

## Role of HER2/neu in tumor progression and therapy

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**Abstract.** HER2 (human epidermal growth factor receptor-2; also known as erbB2) and its relatives HER1 (epidermal growth factor receptor; EGFR), HER3 and HER4 belong to the HER family of receptor tyrosine kinases. In normal cells, activation of this receptor tyrosine kinase family triggers a rich network of signaling pathways that control normal cell growth, differentiation, motility and adhesion in several cell lineages. The first tumor studied for an alteration of the HER2 oncogene is breast carcinoma, and so far the majority of studies have been performed on this oncotype. Although involvement of HER2 as a cause of human cell transformation needs to be further investigated, overexpression of the HER2 oncogene in human breast carcinomas has been associated with a

more aggressive course of disease. It has been suggested that this association depends on HER2-driven proliferation, vessel formation and/or invasiveness; however, poor prognosis may not be directly related to the presence of the oncoprotein on the cell membrane but instead to the breast carcinoma subset identified by HER2 overexpression and characterized by a peculiar gene expression profile, as recently identified. HER2-positive tumors were recently shown to benefit from anthracyclin treatment and to be resistant to endocrine therapy. Despite the fact that many pathways interacting with HER2 are still not fully understood, this tyrosine kinase receptor is, to date, a promising molecule for targeted therapy.

**Key words.** HER2; biological role; prognostic value; predictive value; therapy.

### Description of the normal counterpart and of its normal function

HER2 (human epidermal growth factor receptor-2; also known as erbB2) and its related HER1 (epidermal growth factor receptor; EGFR), HER3 and HER4 belong to the HER family of receptor tyrosine kinases. Individual features of this group of receptors include the ligand-less HER2 receptor and the kinase-dead HER3 receptors [1–4]. A large family of ligands (reviewed in [5]) induces receptor dimerization, with each ligand favoring some dimeric combinations over others in a specific hierarchical order, although a marked preference for HER2 as a dimer partner has been described [6, 7].

Expression patterns of HER receptors and their ligands, as well as targeted inactivation of components of the HER signaling network, have highlighted the importance of short-range ligand-receptor interactions, especially in mid-gestation processes. Apparently, the network is involved primarily in two types of interactions: (i) mes-

enchyme-epithelia cross-talk and (ii) neuronal effects on target cells, including muscle, astroglia, oligodendrocytes and Schwann cells. NRGs (neuregulins, a family of HER3 and HER4 ligands), for example, are synthesized by mesenchymal or neuronal cells and influence adjacent epithelial or non-neuronal cells, respectively, with respect to their differentiation, proliferation and migration. This may explain the crucial role of the HER receptor family in development of the cardiovascular system, nervous system, mammary gland and probably others.

An essential role for the HER receptor family in mid-gestation development was indicated by embryonic lethality of HER2- [8], HER4- [9] and NRG-deficient mice at around day 10 post-fertilization due to aberrant cardiac and peripheral nervous system development. The trabeculae, fingerlike extensions of the ventricular myocardium, fail to develop in these mice, and the resulting mutant heart is characterized by irregular beat, an enlarged common ventricle and reduced blood flow. HER3 knockout mice have less severe heart defects and consequently survive several days longer (to around embryonic day 13.5), displaying normal heart trabeculation but defective

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valve formation. The role of HER4 has been further established in HER4 knockout mice by re-expressing HER4 under the regulatory control of the cardiac  $\alpha$ -myosin heavy chain ( $\alpha$ MHC) promoter [10, 11]. Cardiac-rescued MHC-HER4 HER4<sup>-/-</sup> mice are viable but display abnormalities in the central nervous system and mammary gland. The mid-gestation cardiac defect in HER2 knockout animals has been circumvented through the use of a conditional mutagenesis approach. Conditional mutagenesis of the HER2 gene in murine ventricular cardiomyocytes [12] revealed development of severe dilated cardiomyopathy, with signs of cardiac dysfunction generally appearing by the second postnatal month. Based on these findings, the authors concluded that signaling from the HER2 receptor, which is enriched in T-tubules in cardiomyocytes, is crucial for adult heart function. In light of the adverse cardiac side effects observed in breast cancer patients treated with the monoclonal anti-HER2 antibody trastuzumab [13], an improved understanding of the molecular mechanisms by which HER2 regulates heart function is especially important.

NRGs exert many functions in neural development. Roles for NRG-1-HER signaling in neural development have been demonstrated in mice carrying an HER3-null mutation [14], as well as by selective mutations of NRG-1 ectodomains [15, 16], by MHC-HER-specific expression [10, 11] and by inducible HER2 ablation in Schwann cells [17]. In all of these mice, peripheral motor neuron axons defasciculate as they enter the muscle mass and fail to form mature neuromuscular junctions. Aberrant cranial nerve architecture and increased numbers of large interneurons within the cerebellum have been demonstrated in MHC-HER4 HER4<sup>-/-</sup> mice, while loss of NRG-1-HER signaling led to hypoplasia of the sympathetic ganglion chain and the neural crest-derived portion of the cranial sensory ganglia [14, 18–20]. Moreover, the NRG-1-HER mutants were completely devoid of Schwann cells in peripheral nerves at late development stages [15, 21]. Thus, NRG-1 signaling through the HER2/HER3 heterodimer is required for normal Schwann cell development. The role of HER2 signaling in later development of the Schwann cell lineage was also analyzed using conditional mutagenesis [17]. In that study, the conditional HER2 mutants displayed peripheral nerve hypomyelination associated with neuropathy, a phenotype reminiscent of the pathology in patients with Charcot-Marie-Tooth disease [22]. Thus HER2 is the first signaling molecule for which a role in control of Schwann cell myelination has been demonstrated *in vivo*. Moreover, Kim et al. have very recently demonstrated that HER2 signaling is also critical for oligodendrocyte differentiation *in vivo* [23]. To date however, no data have been reported on nervous system toxicity in patients treated with anti-HER2-targeted therapy. Nevertheless, patients treated long term should be carefully monitored for potential side effects,

