

REVIEW ARTICLE

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The *neu*-protein and breast cancer

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Abstract The *neu*-protein is overexpressed in about 20% of invasive duct cell carcinomas of the breast. The only reliable sign for *neu*-overexpression by immunohistochemistry is membrane staining. Its overexpression is correlated with decreased overall survival and disease free survival due to increased metastatic activity of *neu*-overexpressing tumour cells. This increased metastatic potential is a consequence of the motility enhancing activity of the *neu*-protein, which is exclusively expressed on pseudopodia, and to a lesser extent of its growth stimulating effect. From a clinical point of view, the assessment of *neu*-overexpression in breast cancer might become a useful tool in the future treatment of patients by chemotherapy, since patients whose tumour shows *neu*-overexpression benefit from higher doses of chemotherapy. The molecule plays a key role in the pathogenesis of Paget's disease of the breast. A chemotactic factor which is secreted by epidermal keratinocytes attracts the Paget cells to spread into the epidermis and acts via the *neu*-protein. In ductal carcinoma in situ, the combination of *neu*-overexpression and large cell type is highly correlated with extent of disease and therefore *neu*-overexpression might be a predictive marker for recurrence of disease after tumour resection.

Key words *neu* · *c-erbB2* · Oncogene · Breast cancer · Paget's disease

Introduction

In recent years, the expression of the *neu*-protein has gained increasing importance for the biology and prognosis of breast cancer. This review focusses on the different biological and clinical aspects of the protein. From the biological point of view, its function as a motility factor receptor is stressed. Clinical applications of this motility function are found in the pathogenesis of Paget's

disease, in the predictive value of *neu*-overexpression in the assessment of the extent of disease in ductal carcinoma in situ (DCIS) and in the aggressive metastatic behaviour of *neu*-overexpressing tumours in the first years after diagnosis. The ongoing debate on the correlation of *neu*-overexpression with prognosis and the search for a specific ligand are discussed.

The *neu*-oncogene: structure and function

The *neu*-oncogene is a protooncogene which codes for a membranous protein: the *neu*-protein. The *neu*-protein belongs to the *erbB* protooncogene family or type I family of tyrosine kinase receptors. This *erbB* family of tyrosine kinase receptor molecules is constituted by epidermal growth factor receptor (EGF-R), *neu*, *c-erbB3*/HER3 and *c-erbB4*/HER4. Their molecular weight is determined as 170 kDa, 185 kDa, 160 kDa and 180 kDa, respectively [38, 55]. All of these proteins share considerable homology with EGF-R, the first detected member of this oncogene family [4, 13, 62, 63, 66, 84]. All these transmembrane proteins function by phosphorylation of tyrosine for instance after triggering the intracellular tyrosine kinase domain by a ligand-receptor interaction. This phosphorylation presumably leads to signal transduction by phosphorylating second messenger proteins. Phosphorylation of a particular oncoprotein, triggered by a specific binding with its ligand, may lead to cross-phosphorylation of a related oncoprotein. So, the *neu*-protein can be cross-phosphorylated after phosphorylation of EGF-R by its ligands, EGF or transforming growth factor- α (TGF- α), which are not specific ligands for the *neu*-protein [11].

The *neu*-protein is composed of a cell-external ligand binding domain, a transmembranous component, which keeps the molecule in the lipid bilayer of the cell membrane and a cell-internal domain with tyrosine-kinase activity, involved in signal transduction [13]. The *neu*-oncogene was first described in rat neuro/glioblastoma cells, induced by treatment with a carcinogen [62, 63].

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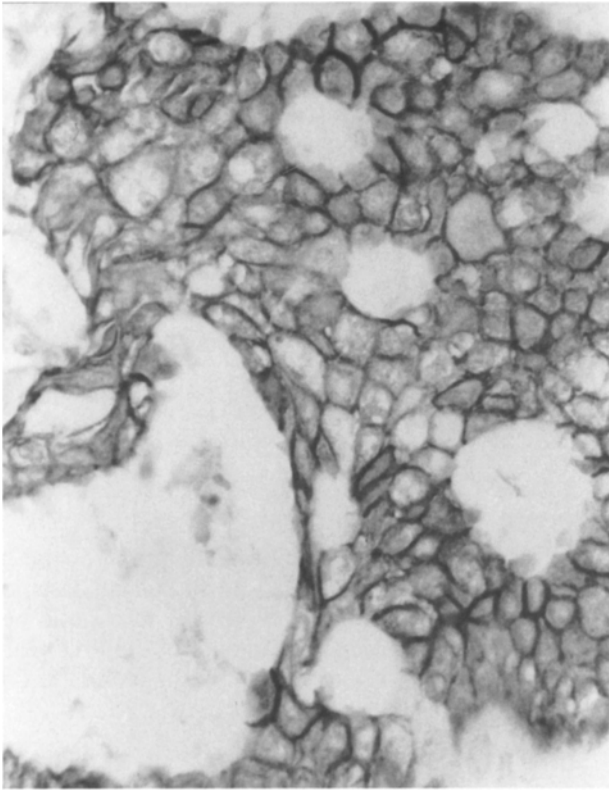


