Neurotensin Receptor 1 (NTSR1) Overexpression in Breast Carcinomas Is Common and Independent of ER/PR/Her2 Expression

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ABSTRACT

Neurotensin (NT) is a 13-amino acid peptide with trophic effects on some neoplasms. Its bioactivities are mainly mediated by neurotensin receptor 1 (NTSR1). Both NT and NTSR1 were found to be upregulated in breast cancer. NT/NTSR1 thus becomes a potential therapeutic target. We studied whether any correlation exists between the expression of NTSR1 in breast carcinomas and the expression of ER, PR, and Her2. A total 85 cases of invasive ductal (62) and lobular (23) breast carcinomas were studied. Based on their ER/PR profiles, the ductal carcinomas (DCs) were subcategorized into ER+/PR+ (21), ER+/PR− (20), and ER−/PR− (21). All of the lobular carcinomas (LCs) were ER+/PR+. 21.57% of all DCs and 5.56% of LCs were Her2 positive. 77.78% of ER−/PR− DCs were also Her2 negative (triple negative). The expression of NTSR1 was detected by immunohistochemistry and was semiquantitated (as negative, 1+, 2+, 3+). Both 2+ and 3+ were collectively defined as overexpression. The expression of NTSR1 was weak and focal in non-neoplastic mammary epithelial cells. It is increased in 74.19% of DCs (80.95% in ER+/PR+, 75% in ER+/PR−, and 66.67% in ER−/PR− group), and in 95.65% of LCs. The overexpression of NTSR1 is similar between ER+ DCs and ER− DCs (75% vs 66.67%, p > 0.05) as well as between PR+ DCs and PR− DCs (80.95% in ER+/PR+ vs 75% in ER+/PR−, p > 0.05). And it was seen in 77.78% of Her2+ DCs, 78.38% of Her2− DCs, 94.12% of Her2− LCs, and 78.57% of triple negative DCs. Overall, NTSR1 is commonly overexpressed in both ductal and lobular breast carcinomas and is independent of the ER/PR/Her2 profiles of the tumors. The present data supports the potential benefit of developing NTSR1 blockers in the adjuvant therapy of breast carcinomas, particularly for those “triple negative” tumors.

Keywords: Neurotensin; Neurotensin Receptor; NTSR1; Breast Carcinoma; Estrogen Receptor; Progesterone Receptor; Her2

1. Introduction

Neurotensin (NT) is a 13-amino acid regulatory peptide that was initially isolated from bovine hypothalamus by Carraway and Leeman [1] and was subsequently found to express highest in the small intestine [2]. Its physiologic secretion is stimulated significantly by fat intake [3]. Over the years, NT was found to possess diverse functions including trophic effects on some normal and neoplastic cells. Most of NT’s effects are mediated via high-affinity neurotensin receptor 1 (NTSR1), a member of G-protein-coupled receptor family. The activation of NT/NTSR1 pathway involves several intracellular signaling events, including increased intracellular calcium release, activation of phosphatidylinositol 3'-kinase (PI3K), mitogen activating protein kinase (MAPK) family member ERK1/2, NF-κB pathway, and Rho family of small GTPases, leading to cell proliferation. The growth-promoting effect of NT/NTSR1 has been found in a variety of carcinomas including those of the colon, pancreas, prostate, and lung [4]. It has also been found in breast carcinomas that near all breast ductal carcinomas over express NTSR1 and about a third coexpress both NT (autocrine) and NTSR1 [5], NT exerted an inhibitory activity on the apoptosis in breast cancer cells [6], and the upregulated expression of NTSR1 is related to the tumor progression [7]. All of these features make NT/
NTSR1 an ideal novel supplementary therapeutic target in the hormonal therapy of breast cancer. Meanwhile, many efforts on development of NTSR1 antagonists (e.g., SR48692, SR142948) have been undertaken [8,9]. Additionally, NT/NTSR1 activation was found to stimulate the expression of epidermal growth factor receptor (EGFR) as well as to transactivate EGFR [10-12]. This is very interesting in view of that a member of the EGFR family, Her2 (cErbB2) is currently the primary hormonal therapeutic target for breast cancer. Moreover, a cross talk between NT and estrogen and between NT and progesterone has been noted, at least in the physiologic regulation of central nervous system [13-16]. In order to better evaluate the possible application and benefits of the anti-NTSR1 treatment, further understanding on whether the NTSR1 expression correlates with the expression of ER/PR/Her2, as well as the histologic type and grade of breast cancer will be important. Little data regarding these questions is currently available. In the present study we have found that NTSR1 overexpression is common in breast carcinomas and it is independent of the expression of ER, PR, or Her2.

