ErbB2 Dephosphorylation and Anti-Proliferative Effects of Neuregulin-1 in ErbB2-Overexpressing Cells; Re-evaluation of Their Low-Affinity Interaction

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Neuregulin-1 binds to ErbB3 and ErbB4 and regulates cancer proliferation and differentiation. Neuregulin-1 had been suggested to also react with ErbB2, but this argument becomes controversial. Here, we re-evaluated the cellular responses and ErbB2 interaction of neuregulin-1 in ErbB2 overexpressing cell lines. In a competitive ligand-binding assay, we detected significant replacement of [35S]-labeled neuregulin-1 with nano molar ranges of cold neuregulin-1 in L929 cells expressing ErbB2 alone and SKOV3 cells carrying sulf-1 cDNA but not in these parental cells. The concentration of neuregulin-1 significantly decreased thymidine incorporation and phosphorylation of ErbB2 (Tyr877, Tyr1396, and Tyr1121) in ErbB2-overexpressing cancer cells as well as in L929 cells expressing ErbB2. A crosslinking assay ascertained the presence of neuregulin-1 immunoreactivity in the ErbB2 immune complexes of L929 expressing ErbB2 alone. These results suggest that the higher concentrations of neuregulin-1 exert an anti-oncogenic activity to attenuate ErbB2 auto-phosphorylation potentially through its low-affinity interaction with ErbB2.

A receptor molecule for neuregulin-1 (neu-differentiation factor or heregulin) was initially suggested to be not only ErbB3 and ErbB4 but also ErbB2 (Her2, Neu)1,2. The following studies demonstrate that the high affinity interaction of ErbB2 with neuregulin-1 involves ErbB3 heterodimerization and tone down the argument for the direct ErbB2–neuregulin-1 interaction3-6. ErbB2 is often overexpressed with or without gene amplification in various cancer cells, such as mammary cancer cells and ovarian cancer cells7-10. ErbB2 overexpression results in ligand-independent self-dimerization and subsequent auto-phosphorylation, leading to downstream signaling linked to cell proliferation and migration. Thus, the extent of ErbB2 overexpression in cancer cells is sometimes correlated with malignancy and/or a poor prognosis11,12. Recently, various anti-cancer drugs that target ErbB2 have been developed, including anti-ErbB2 antibodies that inhibit ErbB2 self-dimerization or induce its down regulation13,14. In this regard, ErbB2 auto-phosphorylation play key roles in the proliferation of the cancer cell lines overexpressing ErbB215-17.

After two initial reports demonstrated an interaction between neuregulin-1 and ErbB2, subsequent studies questioned the receptor function of ErbB2. There are two major reasons why ErbB2 cannot function as a high-affinity receptor for neuregulin-1. First, many ovarian cancer cell lines that express high levels of ErbB2 do not exhibit an increase in ErbB2 phosphorylation in response to neuregulin-11. Secondly, [125I]-labeled neuregulin-1 exhibits a low or no binding affinity for these cells16,18. However, other studies still indicate that the direct interaction between ErbB2 and neuregulin-1 might be real despite the weak affinity16,19. In addition, subsequent studies further increase the complexity of this argument; there are positive and negative regulators for ErbB2 receptor function and subcellular localization. For example, sulfatase-1 (Sulf-1), which removes O-sulfate groups of heparin or heparan sulfate proteoglycans, significantly influence the molecular interaction of neuregulin-1.
with ErbB receptors\(^{20,21}\). Of note, approximately 70% of ovarian cancer cells lost the expression of sulf-1 and exhibit higher grades of malignancy often with ErbB2 overexpression\(^{20,21}\). With this respect, the affinity and biological role of the neuregulin-1–ErbB2 receptor interaction remains to be controversial and needs to be re-evaluated.

To clarify this issue, we carefully assessed the molecular response of an ovarian cancer line, SKOV3, and a breast cancer line, BT-474, to neuregulin-1, both of which highly express ErbB2. We also used L929 mouse fibroblasts that lack the expression of ErbBs or those transfected with \(erbB2\) cDNA. Molecular interactions of ErbB2 with neuregulin-1 were examined with novel procedures for ligand labeling and immunoprecipitation. In particular, we manipulated the expression of heparan sulfatase-1 in the ovarian cancer cells as the O-sulfate structures markedly alter receptor interactions of the heparin-binding growth factors such as neuregulin-1\(^{20,21}\). With the obtained results, we discuss the involvement of ErbB2 in neuregulin-1 signaling and functions.

