Phosphorylation at serines 104 and 106 by Erk1/2 MAPK is important for estrogen receptor- α activity

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Abstract

Phosphorylation of estrogen receptor- α (ER α) at specific residues in transcription activation function 1 (AF-1) can stimulate ER α activity in a ligand-independent manner. This has led to the proposal that AF-1 phosphorylation and the consequent increase in ER α activity could contribute to resistance to endocrine therapies in breast cancer patients. Previous studies have shown that serine 118 (S118) in AF-1 is phosphorylated by extracellular signal-regulated kinases 1 and 2 (Erk1/2) mitogen-activated protein kinase (MAPK) in a ligand-independent manner. Here, we show that serines 104 (S104) and 106 (S106) are also phosphorylated by MAPK *in vitro* and upon stimulation of MAPK activity *in vivo*. Phosphorylation of S104 and S106 can be inhibited by the MAP-erk kinase (MEK)1/2 inhibitor U0126 and by expression of kinase-dead Raf1. Further, we show that, although S118 is important for the stimulation of ER α activity by the selective ER modulator 4-hydroxytamoxifen (OHT), S104 and S106 are also required for the agonist activity of OHT. Acidic amino acid substitution of S104 or S106 stimulates ER α activity to a greater extent than the equivalent substitution at S118, suggesting that phosphorylation at S104 and S106 is important for ER α activity. Collectively, these data indicate that the MAPK stimulation of ER α activity involves the phosphorylation not only of S118 but also of S104 and S106, and that MAPK-mediated hyperphosphorylation of ER α at these sites may contribute to resistance to tamoxifen in breast cancer.

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Introduction

Estrogen receptor- α (ER α) is a member of the nuclear receptor superfamily of ligand-regulated transcription factors, and is required for development and maintenance of the female and male reproductive systems, bone, and cardiovascular system, as well as being significant for certain brain functions (Couse & Korach 1999). Additionally, estrogens stimulate breast cancer progression and $\sim 70-80\%$ of breast cancers express ER α . The presence of ERa correlates with likelihood of response to anti-estrogens, such as tamoxifen (Osborne 1998), and to aromatase inhibitors that inhibit estrogen biosynthesis (Henderson & Piccart-Gebhart 2005), demonstrating the importance of ER α in breast cancer progression. However, a proportion of patients with ERαpositive disease do not respond to endocrine therapy. Further, of those patients who respond, many who initially present with localized disease, and all with advanced breast cancer, relapse. Most resistant tumors remain ERα-positive and frequently respond to alternative antiestrogen treatment, indicative of a continued role for ERa in breast cancer cell proliferation (Ali & Coombes 2002). Augmentation of ER α activity in the presence of estrogen or anti-estrogens, and/or ligand-independent activation of ERα, could provide an important mechanism for resistance to anti-estrogen therapies. In agreement with

this possibility, a recent study shows that levels of S118 phosphorylation are increased following breast cancer recurrence following tamoxifen treatment (Sarwar *et al.* 2006).

ERα regulates gene expression through direct binding to estrogen-response elements within the promoters of estrogen-regulated genes, or by recruitment to DNA through interaction with other transcription factors (Gronemeyer 1991, Bjornstrom & Sjoberg 2005). Transcriptional regulation by ERα is partly mediated by transcription activation function, AF-2, which is intrinsic to the ligand-binding domain (LBD). Estrogen binding to the LBD results in a conformational change that allows recruitment of coactivators to the LBD (Brzozowski et al. 1997, Shiau et al. 1998, Glass & Rosenfeld 2000). A second transcription activation function, AF-1, is encoded within the N-terminal 180 amino acids. AF-1 activity is promoter and cellular context dependent (Tzukerman et al. 1994), and can be influenced by the expression profile of specific coactivators (Smith et al. 1997). Although there is evidence to suggest that phosphorylation of AF-1 alone can enhance activity (Bunone et al. 1996, Ignar-Trowbridge et al. 1996), ligand binding to AF-2 can also induce AF-1 activity (Metzger et al. 1992). This suggests that, in a basal state, AF-1 activity is blocked by AF-2. Finally, AF-1 and AF-2 can act synergistically in a promoter- and cell-specific manner (Gronemeyer 1991, Tsai & O'Malley 1994, Beato *et al.* 1995).

Tamoxifen is an example of a mixed estrogen agonist/antagonist, also termed selective ER modulators, which has been proposed to act by inhibiting the LBD/AF-2, whilst allowing activation of AF-1 (Berry et al. 1990, McDonnell et al. 1995, Brzozowski et al. 1997, Shiau et al. 1998). Structural studies have indicated that the inhibition of the LBD/AF-2 in the tamoxifen-bound $ER\alpha$ is due to the positioning of helix 12 in the hydrophobic cleft to which transcriptional coactivators, such as the p160 family members, are recruited, thereby preventing coactivator recruitment. The same cleft can also be involved in corepressor binding, and tamoxifen has been suggested to allow corepressor recruitment to the LBD (Nettles & Greene 2005). The nature of AF-1 activation by ligands such as tamoxifen, and in the absence of ligand, is unknown, but may involve phosphorylation of ER α . Previous studies have shown that in ovariectomized mice, uterine proliferation (which is normally estrogen- dependent) can be mediated by growth factors such as epidermal growth factor (EGF), and that the EGF-mediated uterine stimulation is ERa dependent (Ignar-Trowbridge et al. 1992, Curtis et al. 1996). ERa may become phosphorylated in response to EGF and other growth factors, resulting in ligand-independent ERα activation (Ignar-Trowbridge et al. 1993).

