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p38 γ Mitogen-Activated Protein Kinase Integrates Signaling Crosstalk between Ras and Estrogen Receptor to Increase Breast Cancer Invasion

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Abstract

Ras is believed to stimulate invasion and growth by different effector pathways, and yet, the existence of such effectors under physiologic conditions has not been shown. Estrogen receptor (ER), on the other hand, is both anti-invasive and proliferative in human breast cancer, with mechanisms for these paradoxical actions remaining largely unknown. Our previous work showed an essential role of p38 γ mitogen-activated protein kinase in Ras transformation in rat intestinal epithelial cells, and here, we show that p38 γ integrates invasive antagonism between Ras and ER to increase human breast cancer invasion without affecting their proliferative activity. Ras positively regulates p38 γ expression, and p38 γ in turn mediates Ras nonmitogenic signaling to increase invasion. Expression of the Ras/p38 γ axis, however, is *trans*-suppressed by ER that inhibits invasion and stimulates growth also by distinct mechanisms. Analysis of ER and its cytoplasmic localized mutant reveals that ER additionally binds to p38 γ protein, leading to its specific down-regulation in the nuclear compartment. A p38 γ -antagonistic activity of ER was further shown in a panel of breast cancer cell lines and was shown independent of estrogens by both ER depletion and ER expression. These results revealed that both Ras and ER use distinct pathways to regulate breast cancer growth and invasion, and that p38 γ specifically integrates their antagonistic activity to stimulate cell invasion. Selective targeting of p38 γ -dependent invasion pathways may be a novel strategy to control breast cancer progression.

Introduction

Two defining hallmarks of cancer are uncontrolled proliferation and increased invasion (1). Although increased proliferation is frequently associated with enhanced invasion (2), recent evidence suggests that these two events are distinct (3–6). Ras protein is involved in human malignancies through multiple processes that include increased proliferation, enhanced invasion, and altered cytoskeleton organization (2). Although studies with Ras effector domain mutants suggest a specific role of individual Ras effectors/pathways in these processes (7,8), the existence of such effectors under physiologic conditions remains unproven. Ras protein is overexpressed in up to 60% of human breast cancer (1,9), and dissecting Ras mitogenic

signaling from its invasive activity is, therefore, critical in understanding its promoting roles in breast cancer progression.

Estrogen receptor (ER α) is a nuclear transcription factor that regulates gene expression through binding to specific estrogen-responsive elements (ERE) on a target gene promoter. ER is proliferative in mammary tissues and consequently implicated in the etiology of breast cancer (10,11). Paradoxically, ER inhibits breast cancer *in vitro* invasion and *in vivo* metastasis (12, 13). Furthermore, higher levels of ER protein expression in human breast cancers correlate negatively with clinical metastasis but positively with the prognosis (14,15). These results together indicate that ER may use distinct pathways to stimulate breast cancer growth and to inhibit its invasion/metastasis.

ER may regulate breast cancer progression through crosstalk with Ras/mitogen-activated protein kinase (MAPK) signaling. Activation of the Ras/extracellular signal-regulated kinase (ERK) pathway downstream of growth factor receptors such as epidermal growth factor receptor (EGFR) increases both cell proliferation and invasion (4,16). However, forced Ras expression in ER⁺ breast cancer cells inhibits ER activity (17), and expression levels of EGFR and/or Ras proteins are higher in ER⁻ and/or metastatic breast cancers (18–20). These results suggest that ER and Ras cooperate in stimulating growth but antagonize each other in regulating invasion, and dissecting their crosstalk may contribute to understanding breast cancer progression.

MAPKs consist of ERK, c-Jun NH₂-terminal kinase, and p38 cascades. ERK activity is essential for Ras-induced proliferation/transformation (21,22), whereas p38 is inhibitory to Ras activity (23–25). In response to Ras-induced invasion, however, both ERK and p38 are required in this process (26). Although these different p38 effects on Ras-induced proliferation and invasion are not understood, some p38 family members may have distinct activity in these regulations. For example, p38 isoform proteins can be either cooperative or antagonistic in regulating gene expression and cell proliferation (27,28). Studies of p38 isoform-specific effects may contribute to understanding different Ras-p38 signaling interactions in regulating cell proliferation and invasion.

The p38 family consists of four isoforms: α , β , γ , and δ (29). p38 γ , also called stress-activated protein kinase 3 or ERK6, shares about 60% identity with p38 α and p38 β (30,31). In contrast to the ubiquitously expressed p38 α , p38 γ is only detectable in certain normal tissues (30,31). Recent studies, however, show that p38 γ protein is highly expressed in several human malignant cell lines (27,32,33), indicating its possible contributing role in cancer development. Indeed, our recent studies showed that Ras activates p38 γ by increasing its expression in rat intestinal epithelial (IEC-6) cells, and induced p38 γ , albeit nonmitogenic by itself, is required for Ras transformation (34). Specific roles of p38 γ in Ras-associated human cancers, however, remain to be established. Because Ras plays critical roles in increasing breast cancer progression (2), p38 γ may function as a Ras effector in this regulation.