2. Methods and Materials

2.1. Breast Carcinoma Patients and Samples

Eighty five cases of invasive breast carcinomas (age 24 - 83, mean 59.98 years old), including both ductal (62) and lobular (23) types, resected by mastectomy or segmental resection were retrieved from the 2010 surgical pathology file at Calgary Laboratory Services. All tumors were graded histologically using the modified Bloom-Richardson (MBR) system.

The study was approved by the Research Committee of Calgary Laboratory Services. The ethics approval was granted by Calgary Health Research Ethics Board.

2.2. Detection of NTSR1, ER, PR, and Her2

One of the most representative sections of tumor was selected from each case for the study. The immunoreactivity of NTSR1 was detected by immunohistochemistry carried out on deparaffinized sections using the avidin-biotin-peroxidase complex method. The formalin fixed paraffin embedded tissue sections were pretreated in CINTec Epitope Retrieval Solution (10 mM Tris/1mM EDTA, pH 9.0) for 20 minutes at 95°C. This is important to retrieve the epitope of NTSR1 protein. The pretreated sections were then incubated with an appropriate dilution of the primary rabbit polyclonal antibody (IMGENEX, California, USA) against the third cytoplasmic domain of human NTSR1. All slides were stained with Ventana Nexes IHC auto-stainer at 1:40 dilution using UltraView Universal DAB Detection (Ventana #760 - 500). The expression of NTSR1 was semiquantitated based on the positivity and intensity of the immunostain as negative (no stain at all), 1+ (faint or less than 10% of tumor cells stained), 2+ (moderate intensity or up to 50% of tumor cells stained), and 3+ (strong intensity and more than 50% of tumor cells stained). A comparison of NTSR1 expressions between these groups was analyzed. Both 2+ and 3+ were collectively defined as increased expression (overexpression) in our results, considering that 1+ was observed in non-neoplastic ductal epithelial cells.

ER and PR expression were routinely assessed by immunohistochemistry (performed at the diagnostic immunohistochemistry laboratory of Calgary Laboratory Services), and Her2 was routinely detected by silver in-situ hybridization (SISH) (performed at the diagnostic laboratory of the Tom Baker Cancer Centre). The methodology of all of the lab tests and the result analysis were validated. The results were reported routinely and included in the clinical pathology reports.

2.3. Statistical Analysis

The rate (%) of NTSR1 overexpression in tumors of different groups was calculated and compared between groups by Fisher Exact test.

3. Results

The details of the patients’ information are summarized in Table 1.

Based on their ER/PR profiles, the ductal carcinomas (DCs) were subcategorized into ER+/PR+ (21 cases), ER+/PR− (20 cases), and ER−/PR− (21 cases). The 23 lobular carcinomas (LCs) were all ER+/PR+. The data of Her2 were available in 69 cases. 11 of 51 (21.57%) DCs and 1 of 18 LCs (5.56%) were Her2 positive. 14 of 18 (77.78%) ER−/PR− DCs were also Her2 negative (triple negative).

The immunoreactivity of NTSR1 appeared in both membranous and cytoplasmic patterns, as shown in Figure 1. It was weak and partial in the non-neoplastic mammary epithelial cells, but it was stronger in both in-situ and invasive carcinomas.