since a possible role for HER2 in mature nervous system tissue cannot be excluded.

In contrast to the embryonic lethality caused by HER2 inactivation, mice carrying a naturally occurring germ-line mutation in the kinase domain of EGFR known as Waved-2 (hypomorphic allele with severely reduced catalytic activity) are completely viable and display only epithelial defects, such as a wavy hair phenotype. Mutant mice display impaired epithelial development in several organs, resulting in phenotypes ranging from peri-implantation death to live progeny with abnormalities in multiple organs, such as liver and skin, depending on the genetic background [24, 25].

The importance of the HER receptor family and ligands in human mammary carcinoma has evoked keen interest in the normal functions of these receptors in the mammary gland, an organ that undergoes considerable postnatal development. Analyses of HER family ligands in mammary development have revealed a complicated picture. RNAs encoding the majority of the HER-specific growth factors such as AR (anphiregulin), BTC (betacellulin), HB-EGF (heparin-binding EGF), EPR (epiregulin), EGF (epidermal growth factor), NRG1 and TGF $\alpha$  (tumor growth factor  $\alpha$ ) are all present, each with a unique temporal pattern of transcriptional regulation during the normal course of mammary development, maturation and involution [26].

HERs play several normal non-oncogenic roles in regulating growth, differentiation, apoptosis and/or remodeling in normal mammary glands. These receptors are differentially expressed in mammary epithelial and/or stromal cells during various stages of development. In the mouse virgin gland, HER1 and HER2 colocalize in all major cell types during ductal morphogenesis but localize differentially in the mature gland. EGFR and HER2 are preferentially expressed in lactating ducts and alveoli, and HER3 and HER4 are more pronounced in alveoli [26]. Interestingly, a switch from HER3 to HER4 expression was observed in the developing mammary gland, suggesting that the two receptors play different roles in mammary morphogenesis. Activated EGFR and HER2 are highly expressed in extracts of mammary glands collected at puberty, suggesting a prominent role of these receptors at this stage of development, while both are expressed to only a minor extent in mammary glands in late-stage pregnancy and in lactation [27]. By contrast, HER3 and HER4 are active in mammary glands mostly during pregnancy and lactation [28]. With respect to the signaling pathways activated by HER2, Niemann et al., [29] demonstrated that formation of branched tubules relies on a pathway involving PI-3K, whereas alveolar morphogenesis requires MAPK.

Mammary functions of HER2 have also been assessed using cytoplasmic, truncated dominant-negative form of the receptor, under the control of the mouse mammary

tumor virus (MMTV) promoter. Transgenic animals expressing dominant-negative MMTV-driven truncated HER2 have significant defects in mammary development late in gestation and early postpartum, with failure of alveolar expansion [30]. Ductal development occurs in these animals, but they have lactation problems, and mammary glands early postpartum show an immature phenotype more typical of late pregnancy [11]. These data do not reveal any indispensable role for a particular member of the family in mammary gland development. Moreover, the cross-talk between HERs and steroid hormone receptors in mammary gland development remains to be established. It seems very likely that these two receptor types act synergistically and that inhibition of both pathways is required for complete ablation of mammary gland development.

### What are the types of cancer related to the described tyrosine kinase?

The first tumor studied for an alteration of the HER2 oncogene was breast carcinoma, and so far the majority of studies have been performed on this onco-type. The literature reveals a wide variation in HER2 levels [31] within a single tumor type, most probably attributable to the lack of standardized methodologies used to assess HER2 status. The advent of standardized, FDA-approved tests for protein overexpression by immunohistochemistry (IHC) (Herceptest) (fig. 1a) or gene amplification by FISH (fig. 1b) has led to overall improvement in HER2 status assessment [32–34], as indicated by several quality control studies performed around the world [35, 36]. However, evaluation even with standardized IHC methodology has demonstrated wide variation in HER2 overexpression rates in different tumor types. The highest frequency of HER2 overexpression is found in inflammatory breast cancers scoring 3+ with Herceptest in more than 60% of the cases [37], followed by Wilm's tumor (50%), bladder cancer (44%) and non-inflammatory breast cancer (25%). In other tumors, including pancreatic, colon, ovary and lung tumors, the frequency of HER2 overexpression differs in different reports and is often greater than predicted by gene-amplification data [38–45], suggesting that overexpression in these tumors is due to gene deregulation rather than gene amplification. Breast carcinomas show the highest concordance between overexpression of the HER2 oncoprotein and amplification of the gene, making it possible to generate clinical data on the role of HER2 overexpression in disease progression.

