

Prognostic Importance of p53 and *c-erbB-2* Oncoproteins Overexpression in Patients with Breast Cancer

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Abstract

Using immunohistochemistry, 119 breast cancer tissues were examined for overexpression of p53 and *c-erbB-2* oncogene proteins. In 46 (38.7%) of the cases p53 was overexpressed, while 35 (29.4%) demonstrated positive *c-erbB-2* immunostaining. Expression of these two oncogene products was closely correlated ($p < 0.01$). There was no significant association between p53 protein expression and age of the patients, clinical stage, tumor size, number of involved nodes or estrogen receptor status. However, we found significant correlation between p53 protein expression and 5-year disease-free survival ($p = 0.0113$). In addition, the findings in this study clearly indicated that the co-overexpression of p53 and *c-erbB-2* proteins was a powerful predictor for early recurrence in the patients with breast cancer.

The 20-kilobase p53 gene (TP53) is present on human chromosome 17p and encodes a 53-kilodalton (Kd) nuclear phosphoprotein⁽¹⁻³⁾. Mutations in the p53 gene have been identified in a variety of human malignancies⁽³⁻⁶⁾, including breast cancer^(3,6). Most mutations in the p53 gene are commonly associated with overexpression of mutated p53 protein which can be immunohistochemically detected in the nuclei of cancer cells⁽⁷⁻⁹⁾. Overexpression of this protein has also been suggested to be of prognostic value in breast cancer patients^(10,11).

The *c-erbB-2* (HER-2, *neu*) gene is carried on chromosome 17q and encodes a 185 Kd glycoprotein that is closely related to the epidermal growth factor receptor protein⁽¹²⁻¹⁴⁾. Both gene amplification and protein overexpression analyses have shown that elevated *c-erbB-2* is a poor prognostic indicator in patients with breast cancer⁽¹⁵⁻¹⁷⁾. In our previous study we also found that overexpression of the *c-erbB-2* protein detected by immunohistochemistry was an effective predictor for disease recurrence in the patients⁽¹⁸⁾. In this article, data of p53 expression correlating with cli-

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nical parameters and five-year disease-free survival were additive. Moreover, we also examined the effects of the two oncogene products coexpression on five-year disease-free survival which were included in the present study.

MATERIAL AND METHOD

Patients

This study included 119 primary tumors from women with breast cancer treated at the National Cancer Institute, Bangkok between 1987 and 1989. None of the patients had distant metastasis at the time of operation. All node-positive patients received six cycles of adjuvant chemotherapy containing cyclophosphamide, methotrexate and fluorouracil and local radiation (if the primary tumor was T3 and the patients had inadequate lymph node dissection). For the node-negative patients treatment varied according to T lesion, hormone receptor and age. After surgery, tissue samples were kept frozen at -70°C until use for biochemical determination of estrogen receptor (ER)(19). Receptor concentration less than 10 fmol/mg was considered negative. A parallel sample was processed using routine techniques for histological examination and immunohistochemical study on paraffin sections. The mean patient follow-up period was more than five years.

Immunohistochemical staining

The method for detection of *c-erbB-2* was performed as previously described(18). To detect p53 in formalin-fixed, paraffin-embedded tissues, an antigen retrieval using microwave technique was employed. Four-micrometer thick paraffin embedded sections were deparaffinized in xylene, rehydrated through alcohol and then treated with 3 per cent hydrogen peroxide in deionized water for 5 min to block endogenous peroxidase activity. The sections were washed with phosphate-buffered saline (PBS) and placed in a plastic coplin jar containing distilled water. The jar was heated in a microwave oven (800 w) at the medium power setting for three 4-min cycles with an interval of one min between cycles to check on the water level in the jar. After heating, the coplin jar was removed from the oven and allowed to cool for 15 min. The slides were rinsed in PBS and preincubated with 1 per cent normal horse serum in PBS for 30 min. The sections were then incubated for 1 hour at room temperature with mouse monoclonal