Here, we sought to test the hypothesis that the Ras/p38 γ axis may exist in human breast cancer and thereby regulate breast cancer progression in coordination with ER protein. Our results revealed that in human breast cancer, Ras dictates p38 γ expression, which in turn mediates Ras nonmitogenic signaling to stimulate cell invasion. The invasive activity of the Ras/p38 γ axis, however, is antagonized by ER protein through *trans*-repression, and ER can additionally disrupt p38 γ protein through direct binding. These results suggest an invasive role of p38 γ in human breast cancer through integrating Ras proinvasive and ER anti-invasive signaling.

Materials and Methods

cDNA constructs

Recombinant adenovirus vector (Ad-Vect) and vector-containing oncogenic H-Ras (Leu⁶¹, L61) or dominant-negative H-Ras (Asn¹⁷, N17) were provided by Dennis Stacey (35). Adenovirus β -gal (ad- β -galactosidase), ad-MKK6, and ad-p38 γ were described elsewhere (32). Retroviral vector pLHCX (provided by Dan Mercola; ref. 36) was used to express p38 γ . To deplete endogenous protein, a retroviral vector pSUPER (pSR; ref. 37) was used (target sequence for sip38 γ was 5'-AAGGAGAT-CATGAAGGTGACG-3' and for siER α was 5'-AAGAGGAGGGAGAATGTT-GAA-3') using a luciferase gene sequence as a control (5'-CGTACGCGG-AATACTTCGA-3'; ref. 34). Human ER cDNA and ERE luciferase reporter (ERE-Luc, four ERE repeats upstream of a luciferase gene) were provided by David Shapiro (38). ER site-directed mutagenesis (T311A) was carried out by a PCR-based technique using the QuickChange mutagenesis kit (Stratagene, La Jolla, CA) with primer sequences as previously published (39). All mutations generated were confirmed by DNA digestion with restriction enzymes and DNA sequencing.

Cell culture, reagents, and Tet-inducible ER expression

MEM and other reagents for cell culture were purchased from Life Technologies (Gaithersburg, MD). Protein-Sepharose G beads and Cy3- and FITC-labeled second antibodies were obtained from Zymed (South San Francisco, CA). p38 isoform-specific antibodies were described previously (27,34), and their specificity was further confirmed by antibodies from BD Clontech (Palo Alto, CA). ERK1/2 and ER antibodies were from Santa Cruz Biotechnology (Santa Cruz, CA), whereas phosphorylated p38 (p-p38) and phosphorylated-ERK (p-ERK) antibodies were from Cell Signaling (Beverly, MA). Mouse pan-Ras antibody was purchased from Oncogene Research Products (San Diego, CA). Human breast cancer cell lines [ER⁺: MCF-7 and 47D; ER⁻: MDA-MB-231 (231) and MDA-MB-468 (468)] were obtained from the American Type Culture Collection (ATCC, Manassas, VA) and maintained in MEM containing 10% fetal bovine serum (FBS) and antibiotics at 37°C with 5% CO₂. The Tet-on expression system (T-Rex) was purchased from Invitrogen (San Diego, CA) and used to express human ER or ER/T311A in 231 cells as previously described (40). To evaluate effects of estrogens, cells were cultured with 10% charcoal-treated serum in phenol-free medium for 48 hours and then treated with 10 nmol/L estrogens for additional 24 hours.

Transfection, viral infection, reporter assays, and small interfering RNA experiments

ERE-Luc was transiently transfected by calcium phosphate into Tet-ER cells, which were then incubated with and without Tet for an additional 24 hours before the reporter assay using dual luciferase kit (41). For adenoviral infection, cells were infected with ad-vector or ad-L61/ad-N17 Ras in no serum medium for 4 hours and cultured in normal medium for 24 hours for protein expression and migration assays. In the case of retroviral infection, pSUPER (pSR-Lucif, pSR-sip38 γ , and pSR-siER) and pLHCX (Vector and p38 γ ; refs. 24,34) were transfected into Phoenix-Ampo retrovirus packaging cells (ATCC), and supernatants were used to infect target cells. Typically, cells were analyzed for protein or RNA expression at 24 to 48 hours.