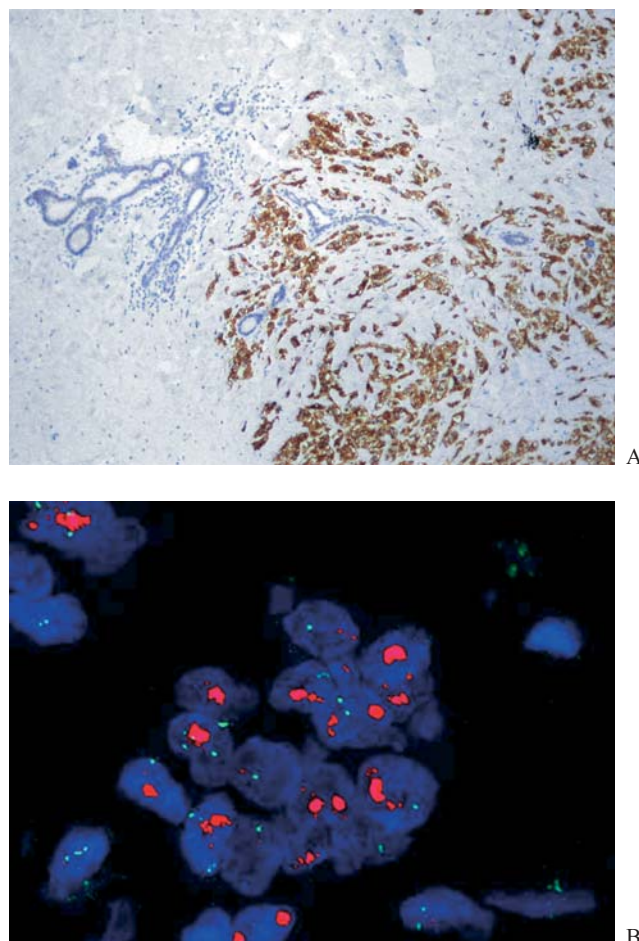


Figure 1. Evaluation of HER2 status in breast carcinomas. Paraffin-embedded sections were analyzed by immunohistochemistry (IHC) using Herceptest KIT supplies (DAKO) (A), or by fluorescence in situ hybridization (FISH) using HER2 Spectrum Orange-labeled probe, centromeric 17 Spectrum Green-labeled probe (Vysis) and DAPI as nucleus counterstaining (B)

### What is known about cause/effect and the generation of certain tumors

Transgenic mice models have shown that the oncogenic rat HER2/neu, with an amino acid change in the extracellular domain is highly tumorigenic in mice with an early onset of mammary carcinomas in both females and males [46]. Also, the rat protooncogene HER2/neu induces mammary carcinomas, but the onset is late, variable and restricted to females [47]. In mice transgenic for this protooncogene, gene deletions involving removal of cysteine residues in the extracellular domain of the receptor were found to mediate transformation [48]. Similar deletion was not found in humans, where, however a splice variant of HER2 messenger RNA (mRNA), excluding the small exon 19 containing two cysteine residues was observed [48]. This physiologic splice variant of HER2 mRNA, whose role in human cell transformation is still not de-

finer, represents about 5% of all HER2 mRNA, and it has been shown to transform *in vitro* cells with high efficiency due to its constitutive activation occurring following spontaneous dimerization (fig. 2).

Despite the involvement of HER2 in cancer, its role in human cell transformation needs to be further investigated. Recent data have shown that HER2-positive breast carcinomas represent a particular tumor group with peculiar behavior. Various clinical studies have evaluated the relationship between HER2 and breast cancer outcome, and most have shown that women with HER2-positive tumors have a poorer prognosis than women with HER2-negative tumors [49–51]. However, while the prognostic value of HER2 amplification/overexpression in node-positive patients has been widely demonstrated, there is no consensus on its value in node-negative cases [52–55]. Although a few studies on small series have shown some prognostic impact of HER2 positivity in node-negative patients, others, including our study of a large cohort of node-negative cases, argue against a prognostic role for HER2 in this patient subset [56]. Moreover, in a recent microarray analysis to identify genes associated with poor prognosis (poor signature) and good prognosis (good signature) in node-negative patients, HER2 was not included in the 70 prognostic genes identified [57].

In any event, the prognostic impact of HER2 positivity is related only to the first 3–4 years after surgery, as indicated by the peak of early recurrences [56]. The reasons for early recurrences in HER2-positive tumors have been suggested to rest in events occurring at time of surgery. Indeed, growth factors released during wound healing [58, 59] have been shown to preferentially stimulate the growth of HER2-positive tumors [60, 61]. These growth factors are more likely to have a stimulatory effect in patients with disseminated micrometastasis (node-positive patients) of an HER2-positive tumor, which might also explain the prognostic impact of HER2 according to nodal status.