antibody against p53 (DO7, Novocastra Lab., New Castle, UK) at a dilution of 1:200 in PBS. After rinsing in PBS, the sections were incubated for 30 min at room temperature with a biotinylated anti-mouse immunoglobulin (Vector Lab., Burlingame, CA) at a dilution of 1:1000, then rinsed again with PBS. Antibody binding was visualized by incubation with avidin-biotin peroxidase complex (Vectastain ABC kit, Vector Lab., Burlingame, CA) for 30 min at room temperature. The sections were rinsed in PBS and immersed in Tris-buffered saline with a pH of 7.4 containing 3, 3'-diaminobenzidine and 0.5 per cent hydrogen peroxide for 5 min and counterstained with Mayer's hematoxylin for 5 min. Finally, they were rinsed in tap water, dehydrated in ethanol, cleared in xylene and mounted in permount.

Staining with 1 per cent normal horse serum as primary antibody was used as a negative control. As positive controls, we included in each run sections from a formalin-fixed, paraffin-embedded breast cancer which had been previously shown to contain mutated p53 protein. Obvious nuclear staining in any number of malignant cells was considered positive.

Statistical analysis

The correlation between the immunostaining of p53 oncoprotein and other clinical parameters including *c-erbB-2* expression was evaluated by χ^2 test. Five-year disease-free survival (DFS) curves were performed by the Kaplan-Meier method(20) and the difference between the curves was assessed using the log rank test(21).

RESULTS

Of the 119 breast carcinomas included in this study, 46 (38.7%) were positive for p53 and 35 (29.4%) were positive for *c-erbB-2*. The relationship between staining of p53 oncoprotein and clinicopathologic features including *c-erbB-2* protein expression is shown in Table 1. Positivity of p53 protein was significantly associated with *c-erbB-2* protein overexpression ($P < 0.01$). No significant association was observed between expression of p53 protein and age, stage, tumor size, number of positive axillary nodes or ER status.

There was a significant difference in the 5-year disease-free survival curves between the patients with p53-positive tumors and the other cases with p53-negative tumors (Fig.1) ($p = 0.0113$).

Table 1. Expression of p53 protein in relation to clinicopathologic parameters.

Parameters	Number of cases Staining of p53 protein		Total	P
	Negative	Positive		
Age at diagnosis (yrs)				
<40	23 (69.7)	10 (30.3)	33	NS
40-60	45 (59.2)	31 (40.8)	76	
>60	5 (50.0)	5 (50.0)	10	
Stage				
I	18 (78.3)	5 (21.7)	23	NS
II	36 (62.1)	22 (37.9)	58	
III	11 (52.4)	10 (47.6)	21	
IV	0 (0)	3 (100.0)	3	
Unknown	8	6	14	
Tumor size (cm)				
≤ 3	61 (66.3)	31 (33.7)	92	NS
> 3	10 (47.6)	11 (52.4)	21	
Unknown	2	4	6	
No. positive axillary nodes				
0	40 (70.2)	17 (29.8)	57	NS
1-3	17 (63.0)	10 (37.0)	27	
>3	16 (45.7)	19 (54.3)	35	
ER status				
-	37 (56.1)	29 (43.9)	66	NS
+	36 (67.9)	17 (32.1)	53	
<i>c-erbB-2</i>				
-	58 (69.0)	26 (31.0)	84	<0.01
+	15 (42.9)	20 (57.1)	35	

Five-year disease-free survival curves were also plotted for the combination of p53 and *c-erbB-2* oncoproteins with four nonoverlapping categories: negative for both, p53-positive; *c-erbB-2* negative, p53 negative; *c-erbB-2* positive, and doubly positive as shown in Fig. 2. Patients with tumors overexpression either p53 or *c-erbB-2* separately did worse than patients lacking both oncoproteins. Patients with tumors that were both p53 and *c-erbB-2* positive had the worst prognosis ($p=0.0001$).

DISCUSSION

Recently, several investigators have agreed that immunohistochemical methods used to detect expression of the p53 and *c-erbB-2* gene products in paraffin-embedded archival samples is reliable in indicating p53 missense mutations and *c-erbB-2*

gene amplification(22-25). The antibodies to p53 and *c-erbB-2* proteins used for immunostaining in this study were clearly demonstrated to provide complete agreement between immunohistochemistry and molecular biology assays(26,27).

