tumors. In addition, Huang HJ et al. [5] reported that there is an inverse association between PR and HER-2/neu in patients older than 45-year-old, and Han Zhang et al. [10]

reported a significant difference of HER-2 gene amplification between ER+PR+ and ER-PR-. In our present results, PR expression and ER/PR status was negatively correlated with HER-2 gene amplification. Meanwhile, ER/PR status, such as ER-PR- or ER+PR-, seemed to have worse biologic characteristics and poor prognosis [19]. The reason could be the complex signaling pathways between ER, PR, HER-2/neu in breast cancer cells. Evidence certified that the expression of ER, PR, HER-2 in breast cancer may classify the patients into subtypes, which closely related to clinic [20, 21].

It is an established observation that HER-2/neu protein positive status is associated with growing tumor size [22, 23] and tumor grade [24]. It similarly showed a positive correlation with HER-2/neu protein expression in our present study. And other clinicopathologic features like Ki67, age and lymph node did not show significant correlation with HER-2 gene amplification and HER-2/neu protein expression, as Ning SF et al. reported [6]. But Mujtaba et al. [23] once reported HER-2/neu over-expression increased with increasing lymph node metastasis, this might be due to different date statistic.

In conclusion, Breast cancer patients, considering clinical therapy, should be detected ER, PR and HER-2/neu expression routine by IHC, but as the basis of targeted therapy, FISH strongly recommends a standard method to evaluation HER-2 gene amplification. The new 2013 guidelines improved consistency between IHC and FISH test, and calculating HER-2/CEP17 ratio combined with the HER-2 gene copy number show a more accuracy when evaluating negative and suspicious status.

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Disclosure of conflict of interest

None.

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