There is considerable interest in biologic markers able to predict the response of cancer patients to therapy. HER2 overexpression has been implicated as a potential indicator of responsiveness to doxorubicin. Indeed, the study by the Cancer and Acute Leukemia Group B (CALGB) of node-positive patients randomly allocated to three dose levels of CAF (cyclophosphamide, doxorubicin and fluorouracil) [62], as well as the study by Paik et al. [63] from the National Surgical Adjuvant Breast and Bowel Project (NSABP) examining the effect of doxorubicin in node-positive patients, and others [64] indicated that administration of doxorubicin was of significant benefit in HER2-positive tumors but without any beneficial effect in HER2-negative tumors.

Contrary to expectations based on most previous studies, HER2-positive tumors were recently shown to benefit

from cyclophosphamide, methotrexate and fluorouracil (CMF) treatment [65, 66]; however, addition of doxorubicin to the CMF regimen further improved survival only in patients with HER2-positive tumors [67]. HER2 positivity has been related to endocrine therapy unresponsiveness, even in hormone receptor-positive patients [68, 69]. The recent observation that the level of expression of estrogen receptor is inversely correlated with HER2 expression [70], together with clinical data indicating that only high estrogen receptor-expressing tumors are sensitive to the anti-estrogen reagent tamoxifen [71], likely explain the tamoxifen resistance of HER2-positive tumors.

### **Is the described tyrosine kinase structurally altered or overexpressed in neoplastic cells?**

#### **Molecular process for such alteration**

In breast carcinomas, gene amplification is the key event for HER2 overexpression, whereas no mutation or genetic recombination has been described in human tumors, as opposed to experimental tumors in HER2/neu transgenic mice.

A new subset classification of breast carcinomas has recently been proposed based on the gene expression profile revealed by microarray analysis [72, 73]. HER2 overexpression, which is associated with overexpression of other genes that are probably co-amplified in the same amplicon, together with a series of other co-expressed genes, has been reported to identify a subset of tumors characterized by the lack of expression of genes associated with hormone receptor signaling pathways; high-level expression of a cluster of genes associated with proliferation; and expression of keratins associated with undifferentiated stem cells [74]. These findings suggest a pattern of gene expression peculiar to HER2-expressing tumors, and raise the possibility that all the clinical features thus far associated with HER2, e.g., prognosis, prediction of response to therapy, might actually be related to the biological behaviors of the subset and not directly to the presence of the oncoprotein. For example, sensitivity to anthracyclins might be related to the growth characteristics of the HER2-expressing subset rather than to the HER2 receptor pathway itself, as also suggested by *in vitro* studies [75].

### **What is known about the signal transduction pathways through which the tyrosine kinase exerts its oncogenic activity?**

The diversification of HERs family during the evolution from one receptor/ligand in worm to four receptors/multiple ligands in mammals has created a complex network able to activate distinct signaling molecules downstream