**Results**

**Expression of ErbB1-4 proteins in L929, SKOV3, and BT-474 cells.** The primary amino acid sequences of the ErbB molecular family members share high structural homology and are often cross-recognized by antibodies raised against different subtypes. As the previous reports demonstrate that mouse fibroblast L929 cells have no binding to neuregulin-1 and harbor the limited expression of ErbB2, we employed parental L929 cells as a negative control and transfected these cells with \(erbB1\), \(B2\), \(B3\) or \(B4\) cDNA expression vectors as positive controls\(^{22,24}\). We examined and confirmed endogenous ErbB1-4 expression in the cell lines (i.e., L929, SKOV3, SKBR3 and BT-474 cells) used in the present study (Fig. 1a). Although L929 cells expressed trace amounts of ErbB2, these cells lacked detectable levels of ErbB1, ErbB3 and ErbB4 as described previously\(^{22-25}\). SKOV3 cells expressed high levels of ErbB2 and ErbB1 and low levels of ErbB4, whereas BT-474 expressed high levels of ErbB2 and low levels of ErbB3 and ErbB1 when we compared those with ErbB1-4 levels in transfected L929 cells.

Although these cell lines contained ErbB2 proteins, they could be nonfunctional if they are not located at the cell surface. We examined the cell surface expression of ErbB2 in the above cell lines by surface biotinylation (Fig. 1b). Except for the parental L929, there were significant surface levels of the ErbB2 protein in all cell lines, which appeared to be correlated with the total ErbB2 levels.

To mimic the ErbB2-overexpression state of the cancer cells in L929 cells, we established the L929 cell line that stably carries \(erbB2\) cDNA through neomycin selection. By transiently transfecting an \(erbB2\)-expression vector, we further boosted ErbB2 expression in this cell line and used these ErbB2-overexpressing L929 cells (referred to as ErbB2-L929 hereafter) in the following experiments. Again, we confirmed that the transfection of \(erbB2\) cDNA and G418 selection did not induce the expression of ErbB3 and ErbB4 in ErbB2-L929 cells (Supplemental Figure S1).

**Responses of ErbB2-overexpressing cells to neuregulin-1.** The ovarian cancer cell line, SKOV3, is reported not to have a high affinity for neuregulin-1\(^{26}\). We re-examined the cellular responses of SKOV3 cells to relatively high concentrations of neuregulin-1, monitoring their cell proliferation and competitive binding activity. The rates of cell proliferation were measured using [\(^{3}H\)] thymidine incorporation in the presence or absence of 10 nM and 40 nM type I neuregulin-1 (a soluble form comprising the extra-cellular domain; ECD) as well as with or without the transfection of sulf-1 (sulfatase 1) cDNA (Fig. 2a). As blood serum contains high levels of neuregulin-1, SKOV3 cells were extensively washed prior to the assay\(^{26}\). The concentration of 10 nM neuregulin-1 had no apparent influence on thymidine incorporation, whereas 40 nM neuregulin-1 significantly decreased incorporation compared with untreated controls. SKOV3 cells lack the enzyme Sulf-1, which selectively removes the 6-O-sulfate group from heparan sulfate\(^{27}\) and regulates neuregulin-1 binding to ErbB receptors\(^{29}\). We also tested the effect of Sulf-1 expression on the thymidine incorporation of SKOV3 cells. The transfection of sulf-1 cDNA further increased the response to neuregulin-1. In addition, we assessed the effects of 40 nM neuregulin-1 on BT-474 cells, ErbB2-L929 cells, and parental L929 cells (Fig. 2c, d). We observed a similar anti-proliferative activity of neuregulin-1 in BT-474 cells and ErbB2-L929 cells, but not in parental L929 cells.