Invasion, cell growth, and data analysis

Invasion assays were carried out using the BioCoat Matrigel Invasion Chamber (BD Biosciences, Bedford, MA) according to the manufacturer's instruction. Briefly, 2.5×10^5 cells in serum-free medium were plated into each chamber, which was placed into the six-well plate containing 20% FBS and incubated for 22 hours, at 37°C, 5% CO₂. Tetracycline (Tet) was added 2 hours later to both inserts and the bottom well to induce ER expression. For infection, cells are replated for invasion assay 24 hours after removal of virus-containing medium.

Following incubation, the noninvaded cells on the upper surface of membrane were removed, and the invaded cells on the lower surface were washed, fixed, and stained with crystal violet. The number of invaded cells from 13 to 20 representative fields was counted manually under a phase-contrast microscope, normalized to the respective control. To further verify these results, some crystal-stained attached cells were solubilized with 10% acetic acid and quantitated at 600 nm (42). For cell growth assay, cells were plated with or without Tet and pulsed labeled with [³H]thymidine, and the incorporated radioactivity was determined as previously described (41). All results were analyzed by Student's *t* test for statistically significant difference.

Immunostaining, immunoprecipitation, glutathione S-transferase pulldown, and cell fractionation/Western and Northern blot assays

For immunostaining, cells were plated on coverslips and fixed in 3.7% formaldehyde. After permeabilized in a buffer containing 0.5% Triton X-100 and 0.5% NP40, cells were blocked in 3% bovine serum albumin in PBS. A mouse monoclonal anti-flag (M2, Sigma, St. Louis, MO) and anti-ER (F10) antibody at 1:100 were used for flag-p38γ and ER staining, as previously described (34,40). For immunoprecipitation, cell lysates were incubated with a goat p38γ antibody, and the precipitates were examined for ER protein by Western blot. For glutathione S-transferase (GST) pulldown assay, GST, GST-ER and GST-ER/T311A proteins were expressed in *Escherichia coli* BL21 and purified using reduced glutathione-agarose beads (23).

To examine protein expression, cells were directly lysed in 1× loading buffer. After heating, samples were separated on SDS-PAGE and transferred to a nitrocellulose membrane for detection of the molecule of interest using enhanced chemiluminescence kit. For cell fractionations, cytoplasmic and nuclear fractions were prepared as previously described (43). For Northern blot, total RNA was prepared by TRIzol (Invitrogen) and separated by agarose gel. p38γ and p38α probe was prepared from their corresponding human cDNAs in pcDNA3 vector by enzymatic digestion and/or PCR (34). The probe was labeled with [³²P]dCTP using High Primer kit (Roche Molecular Biochemical, Indianapolis, IN). The Northern blots were standardized for equal RAN loading using the UV fluorescence of the 18S rRNA band from the same membrane.

Results

Ras positively regulates p38γ expression independent of ERK and/or p38 phosphorylation, whereas ER *trans*-suppresses Ras/p38γ in human breast cancer

Previous studies have shown that both transient and stable Ras expressions induce p38γ protein expression in IEC-6 cells (34). To determine whether Ras regulates p38γ expression in human breast cancer, adenovirus-mediated gene delivery was used to express oncogenic H-Ras (L61) and the dominant-negative mutant (N17; ref. 35) in ER⁺ 231 human breast cancer cells. Following infections, cells were analyzed for protein expression and phosphorylation. Consistent with the positive regulatory effect in IEC-6 cells (34), L61 increases whereas N17 decreases p38γ but not p38α protein expression compared with the control β-gal infection (Fig. 1A and B). In contrast to the p38γ induction in IEC-6 cells, Ras-regulating p38γ protein expression does not associate with significant alterations in ERK and/or p38 phosphorylations (Fig. 1A and B), in agreement with a recent observation of Ras activity dissociating with ERK phosphorylation in human breast cancer cells (44). These results reveal a determinant role of Ras activity in p38γ protein expression in human breast cancer independent of ERK/p38 phosphorylation and indicate a possible role of p38γ in Ras regulating breast cancer progression.

Ras is known to antagonize endogenous ER activity in MCF-7 human breast cancer cells (17,45). To explore whether exogenous ER conversely inhibits endogenous Ras activity, consequently leading to down-regulation of p38 γ protein expression, a Tet-on system was used to express ER protein by including its transcription-deficient mutant (T311A) for comparison (39,40). As expected, Tet-induced ER inhibits Ras protein expression (Fig. 1C). This effect requires ER transcription activity, as levels of Ras proteins remain constant with and without ER/T311A expression (Fig. 1D). Strikingly, ER down-regulating Ras protein expression does not result in persistent alterations in p-ERK and/or p-p38 proteins. Rather, it leads to a significant and sustained decrease in p38 γ protein expression (Fig. 1C and D). This effect is specific, as expression levels of p38 α protein remain constant with and without ER expression. Additional experiments with p-p38 immunoprecipitation and p38 γ /p38 α Western blotting showed that there was no detectable p38 γ phosphorylation in the absence and presence of ER expression (data not shown). These results, by both Ras expression and inhibition, revealed a unique signaling axis from Ras to p38 γ through regulating protein expression in human breast cancer. Because of the ER inhibitory role, the Ras/p38 γ axis may only act to regulate cell invasion/proliferation in ER⁻ breast cancer.