Phosphorylation site mapping has demonstrated that serine 118 (S118) in AF-1 is phosphorylated by Erk1/2 MAPK, resulting in stimulation of ERα activity (Kato et al. 1995, Bunone et al. 1996), whilst phosphorylation of serine 167 (S167) in AF-1 by AKT and p90RSK also stimulates ERa activity (Joel et al. 1998a, Campbell et al. 2001). The ERα AF-1 is additionally phosphorylated at serines 104 (S104) and/or 106 (S106) (Le Goff et al. 1994). Phosphorylation at these sites by Cdk2 (Trowbridge et al. 1997, Rogatsky et al. 1999) and GSK3 (Medunjanin et al. 2005) has been reported, and our own data have previously indicated that these sites might also be targets for MAPK (Chen et al. 2002). Increased ERα activity due to AF-1 phosphorylation mediated by MAPK and/or other kinases could therefore be at least partly responsible for resistance to endocrine therapy in breast cancer. This hypothesis is supported by findings that stimulation of growth factor-regulated pathways can augment the agonist properties of tamoxifen (Smith 1998) and inhibit tamoxifen-induced growth suppression and apoptosis in breast cancer cells (Benz et al. 1993, Pietras et al. 1995, Campbell et al. 2001). Here, we investigate phosphorylation at serines 104 and 106 by Erk2 MAPK in vitro and in response to MAPK activity *in vivo*, and determine their contribution to ERα activity. Transcription reporter assays using ERα phosphorylation site mutants suggest that phosphorylation at all three sites

contribute significantly to AF-1 activity, and in particular are required for tamoxifen to act as an ER α agonist.

Materials and methods

Plasmids

The expression vectors pSG5-HEG0 and pSG5-HE15, encoding full length and AF-2-truncated human ERα (ERα-ΔLBD) respectively have previously been described (Tora et al. 1989a). pGEX1λT-ERα and pGEX1λT-ERα-ΔLBD were generated by cloning ERα encoding EcoRI fragments from pSG5 into pGEX1λT (GE Healthcare UK Ltd, Little Chalfont, UK). ERα mutants were generated by site-directed mutagenesis, according to manufacturer's protocols (Stratagene, La Jolla, CA, USA). pEF-RasV12, expression vector for constitutively active Ha-Ras, was kindly provided by Dr R Treisman (LRI, London, UK). pCMV-Raf CAAX and pCMV-Raf S621A were purchased from Clontech (Saint-Germaine-en-Laye, France).

Antibodies

Peptides corresponding to amino acids 97-112 of human ERα, having the sequence [C]-FPPLNSV^P-SPSPLMLLH (phospho-S104) or [C]-FPPLNSVSP^PSP-LMLLH (phospho-S106), were used to generate rabbit polyclonal antisera, as described (Chen *et al.* 2002). Antibodies specific for ERα phosphorylated at S118 (16J4, New England Biolabs, Hitchin, UK), ERα (6F11, Novocastra Laboratories, UK), MAPK (sc-93, Santa Cruz Biotechnology, Heidelberg, Germany) and P-MAPK (New England Biolabs) were obtained commercially. Horseradish peroxidase-conjugated goat anti-rabbit and anti-mouse immunoglobulins were purchased from DAKO Ltd, Ely, UK.

Immunoblot analysis

Cells were grown in Dulbecco's modified Eagle's medium (DMEM) lacking phenol red and supplemented with 10% dextran-coated charcoal-stripped fetal calf serum (DSS) for 3 days prior to plating in six-well plates at 300 000 cells/well. Cells were transfected using Fugene 6 (Millipore, Wafford, UK), with 100 ng pSG5 empty vector or ER α expression construct, 5 ng empty vector or Ras/Raf expression construct, and 500 ng pBS+ carrier DNA, as appropriate. After 48 h the cells were treated with ethanol carrier, estradiol (E2; 10 nM), 4-hydroxytamoxifen (OHT; 100 nM) or ICI182,780 (ICI; 100 nM) 30 min prior to harvest, phorbol 12-myristate 13 acetate (PMA; 100 nM) was added 15 min, and U0126 (10 μ M) was added 60 min, prior to harvesting, as appropriate. Cells were washed and harvested

directly into Laemmli buffer and immunoblotting was performed, as described (Sarwar et al. 2006). For peptide competition experiments, the primary antibodies were pre-incubated with $10~\mu g/ml$ of the appropriate peptide (Chen et al. 2002). Quantitation of phospho-ER α signal was performed by densitometry of scanned films using ImageQuant software, and expressed relative to the respective total ER level.