Fig. 1 Invasive ductal carcinoma immunostained for the *neu*-protein. All tumour cells show distinct expression at the plasma membrane. ($\times 440$)

The human equivalent of the rat *neu*-gene, which was independently cloned from a cDNA library was called HER-2 [13]. After cloning from genomic DNA it was designated as *c-erbB-2* and after cloning from a breast carcinoma cell line as an *erbB*-related gene [61, 66]. The gene is localized on the long arm of chromosome 17 [26]. So far, no viral oncogenic counterpart has been identified.

The oncogenic potential of the *neu*-protein was initially demonstrated after transfection experiments of the gene in NIH/3T3 cells. The gene lacked transforming activity after transfection despite expression of detectable levels of its protein, but a further five- to tenfold increase in its expression was associated with activation of the gene as a potent oncogene. These findings suggest that overexpression alone may convert the gene protein from a normal receptor into an oncogene protein [22].

Cellular localization of the *neu*-protein

Initially, overexpression of the *neu*-protein was assessed by Southern blot analysis of its gene. Overexpression of the protein can also be evaluated in a reliable way by immunohistochemistry and correlates with gene amplification [74]. Amplification of the gene has been reported in 10–40% of the breast carcinomas [74, 75]. Detected by immunohistochemistry, the *neu*-protein is overexpressed