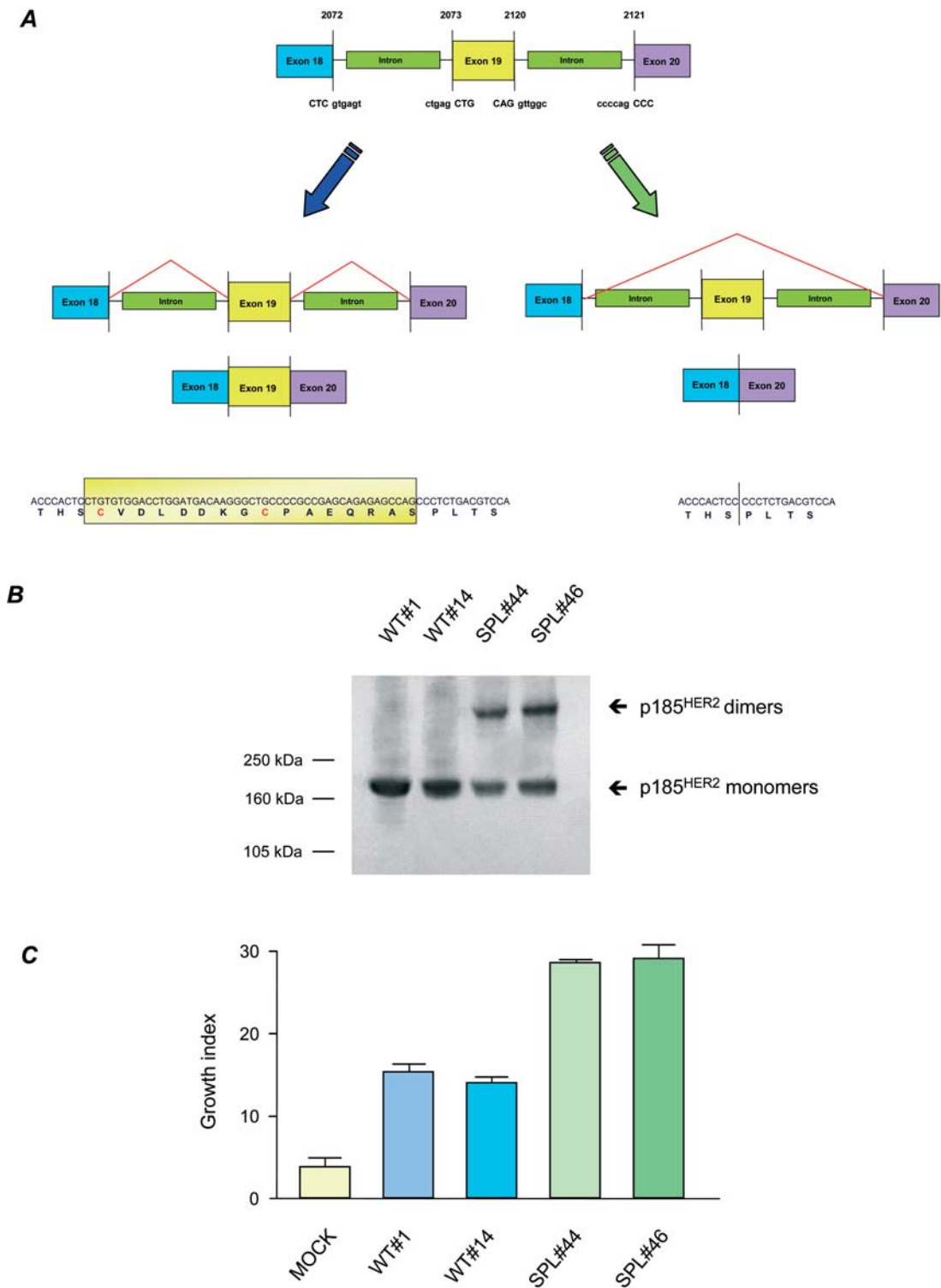


Figure 2. Schematic representation of the aberrant splicing event leading to the deletion of exon 19 (*A*). Western blot analysis of soluble extracts from NIH3T3 cells transfected with wild-type (WT#1 and 14) or splice (SPL#44 and 46) HER2 electrophoresed under non-reducing conditions, using monoclonal antibody against the intracellular domain of HER2 (*B*). Proliferation of the same transfected clones; MOCK represents NIH 3T3 cells transfected with empty vector (*C*).

of each receptor, providing a high degree of signaling diversity [76–78]. In normal tissue, the ligand-less HER2 functions only as a heterodimer with a ligand-bound receptor of the family, resulting in the most prevalent dimers in several human tissues; HER2-containing heterodimer generates a stronger intracellular signal than that originating from other complexes (reviewed in [5]). In normal cells, activation of this receptor tyrosine kinase family triggers a rich network of signaling pathways that control normal cell growth, differentiation, motility and adhesion in several cell lineages [8]. The formation of heterodimers and the ensuing activation are temporary and spatially controlled in normal cells and tissue, but deregulation of this network has been reported in cancer cells, where increased expression levels of HER2 (e.g., in breast cancer) [79, 80] or HER1 (e.g., in lung cancer) [81] and the presence of an autocrine secretory loop of ligands or a paracrine growth induced by ligands secreted by adjacent stroma cells provide a growth advantage. Indeed, overexpression of HER2, which occurs in about 30% of breast and ovarian cancers [82], activates the PI3K/Akt ‘survival’ pathway, whose action favors cell proliferation apparently by inhibiting apoptosis [80, 83–85]. The ‘survival’ signal is normally coupled to the activation of the mitogenic signal involving MAPK-pathway recruitment. Both pathways mediate control on cell cycle progression by regulating effectors such as cyclin D1 and CDK inhibitors (p27 and p21) and by promoting ubiquitination of p53 [86–88]. Increased HER2 expression in cancer enhances and prolongs signaling from both the PI3K/Akt and MAPK pathways [89, 90] and induced deregulation of cell cycle checkpoints, associating upregulation of this receptor to the malignant phenotype. Hyperactivation of the PI3K/Akt pathway, observed in tumors upregulating HER2, and the genetic defect of molecules along apoptotic pathways prevent tumor cell apoptosis in these cancers. Recent proof that HER2 induction of proliferation in a normal mammary cellular context requires cooperation with survival-promoting factors supports the idea that the type of signal originating from overexpressed HER2 depends on additional defects in cell cycle control and suppressor gene activity [91]. In normal tissues, signaling by the HER family is controlled by several mechanisms that ensure the appropriate tuning of signals: dephosphorylation of receptors, dissociation of ligand receptor, downregulation of receptors by endocytosis that can target these molecules either to lysosomal degradation (promoting signal attenuation and decrease in mitogenicity) or receptor recycling to the plasma membrane (resulting in potentiation of receptor signaling). Hyperactivation of HER2 in cancer cells slowed down the major part of these processes (reviewed in [92]). New negative regulatory pathways of the HER family are now emerging, such as the receptor-associated late transducer (RALT)

pathways, able to suppress the mitogenic and survival signal induced by HER receptors by directly interacting with all the different HER dimers with different relative potencies [93].