A wide range of experimental data has been published on p53 expression in breast cancer. Using different monoclonal antibodies and staining conditions, p53 overexpression has been detected in 15 per cent to 50 per cent of breast cancers(28). In the present study, overexpression of p53 was found in 39 per cent of the tumors using the DO7 antibody on paraffin-embedded sections, being similar to the findings of a study by Jacquemier *et al*(29) who used DO1 antibody on paraffin-embedded sections.

The prognostic value of p53 expression in relation to other prognostic parameters in breast

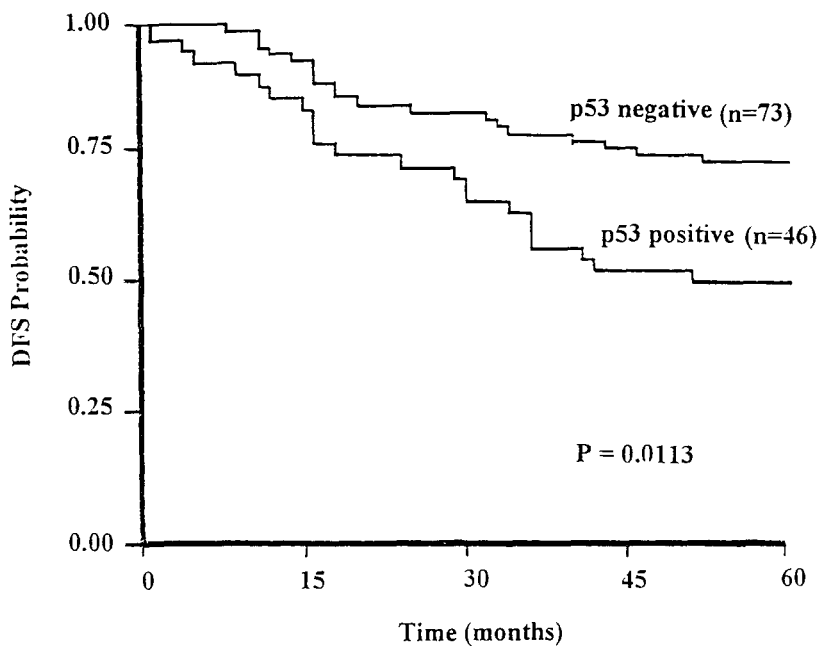


Fig. 1. Five-year disease-free survival curves for patients with breast cancer subdivided by p53 immunostaining.