p38 γ transmits Ras nonmitogenic signaling to promote ER⁻ breast cancer invasion

The Ras determinant role in p38 γ protein expression prompted us next to explore if p38 γ acts as a Ras effector to execute its functions in regulating breast cancer invasion and growth. To assess roles of endogenous Ras in these regulations, ER⁻ 231 cells were infected with Ad- β -gal or Ad-Ras (N17), and its effects on cell invasion/growth were determined. Functional aspects of endogenous p38 γ in these processes, on the other hand, were analyzed by small interfering RNA-mediated p38 γ depletion (34). As expected, inhibition of Ras activity by N17 substantially decreases Matrigel cell invasion and DNA synthesis ($P < 0.001$ for both cases; Fig. 2A–C), indicating that Ras activity is required for both processes (26,46). In contrast to the Ras inhibition, p38 γ depletion only inhibits invasion ($P < 0.01$) without affecting thymidine incorporation ($P > 0.05$, Fig. 2A–C), suggesting a specific role of p38 γ in stimulating cell invasion. These results indicate that although p38 γ is downstream of Ras, they have common and distinct roles in regulating breast cancer invasion and growth.

Because N17 inhibits both p38 γ expression and cell invasion, whereas p38 γ is proinvasive, the p38 γ down-regulation may be required for N17 inhibiting invasion. If this is the case, p38 γ overexpression should rescue N17-induced invasion-suppression. To this end, p38 γ protein is overexpressed by a retroviral vector pLHCX in Ad-N17-infected cells, and its effects on cell invasion and DNA synthesis were determined. Consistent with the inhibitory effect of p38 γ depletion, higher levels of p38 γ protein increase invasion 2.8-fold ($P < 0.01$) without increasing thymidine uptake (Fig. 2D–F). More importantly, p38 γ overexpression almost completely overcomes the N17-induced inhibition of cell invasion ($P < 0.01$) without affecting DNA synthesis ($P > 0.05$), indicating that p38 γ only transmits the nonmitogenic Ras signaling to stimulate invasion. By both depletion and overexpression with and without Ras inhibition, these experiments show that p38 γ is necessary and sufficient for breast cancer invasion through transmitting Ras nonmitogenic invasive signaling.

ER regulates breast cancer growth and invasion by distinct pathways

The above results showed that Ras stimulates invasion and DNA synthesis by p38 γ -dependent and p38 γ -independent mechanisms, indicating Ras using distinct pathways to increase breast cancer invasion and growth. Because ER has well-established activities in breast cancer progression (47,48), we wished next to determine whether ER also regulates invasion and growth through distinct signaling. Previous studies have shown that ER suppresses breast cancer invasion by estrogen-dependent and estrogen-independent mechanisms (13), whereas estrogen can inhibit its own receptor expression (11). Consequently, experiments were done

in normal complete medium without estrogen addition to assess effects of ER protein expression.

Results in Fig. 3A and B showed that Tet induces a similar level of ER and ER/T311A protein expression, but ER transcription activity (ERE-Luc, ref. 38) decreases about 90% ($P < 0.01$) in mutant ER expressed cells. These results are consistent with the previous observation in endometrial adenocarcinoma cells (39). ER expression inhibited invasion $>80\%$ ($P < 0.01$), whereas the mutant ER had no substantial effects ($P > 0.05$, Fig. 3C, *left*). A similar result was also obtained when crystal violet-stained cells were quantitated at 600 nm (ref. 42; Fig. 3C, *right*). Although ER has well-established invasion-inhibitory activity (12, 13), these results suggest that ER may do so through transcriptional regulations. In contrast to the effects on cell invasion, ER and ER/T311A increased DNA synthesis similarly ($P > 0.05$; Fig. 3D), indicating that ER regulates invasion and DNA synthesis by distinct mechanisms. These results together show that both Ras and ER regulate breast cancer growth and invasion by distinct signal transduction pathways.