The tumorigenic action of HER2 is not limited to a potential proliferative effect. In fact, HER2 has been shown to be a metastasis-promoting factor. Changes in HER2 levels and in its activation by different EGF-like and heregulin (HRG) ligands have been associated with increased invasiveness *in vitro* and a more metastatic phenotype *in vivo*. A key role for HER family members in enhancing metastatic potential rests in their ability to promote secretion of basement membrane degradative enzymes, such as the matrix metalloproteases (MMPs) [94, 95], which determine modifications in the tissue architecture through the breakdown of the matrix and consequent perturbations of cell-cell and cell-matrix interactions. These alterations, together with changes in integrin and cadherin function frequently observed in tumor cells with activated HER, facilitate the communication between tumor cells and their escape from control by the microenvironment. In particular, the creation of new interactions of HER family members with integrins, such as HER2 and  $\alpha_6\beta_4$  (fig. 3) and focal adhesion kinases recruit the PI3K pathway [96–98], inducing release of second messengers (phosphoinositides and calcium). These molecules in turn recruit multiprotein complexes that play a role in reorganization of actin-cytoskeleton and thus in tumor cell migration [99, 100]. Moreover, the high expression of HER2 in invadopodia of cancer cells suggests its involvement in cell migration [100, 101]. Although those interactions remain to be studied further,

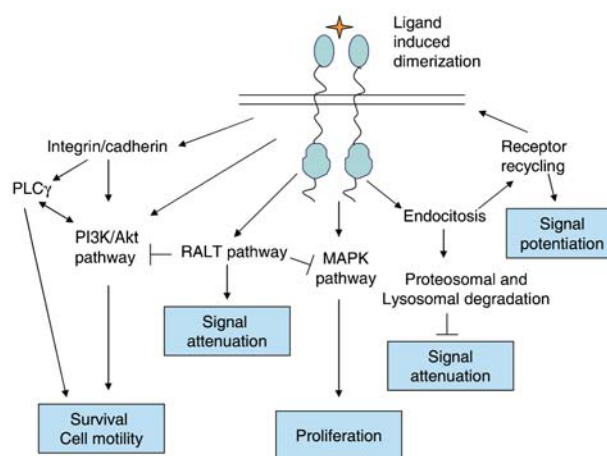


Figure 3. The major signaling pathways stimulated by activation of the HER family. Activation of receptor dimerization by ligand leads to activation of several pathways regulating proliferation, survival and cell motility. Several negative regulatory pathways counterbalance these effects by inducing signal attenuation, such as the pathway controlling HER-receptor degradation and the RALT pathways. Deregulation of HER receptors and their signaling in cancer can promote tumorigenesis.

the current body of information points to a predominant role for HER2 in motogenic and mitogenic signaling, whereas HER signaling appears to control the proteolysis of the matrix, mainly by inducing MMPs, and invasion [94, 99, 100]. By upregulating specific MMPs and angiogenic factors, such as VEGF [102], and by potentiating the adhesion of tumor cells to endothelial cells, activated HER members enhance angiogenesis and vascular invasion.

### Position of the tyrosine kinase early or late

HER2 amplification and overexpression is an early event in breast cancerogenesis. Indeed, while it has never been found altered in breast atypical hyperplasia, 'ductal carcinoma in situ' (DCIS), the pre-invasive lesion for breast cancer, is found to display HER2 overexpression in 50–60% of the cases (fig. 4), whereas the frequency decreases to 25% in invasive carcinomas. A possible explanation for this apparent discrepancy, considering a role of HER2 overexpression in increased invasiveness, is that DCIS is the pre-invasive lesion principally for HER2-positive invasive carcinomas, while the HER2-negative ones might derive directly from atypical hyperplasia.

### Specific drugs available, description of results

#### Passive immunotherapy

Differential levels of HER2 expression in normal versus HER2-overexpressing tumor cells, together with the clear involvement of HER2 in tumor progression, make HER2 an ideal target for therapeutic approaches. Monoclonal antibodies specifically directed against the extracellular domain of HER2 have been shown to selectively inhibit

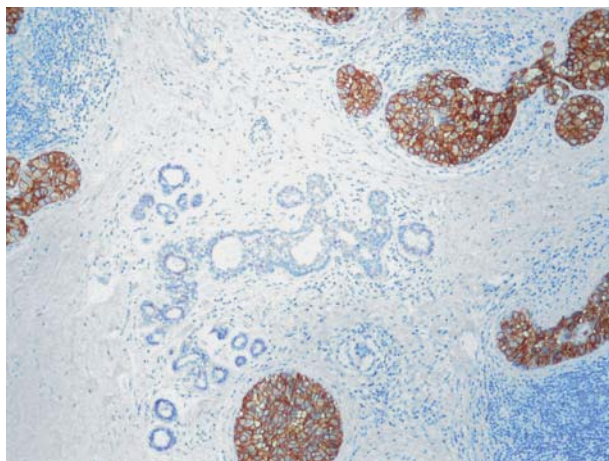


Figure 4. Overexpression of HER2 analyzed by immunohistochemistry (IHC) in ductal carcinoma in situ (DCIS).