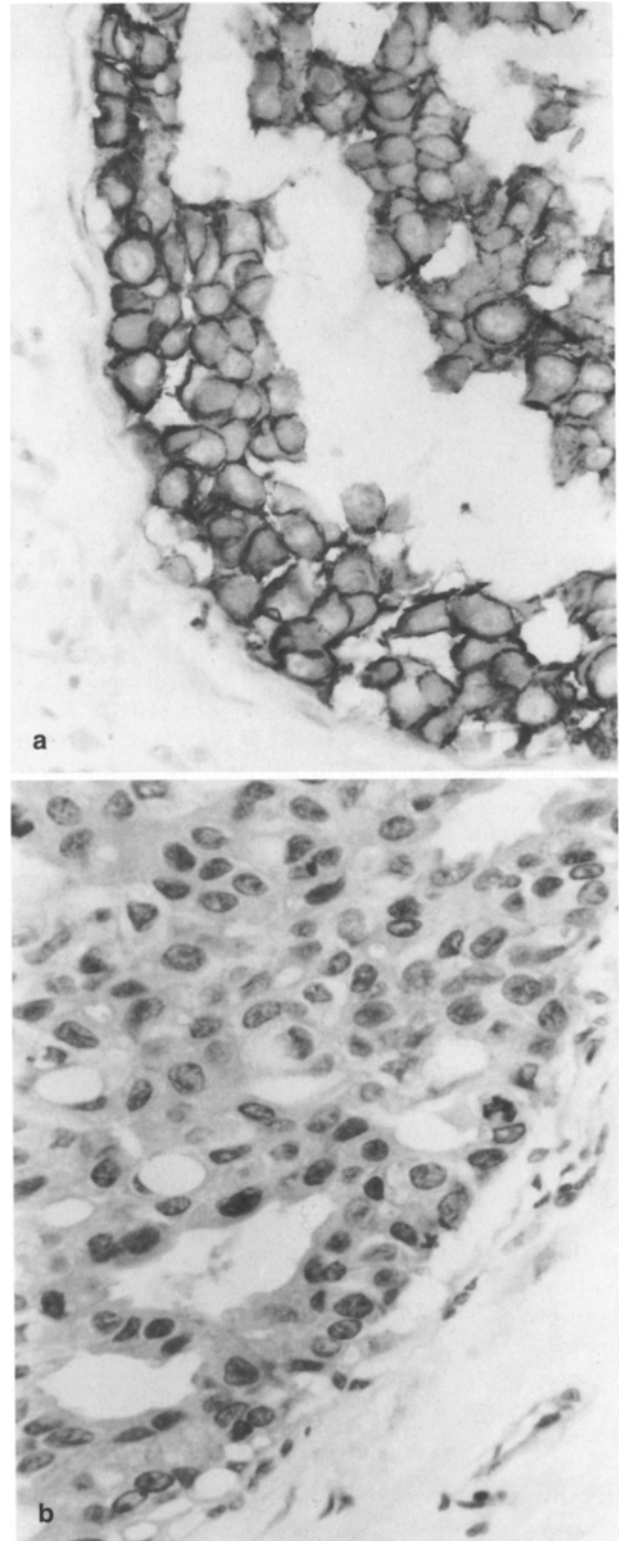


Fig. 2a Ductal carcinoma in situ immunostained for the *neu*-protein. All tumour cells show distinct expression at the plasma membrane. **b** These in situ tumours are typically composed of large cells with abundant cytoplasm. Usually the nuclei are pleomorphic. (Haematoxylin and eosin $\times 440$)

in about 20% of all breast cancers and in 25% of invasive duct cell carcinomas [17]. No overexpression has been described in the invasive lobular carcinoma nor in the in situ lobular carcinoma [31, 59].

In order to investigate overexpression by immunohistochemistry, it should be kept in mind that only membrane staining can be considered as a reliable sign of *neu*-overexpression (Figs. 1, 2). Several authors have observed a cytoplasmic reactivity by light microscopy, although the *neu*-protein is expected to be a plasma membrane protein. This cytoplasmic distribution was demonstrated in kidney cells, oral mucosa, urothelium [29], normal breast tissue [17, 77] and breast carcinoma cells [7]. A cytoplasmic localization of a membrane bound receptor could be explained by internalization of the receptor, but it is difficult to understand why a substantial immunoreactivity can be detected in cells with a barely detectable *neu*-mRNA level as was described in some fetal tissues [37]. The detection of a mitochondrial protein, different from the *neu*-protein but cross-reacting with antibodies directed against the intracellular domain of the *neu*-protein may lead to a false positive granular cytoplasmic staining, which has nothing to do with *neu*-overexpression [18, 68] (Fig. 3). The inconsistent interpretation of *neu*-positivity elucidates several discrepant findings on prognosis in the literature.

At the ultrastructural level, the distribution of the *neu*-protein is not diffuse, but resides on specific cell organelles. In the proximal tubule cells of the kidney, the normal expression was observed on the cell membrane of the microvilli of the brush border [18]. In *neu*-overexpressing breast carcinoma cells obtained from breast biopsy material, the protein was accumulated at plasma membrane protrusions [18]. Since plasma membrane protrusions such as microvilli, filopodia, lamellipodia and lobopodia, collectively designated as pseudopodia are obligatory for functions such as cell spreading, locomotion, cell motility and phagocytosis, it seems reasonable to suggest that the *neu*-protein is involved in these cellular functions.

Subsequently similar results on the distribution of the protein on the cell membrane were obtained from experimental data. In unstimulated *neu*-overexpressing SK-BR-3 human breast cancer cells, the *neu*-protein is similarly localized on small microvilli and plasma membrane protrusions throughout the cell surface as in breast biopsies and normal human glandular tissues. After stimulation of motility and chemotaxis, the small microvilli increase considerably in size. During this period, pseudopodia and long thin plasma membrane extensions are formed and aggregation of the *neu*-protein takes place on these enlarged membrane extensions [16].

The *neu*-protein and cell growth

From the earliest papers linking the *neu*-protein with poor prognosis and due to its similarity to EGF-R and its structural organization, it was postulated that the *neu*-