We recently developed a simple method that enables us to synthesize and label authentic neuregulin-1 with the radioisotope [\(^{35}S\)] methionine\(^{28}\). This procedure does not involve additional chemical modification, such as iodination and/or oxidization of the target protein, we can avoid the structural hindrance of the additional atom in the receptor-binding assay. In fact, we found that the iodination of neuregulin-1 proteins (i.e., NRG1-ECD and an EGF domain peptide of neuregulin-1; eNRG1) significantly reduced their biological activity (Supplemental Figure S2). In the present study, therefore, we prepared a [\(^{35}S\)]-labeled ligand of neuregulin-1 by this in vitro translation system. We performed the competitive ligand-binding assay in...
SKOV3 cells with this [35S]-labeled ligand (Fig. 2b). As reported previously, there was no significant replacement of [35S]-labeled neuregulin-1 in SKOV3 cells. Again, we examined the effects of sulf-1 cDNA transfection on neuregulin-1 binding to this cell line. When the sulf-1 gene was transduced into these cells, there was a significant replacement of [35S]-labeled neuregulin-1 with 40 nM of cold neuregulin-1. The absence of Sulf-1 protein expression in the parental SKOV3 cells as well as its presence after transfection was confirmed by Western blotting (Supplemental Figure S3).

**ErbB2 dephosphorylation following neuregulin-1 challenge.** The above results suggest the biological implication of ErbB2 signaling in the anti-proliferative responses to neuregulin-1. We examined time-dependent alterations in ErbB2 phosphorylation (Tyr877, Tyr1139, and Tyr1196) of SKOV3 cells following neuregulin-1 challenge (Fig. 3a). Western blotting with anti-phospho ErbB2 antibodies suggested that exposing SKOV3 cells to 40 nM neuregulin-1 rapidly decreased the phosphorylation levels of ErbB2 at Tyr877, Tyr1139, and Tyr1196. Statistical analyses revealed that the decrease in Tyr1196 phosphorylation was statistically significant and continued at least for 3 h. In contrast, there were no apparent alterations in the total protein levels of ErbB2 and β-actin.

Similarly, we examined time-dependent ErbB2 dephosphorylation in another ErbB2-overexpressing cell line, BT-474 (Fig. 3b). BT-474 cells exhibited anti-proliferative responses to neuregulin-1 (see Fig. 2) whereas these cells also express ErbB3 whose signaling promotes cell proliferation29,30. Neuregulin-1 treatment indeed elevated ErbB3 phosphorylation in BT-474 cells. In parallel, neuregulin-1 challenge triggered the rapid dephosphorylation of ErbB2 at Tyr877, Tyr1139, and Tyr1196 as was seen in SKOV3 cells. In contrast to the effects on SKOV3, ErbB2 dephosphorylation at Tyr1196 did not continue more than 180 min, however. It was not evident that the observed dephosphorylation might involve the interaction of neuregulin-1 with ErbB4 and ErbB3 receptors expressed by SKOV3 and BT-474 cells, respectively. We examined ErbB2 phosphorylation in ErbB2-L929 cells (Fig. 3c). When we challenged ErbB2-L929 cells with neuregulin-1, we found similar rapid decreases in ErbB2 phosphorylation at Tyr877, Tyr1139, and Tyr1196. Statistical analysis of Tyr1196 phosphorylation verified that there was a significant reduction in ErbB2 tyrosine phosphorylation. In comparison with the duration of the dephosphorylation in SKOV3 and BT-474 cells, the ErbB2 dephosphorylation had a shorter duration and did not continue more than 20 min in ErbB2-L929 cells.
neuregulin-1 is undetectable in parental L929 cells 24. agrees with the report that the cytotoxicity of toxin-conjugated significant replacement with 10 nM cold neuregulin-1. This result

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Figure 4 | Competitive binding assay of [35S]-labeled neuregulin-1 to L929 cells overexpressing ErbB2. Does ErbB2 dephosphorylation stem from the molecular interaction or association of ErbB2 with neuregulin-1? To challenge this question, ErbB2-L929 cells as well as parental L929 cells were examined in a competitive ligand-binding assay (Fig. 4). The total binding of [35S]-labeled neuregulin-1 to ErbB2-L929 cells was gradually diminished with increasing concentrations of cold neuregulin-1. IC50 of the neuregulin-1 binding was approximately 10 nM. In contrast, parental L929 cells exhibited reduced total binding to [35S]-labeled neuregulin-1 and failed to show any significant replacement with 10 nM cold neuregulin-1. This result agrees with the report that the cytotoxicity of toxin-conjugated neuregulin-1 is undetectable in parental L929 cells24.

Since we did not modify the chemical structure of the ligand, [35S]-labeled neuregulin-1 should have the same affinity to ErbB receptors as authentic neuregulin-1. Thus, we can assume that Ki is equal to Kd in the Cheng-Prusoff equation. According to this equation, we estimate that Kd of neuregulin-1 binding was ~12 nM or less with the given IC50 (approximately 10 nM) and the concentration of the tracer (i.e. 1.9 nM) in this assay.