ER requires its nuclear localization to bind to p38 γ protein, leading to its specific down-regulation in the nuclear compartment

Although antagonistic effects between ER and Ras have been described (17,45), ER-suppressing p38 γ activity is novel and warrants further investigations. We sought, therefore, to further explore mechanistic aspects of signaling interactions between ER and p38 γ . Previous studies have shown that ER/T311A is predominantly localized in cytoplasm compared with nuclear wild-type ER protein (39), whereas p38 γ is both in the nucleus and the cytoplasm where it functionally interacts with other proteins (49). We wish next to determine if ER and ER/T311A are differently localized in human breast cancer cells, and whether their distributions affect p38 γ localization through direct binding, thereby additionally contributing to the p38 γ antagonizing activity.

Consistent with previous findings (39), immunostaining analyses showed that ER is mostly localized in the nucleus, whereas ER/T311A is predominantly in the area surrounding the nuclei after a 24-hour incubation with Tet (Fig. 4A). p38 γ signals are detected in the nucleus and cytoplasm in ER⁻ cells. In ER-expressed cells, however, the nuclear p38 γ signal seems negatively correlating with the ER intensity, indicating that nuclear ER is able to disrupt nuclear p38 γ protein, leading to its down-regulation in the nuclear compartment. This conclusion is further supported by the observation in ER/T311A cells in which the mutant ER is mostly outside the nucleus and in which there is no decrease in nuclear p38 γ signals (Fig. 4A).

To further confirm the decreased nuclear p38 γ protein concentration in response to ER expression, cellular fractionation and Western blotting experiments were done (43). Cytoplasmic actin (50) and both cytoplasmic and nuclear p38 α (51) serve as controls in this experiment. Results in Fig. 4B (*top*) showed that wild-type ER is mostly nuclear, whereas its mutant seems to be more abundant in the cytoplasmic fractions, generally consistent with the immunostaining results. Endogenous ER protein was also previously found to be both cytosolic and nuclear in ER⁺ MCF-7 cells by cell fractionation experiments (52). Of great interest, nuclear p38 γ protein is decreased upon ER induction, whereas its levels in cytoplasm remain constant. Because the mutant did not have this effect, Thr³¹¹-directed ER nuclear localization is apparently required for its antagonizing activity against p38 γ protein, leading to its specific down-regulation in the nuclear compartment.

To determine whether there is a physical interaction between ER and p38 γ , thereby leading to p38 γ down-regulation, immunoprecipitation and Western analyses were done. Results in Fig. 4B (*middle*) show that p38 γ binds to ER but not ER/T311A, although both proteins are expressed to a similar level from the input control, indicating ER requiring Thr³¹¹ to bind to

p38 γ protein. To further show whether p38 γ requires its phosphorylation to bind to ER, flag-tagged p38 γ and its nonphosphorable mutant (p38 γ /AGF) were expressed in 293T cells. Lysates derived were incubated with bacterially expressed GST, GST-ER, GST-ER/T311A proteins, and flag precipitates were examined for bound ER proteins. Results in Fig. 4B (bottom) show that ER, but not ER/T311A, binds to both p38 γ and p38 γ /AGF proteins, indicating that the direct ER-p38 γ binding requires ER Thr³¹¹ but not p38 γ phosphorylation. Together with the immunostaining and cell fractionation analysis, these results show that ER requires its nuclear localization to bind and thereby to disrupt nuclear p38 γ protein.

p38 γ overexpression increases invasion but not DNA synthesis in ER⁺ breast cancer cells

Results of ER-inhibiting p38 γ expression and p38 γ -stimulating ER⁻ cell invasion suggest that p38 γ may act downstream of ER to increase invasion. If this is the case, forced p38 γ expression in ER⁺ cells should overcome this genetic restriction and lead to an invasive response. To relate invasion-regulatory effects with cellular localizations, flag-p38 γ was overexpressed in ER⁺ cells (Tet-on), and its cellular distributions and activities on invasion were determined. Results in Fig. 4C show that in contrast to the endogenous p38 γ protein, overexpressed p38 γ is mostly in the cytoplasm, an effect similar to a recent observation in which overexpressed MK2 leads to its own relocalization (53). Although mechanisms are unclear at present, the cytoplasmic localization of exogenous p38 γ proteins in ER expressed cells further supports the notion that ER antagonizes nuclear p38 γ activity, leading to a reduced nuclear p38 γ concentration even under overexpressed conditions. In the cytoplasm, transfected p38 γ may have escaped the nuclear ER targeting as a result of their different localizations. As expected, overexpressed p38 γ led to an increased invasion (3.7-fold) but not thymidine incorporation in these ER⁺ cells (Fig. 4D). These results, thus, further establish the p38 γ -antagonizing activity of ER within the nuclear compartment and reveal that elevation of p38 γ protein concentrations by forced expression can also increase ER⁺ cell invasion.