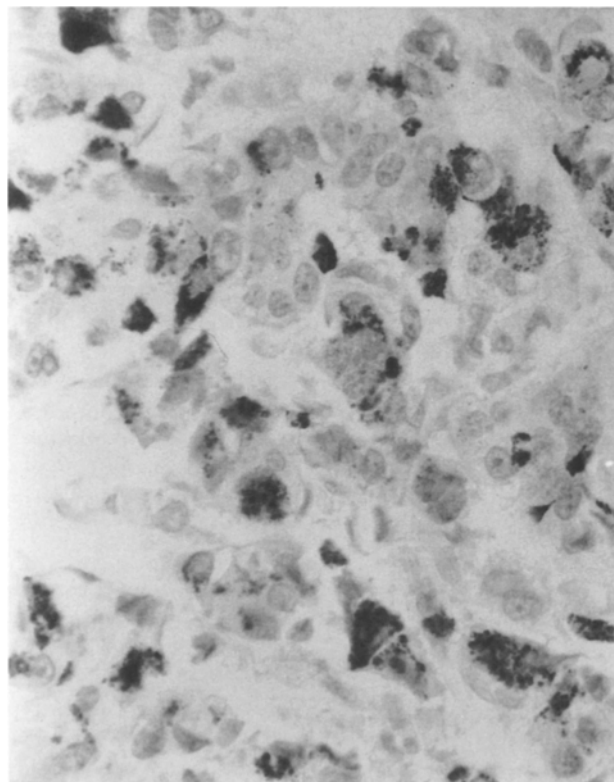


Fig. 3 Invasive lobular carcinoma, showing a granular cytoplasmic reaction after immunostaining for the *neu*-protein. This staining pattern should be considered as nonspecific and has nothing to do with *neu*-overexpression. It is frequently observed after immunohistochemistry with polyclonal and monoclonal antibodies directed against the intracellular domain of the *neu*-protein. ($\times 440$)

protein should be a receptor for a growth factor. The introduction of the *neu*-protein into the genome by transfecting 3T3 cells not only led to an altered genotype with a malignant phenotype, but also resulted in an increased growth potential [22]. The production of a hybrid protein, consisting of the EGF-R extracellular, transmembrane and protein kinase C substrate domains linked to the intracellular tyrosine kinase and carboxyl terminal domain of the rat *neu* cDNA, has evoked a similar result. Upon transfection with this construct and after stimulation of this membrane molecule by EGF and TGF α , tyrosine phosphorylation of the chimeric receptor protein and stimulation of DNA synthesis, resulting in mitogenic and transforming properties of the transfected cells were obtained [40]. Furthermore, anti-*neu* antibodies, regardless of isotype exerted a selective cytostatic effect on the growth of *neu*-transformed cells [23]. In vivo treatment with anti-*neu* antibodies was able to significantly inhibit the tumorigenic growth of *neu*-transformed NIH/3T3 cells implanted into nude mice. The antibody treatment was also able to inhibit the growth of the rat neuroblastoma cells from which the *neu*-oncogene was initially isolated [24]. However, experiments with several putative ligands have led to other conclusions. These putative ligands induced differentiation and led to complete cessation of cell growth (see below).

The *neu*-protein and cell motility

The precise morphological localization of the molecule on the cell membrane was helpful in elucidating its function in motility. The data obtained from experiments carried out with SK-BR-3 cells confirmed the initial morphological observations that the *neu*-protein exclusively resides on microvilli and pseudopodia [16]. The involvement of the *neu*-protein in cell motility is further supported by the detection of a 50 kDa *neu*-protein binding motility factor, which was purified from conditioned medium of malignant keratinocytes. This motility factor causes fast spreading, plasma membrane ruffling, cell differentiation, translocation, stimulation of chemotaxis and at high concentration growth arrest of SK-BR-3 cells. Motility and chemotaxis induced by the 50 kDa motility factor can be inhibited by antibodies directed against the extracellular domain of the *neu*-protein. The inhibition of the provoked motility again proves the function of the *neu*-protein in cell motility. The motility factor is considered as a *neu*-ligand, different from other, previously described putative ligands [15]. The association of the *neu*-protein with microfilaments via a large glycoprotein complex in mammary carcinoma microvilli is consistent with these findings [9].

Although it was generally accepted that the *neu*-protein would be a growth factor receptor, most experiments with putative ligands led to growth inhibition instead of growth stimulation. This finding seems to contrast with the oncogenic activity of the molecule. However, recently reported functions in cell motility are at least as important as a role in cell growth with regard to the oncogenic activity of the molecule. Cancer cells can only exhibit their invasive activity and the destruction of normal tissues by the action of motility factors and motility factor receptors. Both types of molecules render the tumour cell capable of penetrating the normal tissue and of spreading into the body, which may lead to the formation of metastases. The tissue specific routes of spread of *neu*-overexpressing tumour cells, namely, to the liver and not to the lymph nodes, can be explained by the action of specific motility factors secreted by particular organs and attracting the target cells which exhibit the particular receptor.