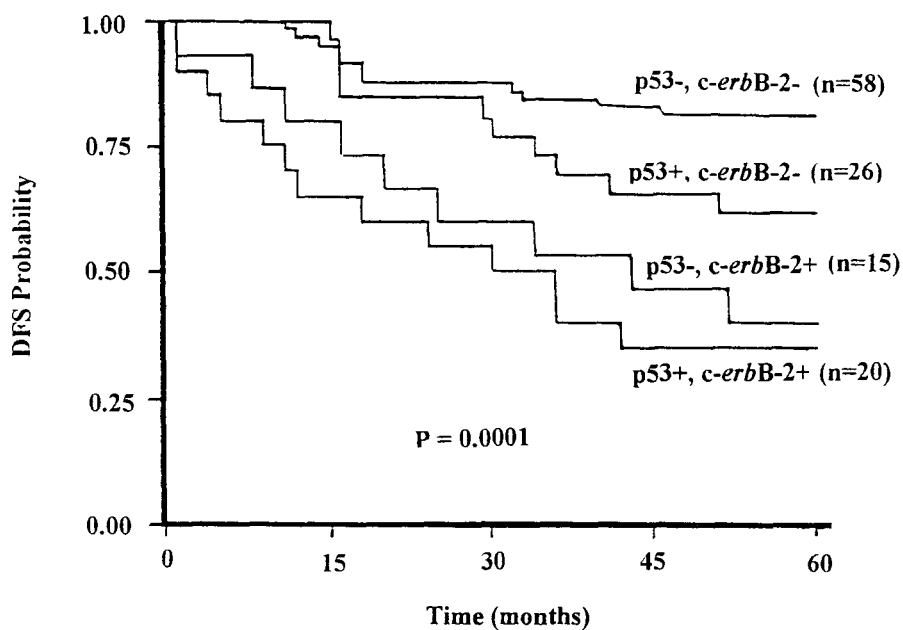


Fig. 2. Five-year disease-free survival curves for patients with breast cancer separated by staining for both p53 and *c-erbB-2*.

cancer is still indefinite. Some reports have suggested that p53 oncoprotein accumulation is associated with clinical stage(29,30,32), tumor size (31-33), nodal status(31,33), estrogen receptor(30, 32-35), and *c-erbB-2* expression(29,33,35). However, others indicate no correlation with age of the patients(31,32,34), clinical stage(36), tumor size (31,34), nodal status(32,34), estrogen receptor(37), or *c-erbB-2* expression(28,30,32). In the current study, we found correlation only with *c-erbB-2* expression.

Thor *et al*(38) examined 253 patients with breast cancer of all stages and found that p53 immunostaining in paraffin-embedded samples was a prognostic factor for disease-free survival in both the node-negative and node-positive groups. Marks *et al*(32) found the significant prognostic effect on recurrence-free survival only in the node-negative subset. In our study, we did not examine the two subsets separately due to the small number of patients. However, our findings indicated that the overexpression of p53 oncoprotein was a prognostic predictor for 5-year disease-free survival in the patients, regardless of nodal status.

Since the study of p53 and *c-erbB-2* oncoproteins can lead to improvements in the management of patients with breast cancer, either as a prognostic indicator or as a predictor of therapeutic responses(32-38), the combined study of these two oncogene products, therefore, could help to improve their clinical value. Our previous study

clearly demonstrated that overexpression of *c-erbB-2* protein was an effective prognostic predictor for 5-year disease-free survival in the patients(18). In the current study, we also found the same prognostic effect in the overexpression of p53 protein. Moreover, when combining p53 and *c-erbB-2* proteins expression, we observed that their coexpression had a more powerful effect on predicting 5-year disease-free survival than p53 protein expression alone. These results are consistent with the other studies(32,39). In addition, one previous study suggested that overexpression of both these oncogenes in a single tumor had an additive effect on the prognosis(32). From these findings and ours, it is suggested that double immunohistochemical detection of p53 and *c-erbB-2* proteins overexpression is likely to help clinicians identify the patients with early recurrence more accurately and arrive at more rational treatment decisions.

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การใช้ p53 และ c-erbB-2 oncoproteins ช่วยพยากรณ์โรคในผู้ป่วยมะเร็งเต้านม

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ผู้วิจัยได้ใช้วิธี immunohistochemistry ในการตรวจหา p53 และ c-erbB-2 proteins ในชิ้นเนื้อมะเร็งเต้านมของผู้ป่วยจำนวน 119 ราย พบ 46 ราย (ร้อยละ 38.7) มี p53 protein และ 35 ราย (ร้อยละ 29.4) มี c-erbB-2 protein การตรวจพบผลผลิตของยีนทั้ง 2 ชนิดนี้มีความสัมพันธ์กันอย่างมีนัยสำคัญทางสถิติ ($p < 0.01$) การศึกษานี้ไม่พบความสัมพันธ์ระหว่าง p53 protein กับอายุของผู้ป่วย, clinical stage, tumor size, number of involved nodes และ estrogen receptor status แต่พบว่า p53 protein มีความสัมพันธ์กับอัตราการรอดโรคในระยะ 5 ปี ($p = 0.0113$) นอกจากนั้นผลจากการศึกษานี้ยังแสดงให้เห็นว่าการตรวจหาทั้ง p53 และ c-erbB-2 proteins จะเป็นตัวช่วยบอกการกลับคืนของโรคในผู้ป่วยมะเร็งเต้านมได้ดี

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