As shown in Figure 2, the overexpression of NTSR1 was detected in 46/62 (74.19%) DCs (80.95% in ER+/PR+, 75% in ER+/PR−, and 66.67% in ER−/PR− group), and 22/23 (95.65%) LCs. The overexpression of NTSR1 appeared to be slightly elevated in tumors with the expression of both ER and PR, although no statistically significant difference existed (p > 0.05) between any of these groups based on our sample size.

As shown in Figure 3, the overexpression of NTSR1 was slightly more common in ER+ DCs than in ER− DCs (75% versus 66.67%, p > 0.05), and was similar between PR+ DCs and PR− DCs (80.95% in ER+/PR+ DCs versus
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Table 1. Summary of breast cancer patients and NTSR1 immunohistochemistry.

<table>
<thead>
<tr>
<th>Type</th>
<th>ER/PR</th>
<th>Number</th>
<th>Age (mean)</th>
<th>MBR Grade 1</th>
<th>MBR Grade 2</th>
<th>MBR Grade 3</th>
<th>Her2 (+/Total)</th>
<th>NTSR1</th>
</tr>
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<tr>
<td>Ductal</td>
<td>ER+/PR+</td>
<td>21</td>
<td>63.67</td>
<td>6</td>
<td>11</td>
<td>4</td>
<td>4/16</td>
<td>0</td>
</tr>
<tr>
<td></td>
<td>ER+/PR−</td>
<td>20</td>
<td>59.12</td>
<td>5</td>
<td>8</td>
<td>7</td>
<td>7/17</td>
<td>0</td>
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<tr>
<td></td>
<td>ER−/PR−</td>
<td>21</td>
<td>60.91</td>
<td>0</td>
<td>3</td>
<td>18</td>
<td>4/18</td>
<td>1</td>
</tr>
<tr>
<td>Lobular</td>
<td>ER+/PR+</td>
<td>23</td>
<td>60.18</td>
<td>2</td>
<td>19</td>
<td>2</td>
<td>1/18</td>
<td>0</td>
</tr>
</tbody>
</table>

Figure 1. The intensity and pattern of NTSR1 expression on immunohistochemistry. (a) Weak (1+) expression in normal ductal epithelial cells (×200); (b) Moderate expression (2+) in ductal carcinoma in situ (DCIS) (×200); (c) Relatively membranous pattern of immunostain (×400); (d) Relatively cytoplasmic pattern of immunostain (×400); (e) Strong expression (3+) in invasive ductal carcinoma (×200); (f) Strong expression (3+) in invasive lobular carcinoma (×200).

75% in ER+/PR− DCs, p > 0.05).

As shown in Figure 4, the overexpression of NTSR1 was seen in 7/9 (77.78%) Her2+ DCs, 29/37 (78.38%) Her2− DCs, and in 16/17 (94.12%) Her2− LCs. In addition, the NTSR1 overexpression was seen in 11/14 (78.57%) triple negative DCs (not shown in the figure). No statistically significant difference existed (p > 0.05) between any of these groups.

Between the tumors with different histologic grades, as shown in Figure 5, the expression of NTSR1 seemed to be negatively correlated with the histologic grade of the tumors. The difference between Grade 1 and Grade 3 was statistically significant (p = 0.032), although the difference between Grade 1 and Grade 2, or between grade 2 and grade 3, was not statistically significant.

4. Discussion

The detection of certain biologic markers in breast cancer has had significant clinical impact and become routine in the clinical pathologic assessment of breast carcinomas. These markers are related either to the aggressiveness, invasiveness, and the extent of tumor spread, or to the response of certain therapeutic agents. Thus they have both prognostic and predictive values. So far, the most useful biomarkers include estrogen receptor (ER), progesterone receptor (PR), Her2, and Ki67. Currently, the application of ER, PR, and Her2 in the selection and prediction of hormonal therapy in breast cancers is critical. The antagonists of ER (e.g., Tamoxifen) and Her2 (e.g., Trastuzumab) play important roles in the management of breast cancer patients. Several new markers have been explored in recent years, including p53, ARF, TBX2/3, Cyclin D1, Cyclin E, BRCA1, BRCA2, and VEGF [17]. The utility of these emerging novel markers are being investigated.