These statements are further supported by experiments carried out on mouse 3T3 cells, which were transfected with the mutation-activated rat *neu*-oncogene. The 3T3 cells exhibited mutative properties both in vivo and in vitro, while parental 3T3 cells did not [87]. Monoclonal antibodies capable of inducing down regulation of the *neu*-protein as well as the adenovirus 5 E1A gene product can suppress the *neu*-induced transformation of 3T3 cells. Both abrogated the metastatic properties of the *neu*-transformed 3T3 cells [88].

The ligand

Several groups of researchers have dealt with one or more putative ligands for the *neu*-protein. Initially, by

employing a series of biochemical assays and particularly by phosphorylation assays, conditioned media from *ras*-transformed fibroblasts were identified as sources of a possible ligand, which affected receptor down-regulation and tyrosine autophosphorylation of the *neu*-protein [86]. From *ras*-transformed fibroblast conditioned medium a 44 kDa *neu* stimulatory factor (NDF) was purified [53]. NDF is considered to be a differentiation factor, which induces altered morphology of breast cancer cells and synthesis of milk components [53]. Although NDF was considered as a specific ligand for the *neu*-protein after cross-linking experiments with radiolabelled NDF resulting in labelling of the *neu*-protein, the molecule is not able to affect tyrosine phosphorylation of *neu*-over-expressing ovarian cancer cells. Neither does NDF bind with high affinity to the ovarian cells, as would be expected from a ligand [54]. From these conflicting results it was concluded that NDF would require an unknown second cellular component in order to affect the *neu*-protein [54].

Almost simultaneously with the reports on NDF, heregulin was identified as a 45 kDa protein, purified from the conditioned medium of MDA-MB-231 human breast cancer cells [35]. Heregulin specifically induces tyrosine phosphorylation of the *neu*-protein. Heregulin is considered as the human equivalent of NDF. Both molecules behave similarly in bioassays. Heregulin has been identified as a ligand for the *c-erbB-3* gene product [10]. It also activates tyrosine phosphorylation of HER4 and is considered as a specific ligand for HER4. It fails to induce phosphorylation of the *neu*-protein in the absence of HER4 [56, 57]. The second cellular component, necessary for binding NDF or heregulin onto the *neu*-protein seems to be the *c-erbB-3* oncogene product. Coexpression of the *neu*-protein and the *c-erbB-3* molecule reconstitutes a high affinity receptor for heregulin [71]. It has been shown that different NDF isoforms are generated by alternative splicing and perform distinct tissue-specific functions, such as differentiation of breast cells and growth of Schwann cells [79].

Several other candidate ligands for the *neu*-protein have been reported. One of these is the 50 kDa motility factor, mentioned above [15]. A 30 kDa factor secreted from MDA-MB-231 breast cancer cells was shown to be a candidate ligand for *neu* [42]. At low concentration, the molecule stimulates growth of *neu*-overexpressing cells while it inhibits growth at higher concentration. Similar results on growth stimulation and inhibition were obtained with the 50 kDa motility factor [15]. A 25 kDa polypeptide purified from bovine kidney was also identified as a ligand for the *neu*-protein. This factor, designated NEL-GF was mutagenic for *neu*-overexpressing cells and stimulates autophosphorylation of the molecule [36]. Its relation with another 25 kDa peptide considered to be a *neu*-ligand and secreted by activated macrophages is unknown [73]. A *neu*-protein-specific activating factor (NAF) was partially purified from medium conditioned by the transformed human T-cell line ATL-2. NAF was able to stimulate tyrosine-specific kinase activity, induce

Table 1 Positive (+) or no (–) correlation of various prognostic factors with *neu*-oncogene amplification or *neu*-protein overexpression (*ER* oestrogen receptor status, *PR* progesteron receptor status, *LN* lymph node status, *OST* overall survival time)