**Appearance of the ErbB2 complexes crosslinked to cold neuregulin-1.** Using chemical crosslinking of cold neuregulin-1 to ErbB2, we attempted to visualize the receptor complexes of neuregulin-1 and ErbB2 using Western blotting with the anti-neuregulin-1 antibody. In our preliminary experiment, we first tested the immunoprecipitation performance of anti-ErbB2 antibodies. SKBR3 cells, which express both ErbB2 and ErbB3, were exposed to a chemical crosslinker. Then we attempted to immunoprecipitate the ErbB2 complexes with four distinct anti-ErbB2 antibodies using the conventional immunoprecipitation procedure of the non-denatured condition (Supplemental Figure S4). However, all the anti-ErbB2 antibodies failed to immunoprecipitate the ErbB2 complexes when the cells were treated with the chemical crosslinker, even though all these antibodies were able to immunoprecipitate authentic ErbB2 protein (Supplemental Figure S4). These results suggest that the amine-reactive chemical crosslinker masks the epitopes for the anti-ErbB2 antibodies in the non-denatured condition.

Alternatively, we took the SDS-denatured condition for ErbB2 extraction and then immunoprecipitated the ErbB2 complexes (Fig. 5a, b). In the ErbB2 immune complexes as well as in the total cell lysate, we detected the neuregulin-1-like immunoreactivities slightly below the 175 kDa marker as well as above the 235 kDa marker (lanes 3 in a) whereas there were no apparent neuregulin-1-immunoreactive bands in the absence of neuregulin-1 (lane 1 in a). In contrast, the neuregulin-1-like immunoreactivity in the complexes disappeared when the chemical crosslinkage was digested, suggesting that these immunoreactivities indeed represent the molecular complexes containing neuregulin-1. When we probed the membrane with the anti-ErbB2 antibody, there were ErbB2 immunoreactivities in the immunoprecipitates (lanes 3 in b). However, the molecular size of the immunoreactivities was smaller than the authentic size of ErbB2. The digestion of the crosslinkers uncovered the masked ErbB2 epitopes and re-emerged the ErbB2 immunoreactivity at the authentic size (lane 4 in b). We found similar gel mobility shift of the ErbB2 immunoreactivity when BT474 cells were treated with a chemical crosslinker (data not shown). The phenomenon agrees with the fact that the intramolecular crosslinkage is known to fold and pack a bulky polypeptide chain and increases its gel mobility21.

Thus, we speculate that that the mobility of ErbB2 was increased by chemical crosslinkage and shifted to a lower molecular range. These results suggest that the high concentration of neuregulin-1 can associate with ErbB2.

**Discussion**

In the present investigation, we demonstrated the biological actions of neuregulin-1 in distinct cell lines harboring high ErbB2 expression. Treating with 40 nM neuregulin-1 resulted in a significant decrease in thymidine incorporation as well as in ErbB2 phosphorylation in cancer cells with erbB2 amplification. The lower dose of

**Figure 5 | A crosslink study of neuregulin-1 in ErbB2-overexpressing L929 cells.** Cells were incubated with or without 40 nM cold neuregulin-1 and then exposed to the amine-reactive crosslinker sulfo-EGS. Total cell lysates (Total) and ErbB2 immunoprecipitates (IP) were subjected to Western blotting with the anti-neuregulin-1 antibody (a) or anti-ErbB2 antibody (b). Prior to the electrophoresis, half of the immune complexes were treated with 1 M hydroxylamine to cleave the crosslinkage (lane 4). Arrowheads indicate the positions of molecular weight markers (kDa) Note; the cleavage of the crosslinker conversely elevated ErbB2 immunoreactivity (lanes 4 in b).
10 nM neuregulin-1 did not significantly affect thymidine incorporation of SKOV3 cells, however, The anti-proliferative reaction was also found in SKOV3, BT-474 and L929 cells overexpressing ErbB2, but not in the parental L929 cells. The dose-dependency of neuregulin-1 in thymidine incorporation roughly agrees with the profile of the competitive ligand-binding assay; nanomolar concentrations of neuregulin-1 were able to replace the cell binding of [35S]-labeled neuregulin-1. The competitive interaction of neuregulin-1 in SKOV3 cells required additional expression of Sulf-1, however. The low-affinity interaction between neuregulin-1 and ErbB2 was ascertained by the crosslink study as well; there was neuregulin-1 immunoreactivity in the ErbB2 immune complexes in ErbB2-L929 cells. Therefore, the present findings suggest that neuregulin-1 challenge exerts anti-proliferative effects, which presumably involve the direct or indirect interaction of neuregulin-1 with ErbB2. However, we confirmed that the biological interaction of neuregulin-1 is significantly influenced by the structures of heparan sulfate moiety on cell surfaces of ErbB2-overexpressing cells.