growth of HER2-overexpressing cancer cells. One such antibody, 4D5 [103], recognizes an extracellular epitope in the cysteine-rich II domain residing very close to the transmembrane region [104]. To facilitate clinical use, 4D5 has been humanized at Genentech (South San Francisco, CA) by inserting its complementarity-determining regions (CDRs) into the human immunoglobulin G1 framework [105]. The therapeutic activity of the humanized antibody, known as trastuzumab, has been evaluated as a single agent given before [106] or after [107] traditional chemotherapy, and in combination with a variety of chemotherapy agents [13], in women with HER2-overexpressing metastatic breast cancer. The results indicated a therapeutic benefit of addition of trastuzumab to the therapeutic protocol. A recent study has also shown that pre-operative trastuzumab in combination with other chemotherapeutic agents was active against HER2-overexpressing early-stage breast cancer [108]. Thus, trastuzumab has become a standard of care for women with HER2-overexpressing metastatic breast cancer, and its clinical efficacy seems to be clear. However, not all patients with HER2-overexpressing tumor found benefits by trastuzumab treatment, indicating the necessity to optimize the use of this therapeutic tool through the definition of its mechanism of action in vivo, which it is not yet completely understood. Analyses of trastuzumab activity in experimental models have evidenced at least three different possible mechanisms: HER2 downregulation, antibody-dependent cell cytotoxicity (ADCC) and alteration of vessel development. In vitro treatment of HER2-overexpressing breast carcinoma cell lines with trastuzumab resulted in downmodulation of the receptor and inhibition of tumor growth [109, 110]. Indeed, trastuzumab induces obligate formation of HER2 homodimers, leading to an increase of ligand-mediated endocytosis of the receptor and, consequently, to significant removal of HER2 from the plasma membrane and decreased receptor-initiated constitutive signaling [104, 109, 111]. Several other phenotypic changes accompany trastuzumab binding to HER2-overexpressing cells in vitro, including induction of p27KIP1 and of the Rb-related protein, p130, with consequent reduction of the number of cells in S-phase [112]. In keeping, we also reported that ectopic expression of Rb2/p130 suppresses the tumorigenicity driven by HER2 overexpression in ovarian cancer cells [113]. To date, experiments evidencing a trastuzumab dose-dependent antitumor effect on human tumor cell lines xenografted in athymic mice have not included analysis of disappearance of the receptor [114]. In contrast, the activity of trastuzumab examined in animal models was found to depend on the engagement of Fc receptor-expressing lymphocytes [115], indicating ADCC as the major mechanism of antibody action. In FcR $\gamma^{+/+}$  nude mice injected subcutaneously with HER2-overexpressing human breast carcinoma cells, trastuzumab treatment re-



sulted in near-complete (96%) inhibition of tumor growth. This protective effect was reduced more than 50% after disrupting the antibody's ability to engage cellular Fc $\gamma$  receptor or in antibody-treated FcR $\gamma^{-/-}$  mice. A recent study in a preclinical model consisting of immunodeficient SCID mice transplanted with human breast carcinoma cells overexpressing HER2 showed that trastuzumab treatment inducing a 30% reduction of tumor volume was accompanied with a reduction in vessel volume, introducing a new mechanism of action for this antibody [116]. Indeed, HER2 signaling is known to control the expression of pro- and anti-angiogenic factors, including VEGF [102, 117], TGF $\alpha$  [118] and TSP-1 [116]. However, this finding does not necessarily exclude ADCC, since in a model such as the SCID mouse, which is profoundly deficient in T and NK cells, inhibition of HER2-driven pathways by the anti-HER2 antibody may contribute significantly to anti-tumor efficacy. The anti-tumor activity through vessel regression, together with the low protection exerted by trastuzumab found in mice knocked out for the Fc receptor, suggest that trastuzumab can utilize different mechanisms *in vivo*.

Clearly, the precise delineation of trastuzumab's mechanism of action in patients is essential for the design of more successful antibody treatment protocols in breast carcinoma patients. If, in fact, the major mechanism is ADCC, the combination of trastuzumab with immunosuppressive chemotherapeutic drugs requires optimal timing of trastuzumab delivery to enable rescue of ADCC effectors such as NK cells after chemotherapy. In that context, clinical studies have demonstrated a synergistic action between trastuzumab and the drug taxane [119], which induces suppression of adaptive immunity, but selectively increases NK activity [120]. To improve therapeutic activity, stimulation of NK activity might also be considered, utilizing cytokines or 'danger signals' such as unmethylated CpG-oligodeoxynucleotides, which enhance innate immunity [121].

### Active immunotherapy

Cancer vaccination is based on the induction of long-lasting immunologic memory provided by expanded populations of T- and B-lymphocytes with cytotoxic potential and tumor antigen specificity.

Although HER2 is generally indicative of a poor prognosis, its overexpression is associated with a better outcome when inflammatory infiltrates are present in the tumor (fig. 5), suggesting a role for HER2 in tumor immunosurveillance [122–124]. In keeping, development of HER2-specific antibodies has been documented in some patients with primary HER2-positive tumors [125–127], and T-cells reactive to HER2 were found to occur naturally in patients with HER2-positive tumors [128], confirming the immunogenicity of the molecule.

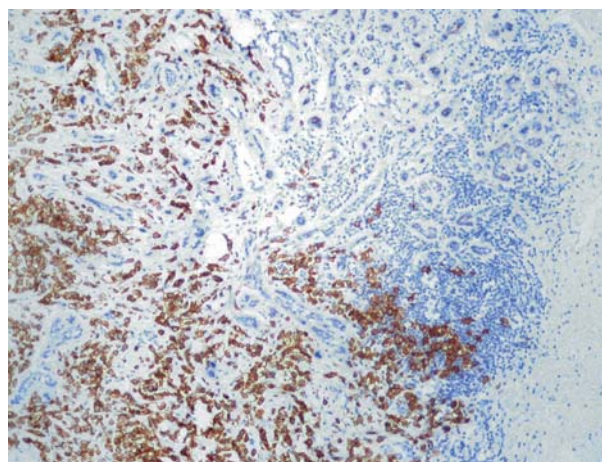


Figure 5. Presence of a marked inflammatory infiltrate in HER2-overexpressing infiltrating breast carcinoma. Paraffin-embedded tissue was probed with DAKO kit.