First author of reference	Age	Premeno-pause	ER ^a	PR ^a	Grade	S-phase	LN	Aneu-ploidy	DFS	OST	Tumour size
Allred [1]	+	+	+	+	+						
Anbazaghan [2]			+	+	+	+	+	+	+	+	
Barnes [5]			+	+	+		–		–	–	
Berger [7]			+		+		+				
Borg [8]			+		+		+		+		–
De Potter [19]			+	+			–		+		
Guérin ^b [28]							+				
Gusterson [31]							–				
Heatley [34]			+				+				
Lovekin [41]			+		+		–			+	
Marks [43]									+	+	
McCann [45]	–	–	–		+		–		+	+	–
Mizukami [47]			–	–			–	–	–		
Paterson ^b [52]									+		
Poller [58]			+								
Richter King ^b [61]							–				
Schroeter [65]	–		+		+		–		+	+	+
Seshadri [67]							+				
Slamon ^b [69]	–		–	–					+	+	–
Slamon [70]									+	+	
Van De Vijver [74]							–		–	+	+
Varley ^b [76]									+	+	
Wilbur [80]			+	+	+						
Wright [82]			+		+		–		+	+	–
Zeillinger [89]			+	+							
Zhou ^b [90]			–	–			–		–		–
Zhou ^b [91]	–		–		–		–		+		–

^a + means inverse correlation between the factor and *neu*-expression

^b Southern blot

dimerization and internalization, and increase the growth of cells bearing the *neu*-protein [25]. Since none of these molecules has been sequenced and cloned, the structure and relation with NDF or heregulin is unknown. It is still waiting for the full identification of a molecule which acts as an unequivocal ligand for the *neu*-protein.

The *neu*-protein and prognosis of breast cancer

The relationship between the *neu*-protein and prognosis of breast cancer has been examined extensively, with considerable attention to tumour recurrence and patient survival. From the earliest investigations, several authors found that overexpression of the *neu*-protein is an indicator for poor prognosis in breast cancer [7, 69, 76]. In some multivariate analyses *neu*-overexpression retained its prognostic significance, being second to the classical prognostic indicators in breast cancer such as the number of invaded axillary lymph nodes, tumour size, tumour differentiation and receptor status [30, 75]. According to some investigators, *neu*-overexpressing tumours seem to behave worse particularly the first years after diagnosis [19, 75], and these patients suffer from a shorter disease free survival time and shorter overall survival time between the 3rd and 4th year after diagnosis [65, 75]. The difference in prognosis seems to vanish from the 5th year after diagnosis. However, other authors failed to find a

significant relation between overall survival time, recurrence rate and the *neu*-protein, when examined in a multivariate analysis. The most relevant prognostic factors and their relation with *neu*-overexpression are summarized in Table 1.

A strong negative association between *neu*-overexpression and oestrogen or progesteron receptor status has also been described [41, 65, 89]. The fact that most *neu*-overexpressing tumours lack steroid receptors can be explained on the one hand by the repressed expression of the *neu*-protein during oestrogen-induced proliferation and on the other by its enhanced expression during growth arrest and/or differentiation of mammary cells. Breast tumours constitutively expressing the *neu*-protein may escape oestrogen growth regulation and this condition may be associated with a particular invasive potential [14]. Experiments with human breast cancer cell lines indicate that oestrogen but not progesteron decreases *neu*-protein expression [60].

Most authors did not consider any significant relationship between *neu*-overexpression and nodal status [17, 30, 75], but, haematogenous metastases, organ metastases and liver metastases in particular, are more frequently observed in *neu*-overexpressing breast cancer patients [19, 65].

Apart from its role in the assessment of prognosis, evaluation of the *neu*-protein in breast cancer patients from a clinical point of view might become important in

the future treatment of patients by chemotherapy. An analysis of molecular markers in breast cancer suggests that increasing the dose intensity of adjuvant chemotherapy may not result in a similar benefit for all patients with positive nodes. It appears that patients whose tumours overexpress the *neu*-protein may have the greatest benefit from higher dosage of chemotherapy. Among patients who received 6 months of chemotherapy, disease free survival was significantly longer when the tumours did not express *neu*-protein. There is a significant dose-response effect of adjuvant chemotherapy with cyclophosphamide, doxorubicin and fluorouracil in patients with overexpression of the *neu*-protein but not in patients with no *neu*-overexpression. Overexpression of the *neu*-protein may therefore be a useful marker to identify the patients who are most likely to benefit from high doses of adjuvant chemotherapy [50].