There were significant levels of basal tyrosine phosphorylation of the ErbB2 kinase domain in ErbB2-overexpressing cell lines. The high level of ErbB2 phosphorylation in L929 cells was achieved by both transient and permanent transfection of erbB2 cDNA. ErbB2-overexpressing L929 cells responded to neuregulin-1 and exhibited a consistent decrease in ErbB2 phosphorylation at multiple tyrosine residues. In contrast, the parental L929 cells exhibited almost no response to neuregulin-1 in the cell proliferation test and competitive binding assay. There results support the previous findings that the parental L929 cells express a limited level of ErbB2 but no detectable levels of authentic receptors for neuregulin-1. With the given proliferative action of ErbB2 auto-phosphorylation, therefore, we assume that the neuregulin-1-triggered ErbB2 dephosphorylation plays a key role in the attenuation of the cell proliferation observed.

However, these findings and/or the data explanations appear to be contradictory against the previous report that the transfection of erbB2 cDNA to L929 cells fail to induce their interaction with [125I]-labeled neuregulin-1 or alter neuregulin-1-triggered ErbB2 phosphorylation. The differences from the present results might stem from the fact that serum contains so high concentrations of neuregulin-1 (i.e. ~200 pM for human serum) that serum-supplemented culture media can perturb these assays. In fact, extensive cell wash or starvation prior to these assays was essential in the present study. Alternatively, it might be illustrated by the fact that the iodination of neuregulin-1 limits its interaction to ErbB2 receptors and perturbs the ligand-binding assay (see Supplemental Figure S2).

Still there is another argument against our conclusion; ErbB4 signaling has been suggested to underlie the anti-proliferative action of neuregulin-1 in some cancer cell lines and might contribute to the observed anti-proliferative phenomena. This scenario might potentially illustrate the results from the SKOV3 cells harboring a low level of ErbB4. However, the lack of ErbB4 expression in ErbB2-overexpressing L929 and BT-474 limits this explanation. Alternatively, the anti-proliferative action of neuregulin-1 might be ascribed to ErbB2 down-regulation. However, we did not find decreases in total ErbB2 immunoreactivity in SKOV3, BT-474, and ErbB2-L929 cells following acute neuregulin-1 challenge when we used fresh blotting membranes for ErbB2 detection (see Material and Methods). Accordingly, we assume that the neuregulin-1-dependent decrease in the proliferation of SKOV3, BT-474, and ErbB2-L929 cells involves attenuated ErbB2 auto-phosphorylation.

There are many novel molecules identified that regulate homo or hetero dimerization of ErbB2 and potentially modify direct and indirect interaction with neuregulin-1 as well; erbin, decorin, and RALT (receptor-associated late transducer). Erbin binds to the carboxyl terminal of ErbB2 and anchors ErbB2 to cytoskeletal networks to regulate its dimerization and protein stability. RALT also interacts with the tyrosine kinase domain of ErbB2 and promote its lysosomal degradation, although Anastasi et al. report the limited expression of RALT in BT-474 and SKBR3. Decorin is suggested to attenuate ErbB2 phosphorylation but involving ErbB4 interaction. These findings indicate the possibility that the homomeric interaction of ErbB2 or its autophosphorylation can be markedly influenced by these ErbB2-interacting molecules. Thus, the involvement of these ErbB2-interacting molecules might illustrate the previous controversy of the anti-proliferative action of neuregulin-1 among literatures.