ER negatively regulates p38 γ gene expression in a panel of human breast cancer cell lines

ER-inhibiting p38 γ protein expression dependent of ERE activity suggests that this regulation may be transcriptional. We would like next to determine if ER antagonizes p38 γ at RNA levels to further establish the *trans*-suppression mechanism. Total RNA in this case was prepared from Tet-on cells and analyzed for p38 γ RNA expression (41) by including p38 α as a control. As shown in Fig. 5A (left), Tet-induced ER decreases p38 γ but not p38 α RNA expression, thereby consolidating our conclusion of ER *trans*-suppressing p38 γ expression. To show if this *trans*-suppression is a general phenomenon, a panel of natural ER⁺ and ER⁻ breast cancer cell lines were analyzed for p38 γ RNA expression. ER protein expression has been previously confirmed in ER⁺ MCF-7 and T47D but not in ER⁻ 231 and 468 human breast cancer cells by Western analyses (40). Consistent with Tet-induced ER expression, higher levels of p38 γ RNA were detected in ER⁻ than in ER⁺ cells (Fig. 5A, right), ruling out cell type-specific effects of the p38 γ antagonistic activity.

To determine whether endogenous ER is also inhibitory to p38 γ expression, ER⁺ human breast cancer cells were depleted of ER proteins by retroviral-mediated gene silencing (34), and its effects on p38 γ expression were determined. Results in Fig. 5B show that ER depletion increases p38 γ expression, further establishing ER-inhibitory role in p38 γ expression. To assess if ligand can further enhance the *trans*-repression, estrogens were added after the retroviral infection and p38 γ protein contents were determined. Similar to the RNA up-regulation, ER depletion leads to an increase in p38 γ protein expression (Fig. 5C, left). Although estrogen treatment can further lower ER protein concentrations as a result of its well-known ER-depleting activity in MCF-7 cells (11), it fails to further increase p38 γ protein expression, indicating a ligand-independent p38 γ antagonistic activity of ER protein. Furthermore, addition of estrogens to steroid-depleted culture did not further increase the ER suppressive effects on

Ras/p38 γ protein expression in Tet-on 231 cells (Fig. 5C, *right*). These results together show that ER *trans*-suppresses p38 γ and/or Ras by ligand-independent mechanisms. Because loss of ER expression frequently occurs in late-stage breast cancer (54), demonstration of a p38 γ antagonistic activity of ER independent of ligand further highlights the role of p38 γ proinvasive activity in breast cancer progression.

Discussion

Uncontrolled proliferation and increased invasion are key characteristics of human cancer, and dissecting these two events is consequently critical for understanding and controlling malignant progression (1). Here, we showed that Ras increases breast cancer invasion and DNA synthesis by p38 γ -dependent and p38 γ -independent pathways. A specific role of p38 γ in increasing invasion independent of growth was shown by both its depletion and overexpression. Moreover, we provide evidence indicating that ER also inhibits invasion and increases DNA synthesis by distinct pathways, where p38 γ is specifically involved in regulating cell invasion. These results together show that there exist distinct signal transduction pathways for both Ras and ER to regulate breast cancer invasion and growth in which p38 γ specifically integrates their antagonistic activity to stimulate invasion (Fig. 6). Because invasion is a major characteristic of tumor but not normal cells (1), inhibiting p38 γ -dependent invasive pathways may be a novel strategy to control breast cancer progression.

Our results presented herein have revealed a unique Ras/p38 γ axis that acts to regulate breast cancer progression in coordination with ER protein. Here are the major characteristics of the Ras/p38 γ axis: (a) Ras, by both its activation and inhibition, primarily signals to regulate p38 γ expression independent of ERK/p38 phosphorylation; (b) Ras and its effector p38 γ have common and individual roles in stimulating invasion and growth; and finally, (c) the Ras/p38 γ axis is *trans*-repressed by ER and p38 γ is additionally down-regulated by ER in the nuclear compartment through direct binding. These results together suggest that p38 γ may serve as an integrating point for Ras proinvasive and ER anti-invasive activity through programmed protein expressions. In ER⁺ cells, invasion signals may be genetically silent in part through ER-induced suppression of Ras/p38 γ expression. In response to loss of ER expression during breast cancer progression, Ras/p38 γ will be activated by increased expression. The activated Ras will increase both proliferation and invasion, whereas the induced p38 γ will only confer proinvasive signaling, leading to increased breast cancer progression (Fig. 6).