The *neu*-protein and DCIS

The expression of the *neu*-protein in DCIS has been investigated by several authors. A correlation with comedo-subtype was reported [6, 75]. A relation between *neu*-overexpression and DCIS, characterized by pleomorphic nuclei and large cell type was also found [6, 21, 49]. Recently, a highly significant relationship between *neu*-overexpression, cell type and extent of disease is described [21]. The extent of disease of the *neu*-positive DCIS is significantly larger than the extent of disease of the *neu*-negative cases. *neu*-positive DCIS always belongs to the large cell group and never to the small cell group [21] (Fig. 2). According to these authors, the combination of large cell type DCIS and *neu*-overexpression particularly, is highly correlated with tumour size and might therefore be of value in predicting recurrence after tumorectomy, due to occult DCIS in the remaining breast tissue.

This last hypothesis remains to be investigated in future prospective studies.

The *neu*-protein and Paget's disease of the breast

In Paget's disease of the breast, the epidermis of the nipple shows invasion by large neoplastic cells with a large nucleus and ample pale-staining cytoplasm. They are devoid of intercellular bridges [3]. It is most widely accepted that the intra-epidermal Paget cells originate from an underlying ductal carcinoma [51]. The cytoplasmic expression of low molecular weight cytokeratin in Paget cells and the absence of high molecular weight cytokeratin correspond with the glandular origin of these malignant cells. Low molecular weight cytokeratins are characteristic of simple and glandular epithelia and of tumour cells derived therefrom, while high molecular weight cytokeratins are cytoskeletal proteins of cornifying keratinocytes and of tumour cells originating from this epithelium [12, 27, 48].

Overexpression of the *neu*-protein was found in an unusually high percentage of cases in Paget's disease.

Almost a 100% positivity for the *neu*-protein has been described in Paget's disease of the breast [39, 46, 81]. The unusually high expression of an oncogene in a particular disease is highly suggestive for its causative role in the pathogenesis of this disease. Therefore, it was hypothesized that the *neu*-protein might play an important role in the pathogenesis of Paget's disease and that it might induce the spreading of the Paget cells into the epidermis.

Experiments with *neu*-overexpressing SK-BR-3 human breast cancer cells have proven that the Paget breast cancer cells spread into the epidermis because of the activity of a chemotactic factor, which is released by the epidermal cells and acts through the *neu*-protein. The chemotactic motility factor was able to induce motility of the *neu*-overexpressing SK-BR-3 cells. The factor secreted in the conditioned medium by the normal keratinocytes also provoked chemotaxis of SK-BR-3 cells and not of MCF-7 cells which are human breast carcinoma cells lacking *neu*-overexpression. Again, antibodies against the extracellular domain of the *neu*-protein were able to inhibit the chemotactic activity of the motility factor in a specific way. Taken together, these experiments suggest that the chemotactic activity of the epidermal motility factor is mediated through the *neu*-protein. The action of the epidermal motility factor through the *neu*-protein explains the pathogenesis of Paget's disease of the breast [20].

The *neu*-protein and other malignancies

neu-overexpression was found in tumours other than breast tumours. In ovarian adenocarcinomas amplification of the *neu*-gene detected by southern blotting was described in 26% of cases, which correlated with amplification detected by immunohistochemistry. Similarly to the correlation of prognosis and *neu*-overexpression in breast cancer, a statistically significant correlation between gene amplification and survival was found [70]. Again, other authors failed to demonstrate any independently adverse effect [32].

In pancreatic epithelial malignancies, *neu*-overexpression evidenced as membrane staining was only noticed in 2% of cases [33]. However, a 45% *neu*-immunoreactivity was described recently, especially in well differentiated adenocarcinomas of the pancreas [85]; the first group [33] considered cytoplasmic staining to be non-specific, the latter also included cytoplasmic staining as a sign for *neu*-positivity. *neu*-overexpression is not only confined to adenocarcinomas [72]. Membrane staining was also observed in 2–36% of bladder carcinomas and in 1–36% of squamous cell carcinomas of the bronchus [44, 78, 83]. By Southern blot *neu*-expression was shown to be very rare in small cell lung cancer [64].

Conclusion

Immunohistochemical expression of the *neu*-protein on the tumour cell membrane in nearly 20% of breast carci-

nomas is correlated with a decreased overall and disease free survival and a negative oestrogen and progesterone receptor status. *neu*-overexpression may increase the metastatic potential due to a growth enhancing effect and, particularly, to a motility stimulating effect. The *neu*-protein plays a crucial role in the pathogenesis of Paget's disease of the breast and may predict larger extent of disease in DCIS.

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