The present immunoprecipitation results provide circumstantial evidence that ErbB2 functions as a low-affinity receptor for neuregulin-1. The given epitope masking of ErbB2 by chemical crosslinkage made our experiments difficult. However, this argument is consistent with a previous study using the ligand-binding assay in SKOV3 cells. [125I]-labeled type 1 neuregulin-1 is indeed replaced with 40 nM cold neuregulin-1. The surface plasmon technique demonstrates the direct interaction between ErbB2 and neuregulin-1; immobilized neuregulin-1 on a solid phase bound to the recombinant extracellular domain of ErbB2 with a calculated dissociation constant (Kd) of 850 nM, which is one digit larger than the present value, however. The variation in Kd is presumably ascribed to the difference of the experimental procedures; the in vitro solid phase system vs. the natural cell-based binding system. In the ErbB2 immunoprecipitates, we additionally detected the neuregulin-1-like immunoreactivities that carried much larger molecular sizes (>250 kDa). In this context, we do not rule out that the observed dephosphorylation of ErbB2 or the attenuation of cell proliferation may involve unknown receptors for neuregulin-1, which might form the molecular complexes with ErbB2. For example, cell-ubiquitous proteins, integrins are reported to interact directly with neuregulin-1. Alternatively, ErbB2 forms the hetero-oligomer with the other receptor tyrosine kinases as well. Thus, several possibilities and controversies still remain regarding the molecular nature of the neuregulin-1 receptor complexes found in the ErbB2-expressing cells. However the present results, at least, suggest that neuregulin-1 triggers ErbB2 dephosphorylation and the anti-proliferative action in an ErbB2 expression-dependent manner. Thus, our present study reveals and confirms an anti-oncogenic role of neuregulin-1 in ErbB2-overexpressing cancer cells.

Methods

Plasmid construction. cDNA for the human type 1 neuregulin-1 beta3 isoform (i.e. extracellular domain neuregulin-1; NRGI-1CD) was chemically synthesized by GenScript USA Inc. (Piscataway, NJ, USA) and subcloned into the pET-22b (+) plasmid (EMD Chemical Darmstadt, Germany) or pcDNA TM 3.1-TOPO TA plasmid (Invitrogen Corporation, Carlsbad, CA, USA), both of which carry T7 and/or cytomaglovirus promoters. cDNA for human Sulf-1 was synthesized from human brain RNA by RT-PCR and subcloned into the pcDNA TM 3.1-TOPO TA plasmid (Invitrogen). The expression vectors carrying human erbB1 cDNA and erbB2 cDNA were obtained from a Riken genome bank (Wako, Saitama, Japan) and those for erbB3 and erbB4 cDNAs were prepared as reported previously.

Cell culture. We obtained cell lines, SKOV3, SKBR3, BT-474 and L929 cells, from the American Type Culture Collection (ATCC, Manassas, VA, USA) and SKBR3 cells from Dr. Sakamaki (Niigata University of Pharmacy and Applied Life Sciences). Cells that overexpress ErbB2 were grown in DMEM/F12 medium (Invitrogen, Tokyo, Japan) with 10% heat-inactivated fetal calf serum (HyClone; Thermo Scientific, Yokohama, Japan). These cells were subjected to experiments within six passages. ErbB2(L929) cells were transfected with a human erbB2 expression vector and grown in Dulbecco's modified Eagle's medium (DMEM) supplemented with 5% heat-inactivated calf serum (CS) (Invitrogen, Tokyo, Japan) for two days (see below). Some of the cells were exposed to genistein G418 (0.4 mg/ml; Sigma-Aldrich Japan, Tokyo, Japan) for three weeks. Cells that were resistant to G418 and thus expressed ErbB2 were cloned. The expression of ErbB2 in the erbB2 transformants was further boosted with the transient transfection of a human erbB2 expression vector for the binding assay or phosphorylation assay.

Cell surface biotinylation. Cells were extensively washed with Hank's balanced salt solution (HBSS) and incubated with 2 mM Sulfo-NHS-biotin (Pierce/Thermo Scientific, Yokohama, Japan) at room temperature for 20 min, after which the reaction was quenched with 50 mM Tris HCl and 10 mM ethanol amine (pH 8.0). Cells were lysed with PIPA buffer [25 mM Tris HCl pH 7.6, 150 mM NaCl, 1%
NP-40, 0.5% sodium deoxycholate, and 0.1% SDS) containing a protease inhibitor cocktail (Boehringer Ingelheim Pharma Japan, Tokyo, Japan) and mixed with streptavidin beads (Sigma-Aldrich). The biotinylated proteins that were trapped with the beads were liberated by boiling with SDS sample buffer (see below).

### Competitive binding assay
According to our method,[6] 0.1% 1 mg/ml poly-D-lysine hydrobromide (Sigma-Aldrich, Tokyo, Japan). The intensity of an immunoreactive band, whose size matched the authentic molecular weight, was measured by a CCD camera and an image processing software, GENETOOLS (Syngene, Cambridge, UK). Comparable protein loading was confirmed by Western blotting using an anti-β-actin antibody (Merck Millipore, Tokyo, Japan).