The central mechanism by which ER antagonizes the Ras/p38 γ activity is likely through its nuclear localization that integrates transcriptional and posttranscriptional regulatory processes. This was shown by the fact that inhibition of ER nuclear localization through Thr³¹¹ mutation both reduces ERE-dependent transcription and disrupts its p38 γ binding activity. These results are consistent with the notion that nuclear localization is essential for activities of nuclear receptors (55). Although a decreased ERE by ER/T311A correlating with the loss of p38 γ binding activity suggests a role of this interaction in ER transcription, a decreased p38 γ expression in ER⁺ breast cancer cells may argue against its physiologic roles in regulating ER nuclear localization and ER activity. Additional studies, however, are needed to explore mechanisms by which the ER direct binding leads to a p38 γ down-regulation in the nuclear compartments.

Our results may offer an explanation about paradoxical ER effects in stimulating growth and in inhibiting invasion, which may otherwise antagonize each other in regulating breast cancer progression. Although loss of ER expression may reduce breast cancer growth, the resultant Ras activation can compensate for this effect by a proliferative response through increased expression (Fig. 6). This theory is supported by the observation of higher levels of Ras protein

expression in ER⁻ and/or metastatic human breast cancers (19,44,54,56,57). Because ER inactivation and Ras activation frequently occur in late stage of the disease (54,56), these results may suggest a stage-specific role of ER and Ras in promoting breast cancer progression through a coordinative signaling integration.

It is well accepted that Ras uses distinct effector pathways to induce malignant transformation/progression (2,46,58). Our studies here are the first to show the existence of such pathways under physiologic conditions in which induced p38 γ transmits Ras nonmitogenic signaling to amplify its invasion signals without affecting its proliferative activity. This specific pathway was first suggested by positive regulation of p38 γ expression through Ras activation and inactivation, then confirmed by ER inhibiting both Ras and p38 γ expression, and finally proven by p38 γ -mediated rescue of N17-induced invasion inhibition. Although ERK and/or p38 phosphorylations are involved in regulating breast cancer invasion (26,42), these effects could be secondary to altered cell proliferation. Here, we show that Ras dictates p38 γ expression without affecting ERK/p38 phosphorylation in human breast cancer, and p38 γ in turn increases invasion without affecting Ras mitogenic activity. These results together suggest that p38 γ act as a Ras effector to stimulate invasion in response to ER inactivation/Ras activation.

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Figure 1.

Ras positively regulates p38γ protein expression independent of ERK phosphorylation, whereas both Ras and p38γ protein expression is *trans*-suppressed by ER in human breast cancer cells. *A*, Ras inhibition suppresses p38γ expression without affecting ERK/p38 phosphorylation. ER⁻ 231 cells were infected with control adenovirus (Ad-β-gal) or virus expressing dominant-negative Ras (Ad-N17) for 4 hours and incubated for an additional 24 hours before analyzed for protein expression by Western blot. *B*, Ras activation induces p38γ expression without stimulating ERK phosphorylation. Cells were infected with Ad-β-gal or virus expressing oncogenic H-Ras (Ad-L61) as in (*A*) and examined for protein expression. *C* and *D*, ER inhibits Ras/p38γ protein expression dependent of its transcription activity. Cells were cultured with and without Tet for the indicated time to induce wild-type (*C*) and the mutant ER expression (*D*), and their effects on Ras/p38γ protein expression were examined by Western. Representative from three separate experiments.

Figure 2.

Endogenous Ras activity is required for both proliferation and invasion, whereas p38γ only transmits nonmitogenic Ras signaling to stimulate invasion. **A**, adenovirus-mediated N17 expression and small interfering RNA-induced p38γ depletion. Cells were infected with either adenovirus (Ad-β-gal or Ad-N17) or retrovirus (pSR-Lucif, control retrovirus; pSR-sip38γ, small interfering RNA retrovirus) and incubated for 24 hours before Western analysis. **B**, both Ras inhibition and p38γ depletion inhibit cell invasion. Cells were plated for invasion assay after infection, and invaded cells were counted and normalized to respective controls. *Columns*, mean of 15 fields ($P < 0.01$ for N17 and sip38γ versus their respective controls); *bars*, SD. Similar results were obtained from additional two experiments. **C**, Endogenous Ras but not p38γ is required for cell proliferation. Cells were infected as above, and cell proliferation was estimated by thymidine incorporation. *Columns*, mean of three experiments ($P < 0.05$ only for Ad-N17 versus Ad-Vect); *bars*, SD. **D**, N17 and p38γ overexpression. ER⁻ 231 cells were infected either with adenovirus (*Vect*, *N17*) and/or pLHCX retrovirus (*Vect*, *p38γ*) and analyzed for protein expression 24 hours later. **E**, p38γ overexpression rescues N17-mediated invasion inhibition. Cells were coinfecting with Ad-N17 with and without pLHCX-p38γ and assessed for cell invasion as described above. *Columns*, mean of 13 fields ($P < 0.01$ for p38γ versus Vect, N17 versus Vect, and for N17 versus p38γ + N17); *bars*, SD. Similar results are obtained from one additional experiment. **F**, high levels of p38γ protein expression do not overcome N17-induced growth inhibition. *Columns*, mean of three separate experiments ($P < 0.05$ for N17 versus Vect, but $P > 0.05$ for p38γ versus Vect and N17 + p38γ versus N17); *bars*, SD.