HER2-derived vaccines have been used in efforts to redirect immunity to induce rejection of HER2-positive tumors [129, 130]. Immunization regimens of active immunotherapy have been devised that generate specific T-cell responses with or without accompanying antibody responses and are currently being tested in animal models or in clinical trials [129–134].

Animal studies have shown that cancer vaccines generally display only limited or no efficacy in curing established tumors in syngeneic models [135]. Furthermore, the limited therapeutic efficacy of vaccines was lost when they were not given in the first few days after tumor cell implantation when tumor burden is still very small [136, 137]. These experimental findings point to the need for better selection of patients for enrollment in immunotherapy trials, such as patients with a minimal residual disease after conventional treatments or patients bearing tumors at an early stage of disease, to allow more reliable and objective clinical results [138–140].

Immunotherapeutic strategies such as immunization with MHC class I- and class II-restricted HER2-specific peptides with or without adjuvants, HER2 DNA, HER2 recombinant protein and dendritic cells loaded with HER2 peptides are now being tested in animal models and phase I clinical trials [129, 130, 141, 142]. In the majority of immunized patients, long-lasting T-cell-mediated immunity was revealed by *in vitro* and *in vivo* monitoring and, most importantly, immunity to the HER2 protein was significantly associated with epitope 'spreading', reflecting the initiation of an endogenous immune response [142].

In preclinical studies, the majority of cancer vaccines have been found to induce both T- and B-cell responses, even when no therapeutic benefits on experimental implanted tumors are observed.

While poorly active in inducing an anti-tumor therapeutic response, most of the vaccines are effective in preventing



spontaneous tumor occurrence [139]. Most data on the preventive potential of HER2 vaccines have come from studies in genetically predisposed animals transgenically expressing the rat HER2/neu proto-oncogene or its mutated activating form, and which are tolerant to self neu [143]. Different vaccination strategies include rat HER2/neu-positive allogeneic cells alone [144] or in combination with interleukin-12 [133]; the extracellular domain of the rat oncoprotein [145] or peptides derived from it [146]; and DNA plasmids encoding different fragments of HER2 or the full-length gene [131, 134, 147]. Immunological interventions targeting the HER2 protein, which is expressed at basal levels on adult normal tissues, have raised safety concerns with respect to the potential induction of autoimmune toxicity [148]; however, at present, no pertinent data in tumor-bearing hosts have been obtained [130, 149].

While therapeutic and prophylactic cancer vaccines may offer the advantage of inducing the development of both T- and B-cell-mediated long-term immunity, unlike the apparently short-lived therapeutic effects of passive therapies, an optimal vaccine formulation with objective therapeutic benefits remains to be defined. This goal awaits an improved understanding of the molecular mechanisms underlying the immunological escape of HER2-positive tumors.

### Anti-receptor therapy

As previously reported, signals induced by HER2-containing heterodimers have the strongest biological activity with respect to the other HERs dimers [5], and high expression of HER2 in tumors leads to amplification of the signaling cascade induced by the EGF-family ligands [92, 150]. Since HER2 is also the preferred partner of HER1, inhibition of HER1 activity has been proposed as an avenue to block HER2 activity. The recent availability of tyrosine kinase inhibitors (TKIs) that specifically block the phosphorylation and function of HER1 has provided a powerful new tool for therapy of tumors whose proliferation depends on HER1 or HER2. One such inhibitor is ZD1839 (Iressa), an orally active, selective HER1 TKI that induces a reversible inhibition or a delay in the growth of cancer cell lines and human tumor xenografts expressing high levels of HER1 [151]. Following studies showed that tumors expressing low levels of HER1 or overexpressing HER2 in presence of normal/low HER1 are also sensitive to ZD1839 [152–155], whereas resistance to this TKI has been observed in cell lines with a high to low level of EGFR, suggesting that expression of the target receptor is not a marker of sensitivity to ZD1839 [154, 155]. In addition, treatment of breast carcinoma cell lines expressing both HER2 and hormone receptors with ZD1839 was found to restore the inhibitory activity of tamoxifen [156], suggesting the

usefulness of TKI-mediated disruption of HER2 signaling in abrogating the anti-estrogen resistance observed in HER2-positive breast tumors. The precise mechanism through which Iressa exerts its antitumor effect remains unclear. The first step results in the inhibition of the EGFR tyrosine-kinase activity, leading to a blockage of the activation pathways that mediate EGFR functions. The failure of ZD1839 with conventional chemotherapeutic agents in phase III trials on unselected NSCLC have highlighted the lack of knowledge of the ZD1839 mechanism of action as well as the absence of reliable markers of response to this drug (clinical endpoints). Elucidation of these points is mandatory to further design of new schedules of treatment with target-specific and conventional chemotherapies.

Despite the abundance of studies on the biology of HER2 and its receptor family, many pathways interacting with HER2 are still not fully understood or exploited for therapeutic purposes. However, HER2 is, to date, one of the most promising molecules for targeted therapy.

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