### Cell proliferation assay
L929, SKOV3, and BT-474 cells were maintained at a density of 1 × 10^6 cells/ml in growth phase were plated at a density of 5 × 10^4 cells/well in poly-D-lysine-coated 48-well plates. Cells were extensively washed with Leibovitz-15 Medium (L-15; pH 7.4; Sigma-Aldrich, Japan) containing 0.1% bovine serum albumin (immunoglobulin-free grade; Sigma-Aldrich, Japan) and 10 mM HEPES. The cultures were treated with fresh medium and 1% FBS. The cells were quenched with 50 mM Tris HCl and 10 mM ethanol amine (pH 8.0). The cells were lysed with 0.5 N NaOH. The radioactivity of the cell lysates was determined using the Liquid Scintillation System LS-3050 (Aloka, Tokyo, Japan).

### Western blotting
Protein samples were prepared from cultured cells for Western blot analyses. These samples were homogenized in cell lysis buffer (25 mM Tris HCl pH 7.6, and 2% SDS) by sonication and denatured in the presence of 100 mM dithiothreitol (Wako Chemicals, Tokyo, Japan) at 85°C for 5 min. In parallel, the protein concentrations were determined using a micro-BCA kit (Protein Assay Reagent; Pierce/Thermo Scientific). Typically, 20 μg of protein was subjected to SDS-polyacrylamide gel electrophoresis and then transferred onto a nitrocellulose membrane (Advantec, Tokyo, Japan). The membrane was probed with anti-ErbB1 (Eptomine Inc., Burton CA, USA), anti-ErbB2 (Eptomine Inc.), anti-ErbB3 (Santa Cruz Biotechnology, Santa Cruz, CA, USA), anti-ErbB4 (Cell Signaling Technology, Danvers, MA, USA), or anti-phospho ErbB2 or anti-phospho ErbB3 antibodies (Cell Signaling; Eptomine Inc.). We used fresh blotting membranes to detect total and phosphorylated ErbB2 proteins as the conventional stripping procedure did not allow us to reuse the membrane that had been treated with the anti-phospho ErbB2 antibodies (data not shown). A blotting membrane was also probed with the anti-Sulf-1 antibody (Abcam, Cambridge, UK). After extensive washing, the immunoreactivity of the membrane was detected with a horseradish peroxidase-conjugated anti-rabbit immunoglobulin followed by a chemiluminescence reagent (ECL kit; ECL HealthCo, Tokyo, Japan). For the in vitro production of an active neurotrophic factor, neuregulin-1 (bissulfosuccinimidyl) substrate (BS3; Pierce/Thermo Scientific) or ethylene glycol bis(sulfosuccinimidyl) substrate (BSulf-ECS; EGS, Pierce/Thermo Scientific) and then quenched with 50 mM Tris (pH 8.0). The cells were lysed with PiPA buffer and then immunoprecipitated for ErbB2. Alternatively, cell lysates were prepared by homogenizing cells with a 1% SDS solution at 70°C and then incubated with 0.5M potassium chloride in the presence of 1% NP-40 and a protease inhibitor cocktail (Boehringer Ingelheim) to reduce the detergent. The complexes in PiPA buffer were treated with anti-ErbB2 antibodies (C-18; Santa Cruz Biotechnology, Ab-3; Merck Millipore Japan, trastuzumab; Roche, Swiss, or polyclonal 06-562; Merck Millipore Japan) for overnight and precipitated with Protein-G Sepharose (GE Healthcare, Tokyo, Japan). Alternatively, the complexes in the denaturing condition were immunoprecipitated with the ErbB2 antibody used in Western blotting (Eptomine Inc.). The total cell lysates and the immunoprecipitates were separated using 6% SDS-PAGE and probed with an anti-neuregulin-1 antibody (Santa Cruz Biotechnology) or with an anti-ErbB2 antibody (Eptomine Inc.). To control the efficacy of the ErbB2 immunoprecipitation, some cells were not exposed to the crosslinker and directly subjected to Western blotting for ErbB2. To che
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R.W. and H.N. conceived and designed the experiments and coordinated the work presented. Y.I., K.A. and N.T. performed the experiments. K.M., M.M., X.W. and S.H. provided materials and commented on the manuscript.

Additional information
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