Figure 3.

ER inhibits invasion and stimulates growth by distinct mechanisms. *A*, ER protein expression in Tet-on cells. Tet-on 231 cells were incubated with and without Tet for 24 hours and examined for ER and ER/T311A protein expression. *B*, Thr³¹¹ is required for ER transcription activity. Cells were transfected with ERE-Luc, incubated with and without Tet for 24 hours, and assessed for luciferase activity. *C*, ER requires its transcription activity to inhibit invasion. Cells were cultured $\pm$ Tet for 22 hours in the invasion chamber for the assay (see Materials and Methods). *D*, ER does not require its transcription activity to stimulate DNA synthesis. Cells were cultured for 24 hours with or without Tet and cell growth was estimated by thymidine incorporation.

Figure 4.

ER antagonizes nuclear p38 γ activity through direct binding. **A**, ER is nuclear while ER/T311A is predominantly cytoplasmic. Cells were incubated for 24 hours ($\pm$ Tet) and double stained for ER (detected with Cy3-labeled second antibody) and for p38 γ expression (detected with FITC-labeled second antibody) with 4',6-diamidino-2-phenylindole (DAPI) staining as a control for nuclear signals. **B**, ER disrupts nuclear p38 γ protein through Thr³¹¹-dependent binding. Cells were prepared for nuclear and cytoplasmic fractions, which were analyzed for protein expression (*top*). p38 γ binding activity of ER was assessed by immunoprecipitation with a p38 γ antibody followed by Western analyses (*middle*). For pulldown assays (*bottom*), the purified GST, GST-ER, and GST-ER/T311A proteins were incubated with equal amounts of lysates prepared from 293T cells transfected with flag-p38 γ and flag-p38 γ /AGF, and the precipitates were analyzed for p38 γ -bound ER protein. **C**, overexpressed p38 γ is cytoplasmic in ER⁺ cells. ER⁺ 231 cells (Tet-on cells incubated with Tet for 24 hours) were infected with Ad-Vect (*Vect*) or Ad-p38 γ for 24 hours and stained with anti-flag antibody for transfected flag-p38 γ . **D**, higher levels of p38 γ proteins increase invasion (*right*) but not DNA synthesis (*left*) in ER⁺ breast cancer cells. Tet-on 231 cells were plated for invasion and thymidine incorporation after infection. *Right, columns*, mean from 18 fields of one representative experiment; *bars*, SD. Similar results were obtained from one additional experiment. *Left, columns*, mean of four experiments; *bars*, SD.

Figure 5.

ER negatively regulates p38γ expression in a panel of human breast cancer cell lines. *A*, ER⁺ phenotype correlates with lower levels of p38γ RNA expression. Total RNA was prepared, separated, and transferred to a nitrocellulose membrane that was hybridized with a specific radioactive-labeled probe for p38γ expression. *B*, ER depletion increases p38γ expression. ER⁺ cells were infected with pSR-Lucif (*Lucif*) or pSR-siER (*siER*) for 24 hours and examined for p38γ RNA (*left*) or protein (*right*) expression. *C*, ER negatively regulates p38γ expression by ligand-independent mechanisms. Cells in steroid-depleted medium were incubated for 24 hours (± 10 nmol/L E2) after the pSR infection (*left*) or in the presence or absence of Tet (*right*) and analyzed by Western analysis for protein expression.

Figure 6.

An experimental model illustrates that ER and Ras regulate breast cancer progression through distinct invasive and proliferative pathways where p38 γ only integrates their antagonistic activity to increase invasion. ER requires transcription activity to suppress Ras/p38 γ expression and to bind p38 γ protein, whereas it increases DNA synthesis by ERE-independent mechanisms. Ras, on the other hand, increases invasion and DNA synthesis through p38 γ -dependent and p38 γ -independent pathways in ER⁻ cells. Although loss of ER expression may reduce cell growth, resultant Ras activation will provide mitogenic signaling, leading to increased DNA synthesis [and possible cell proliferation (?)]. Induced p38 γ following ER inactivation and/or Ras activation will specifically mediate Ras nonmitogenic signaling to stimulate invasion. Through these two mechanisms, Ras/p38 γ activation will lead to an increased breast cancer progression. This model suggests a feasibility to individually target invasive and proliferative signal transduction pathways in human breast cancers.