NT/NTSR1 is still a relatively little-known marker

In summary, the overexpression of NTSR1 is less common in high grade carcinomas than in low grade ones. There seems to be a trend of higher expression of NTSR1 in ER+/PR+ tumors than in ER−/PR− tumors. However, there is no statistically significant difference between ER+ and ER−, PR+ and PR−, and between Her2+ and Her2− tumors.
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Figure 3. NTSR1 expression (% of high expression) in breast carcinomas with regards to ER and PR expression. (a) Relationship of NTSR1 and ER expression; (b) Relationship of NTSR1 and PR expression. No significant difference existed between ER+ and ER− tumors or between PR+ and PR− tumors.

Figure 4. NTSR1 expression (% of high expression) in breast carcinomas with regards to Her2 expression. No significant difference existed between Her2+ and Her2− tumors. (a) Her2+ vs Her2− ductal carcinomas; (b) Her2+ vs Her2− lobular carcinoma.

with regards to breast cancer. Although the tumor growth-promoting effects of NT/NTSR1 on breast cancer cells and the overexpression of NTSR1 in breast cancer tissue have been reported for years, the prognostic and predictive values of detection of NTSR1 in breast cancer have not been studied. In a study reported by Dupouy et al., the overexpression of NTSR1 seemed to be associated with the histological grade, size of tumor, and number of positive axillary lymph nodes, and it was also found to be an independent prognostic factor associated with the poor outcome [6]. In our data, however, the NTSR1 overexpression appeared to be negatively correlated with the histologic grade of the tumors. Further study is required to clarify this discrepancy.

In the present study, NTSR1 was found to be universally expressed in both ductal and lobular breast carcinomas, and was independent of the ER/PR/Her2-neu profiles of the tumors. The study was not designed to further correlate the NTSR1 expression with the patient outcome and many other clinopathologic features. The data of the potentially prognostic value of NTSR1, unfortunately, is not available.

Our finding that NTSR1 is expressed in all kinds of breast carcinomas further supports the potential benefits of developing NTSR1 blockers in the adjuvant therapy of breast carcinomas, particularly for the “triple negative” tumors. It is reasonable to postulate that the anti-NT/NTSR1 approach would be a supplement and/or alternative to the established hormonal treatment of breast carcinomas. In addition, in view of the transactivation of NT/NTSR1 pathway on EGFR and the interaction between NT and estrogen/progesterone, the intervention of NT/NTSR pathway may also modulate the therapeutic effects of ER−/Her2 antagonists.

Several experimental studies using breast cancer cell
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lines have demonstrated the significant growth-inhibitory effects of various means of NTSR1 blockage. Earlier studies using SR48692, the first NTSR1 antagonist has shown a significant reduction of cancer cell proliferation in ex vivo cell culture or xenografted model [18,19]. In the study of Souaze et al., small interfering RNA knockdown of NTSR1 in xenografted MDA-MB-231 cells resulted in a 70% decrease in tumor growth as compared to wild-type cells [5]. In the study of Heakal et al, nanoliposomal short-chain ceramide, a chemical that can inhibit agonist-dependent translocation of NTSR1, was found to significantly inhibit NTSR1-mediated MDA-MB-231 breast cancer progression (mitogenesis, migration, and matrix metalloproteinase activity) [8]. Castillo-Rodriguez et al. recently developed a neurotensin-polyplex that transfects therapeutic genes into NTSR1-expressing breast cancer cells (MDA-MB-231 cells) and kills tumor cells in both in vitro and in vivo models [20]. Unfortunately, to the best knowledge of the authors, a clinically usable NTSR1 antagonist is still not available. The exact efficacy of the proposed anti-NTSR1 treatment awaits validation through future study, provided a clinically applicable drug becomes a reality in the near future with the recent better understanding of the chemical structure of NTSR1 [21].

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REFERENCES


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