Kinase assays

The glutathione-S-transferase (GST) fusion proteins, GST-ERα and GST-ERα-ΔLBD were expressed in Escherichia coli (Rosetta strain, Merck Chemicals Ltd, Nottingham, UK) following induction with 0. 1 mM isopropyl-beta-D-thiogalactopyranoside (IPTG) for 3 h at 25 °C. Cells (50 ml) were lysed with 10 ml XTractor buffer (BD Biosciences, Oxford, UK) containing 1 mM dithiothreitol (DTT) and protease inhibitors, and the lysates cleared by high-speed centrifugation. The lysates were incubated with glutathione-sepharose beads at 4 °C for 1 h, and the beads were washed thrice with TBS containing 1 mM DTT and protease inhibitors. GST-ERα beads were resuspended in 1× kinase buffer, 200 μ M ATP, $\pm E_2$ (10 nM) and kinase (Erk2 and GSK3, New England Biolabs, UK; Cdk2/cyclin A, Cdk2/cyclin E, Cdk4/cyclin D1, Cdk7/cyclin H/MAT1, AKT1, and AKT3; New England Biolabs), according to manufacturer's instructions. For radioactive kinase assays, 50 μM cold ATP and 10 μM ³²PγATP (Amersham) was used. Reactions were incubated at 30 °C for 30 min and processed by SDS-PAGE followed by autoradiography or immunoblot.

Reporter assays

Cells were grown in DMEM lacking phenol red and supplemented with 10% DSS for 3 days prior to plating in 24-well plates at 50 000 cells/well. Cells were transfected using Fugene 6 (Roche), with 100 ng pERE3-TATA-luc and pRL-TK reporters, 10 ng pSG5 empty vector or ERa expression construct, 50 ng empty vector or Ras/Raf expression vector, and 500 ng pBS+ carrier DNA. After 4 h, the medium was replaced with fresh media containing ethanol carrier, E2, OHT or ICI 182 780, at concentrations indicated in figures. After a further 20 h, the cells were harvested and luciferase levels determined using Dual-Glo reagents (Promega). For experiments in which U0126 was used, 10 nM U0126 was added 1 h prior to the addition of ligands and the cells were harvested after a further 7 h. Firefly luciferase levels were corrected for transfection efficiency using corresponding renilla luciferase levels. The activity for wild-type ERa in the absence of ligand was taken as one, with all other activities shown relative to this. All experiments were

independently repeated at least four times, and the data presented as mean values with s.E.M. error bars.

Results

Antisera display specificity for ER α phosphorylated at S104 and S106

Serines 104 and/or 106 have been shown to be phosphorylated by Cdk2/cyclin A and Cdk2/cyclin E (Trowbridge et al. 1997, Rogatsky et al. 1999), and GSK3 (Medunjanin et al. 2005). To further characterize ERa phosphorylation at these residues, we generated rabbit antisera specific for phosphorylation at S104 (α-PS104) or at S106 (α-PS106). S104 phosphorylation was stimulated following treatment of ERa-transfected COS-1 cells with E_2 and PMA (Fig. 1A). ER α mutants in which S104 was substituted by alanine were not detected by α -PS104, whilst mutation of S106 (106A) reduced but did not abolish S104 phosphorylation. S106 phosphorylation was also stimulated by E₂ and PMA, and blocked by mutants of S106 but not S104. Competition with phosphorylated peptides further confirmed that \alpha-PS104 and \alpha-PS106 are specific for ERα phosphorylated at S104 and S106 respectively. Detection of phopho-S104 ERα and phospho-S106 ER α were blocked by pre-incubation of the respective antiserum with a 100-fold excess of peptides containing a phosphorylated S104 (PS104 and PS104/6), and phosphorylated S106 (PS106 and PS104/6) respectively.

Investigation of phosphorylation using mutants in which S104, S106, and S118 were substituted by alanine or glutamic acid, either singly or together, showed that S104 phosphorylation was influenced by the status of S106 and S118 (Fig. 1B). This result could potentially be explained by a reduced efficacy of α-PS104 binding when S106 or S118 are not phosphorylated, but this is not supported by the fact that α-PS104 binding was competed similarly by both the PS104 and PS104/6 peptides (Fig. 1A). Hence, phosphorylation at S106 and S118 may be important for subsequent phosphorylation of S104. S106 phosphorylation was lower in the case of S118A, but not S104A, suggesting a role for S118 in S106 phosphorylation. In the case of S118, the data suggest that S104, but not S106 influences S118 phosphorylation. Together, these findings are indicative of crosstalk between S104, S106, and S118, which regulates phosphorylation at these sites.

Phosphorylation at S104 and S106 is induced by $ER\alpha$ ligands and PMA, and by activated Raf/Ras

S118 phosphorylation is stimulated by E_2 , as well as anti-estrogens (Joel *et al.* 1995, 1998*b*) and can be mediated

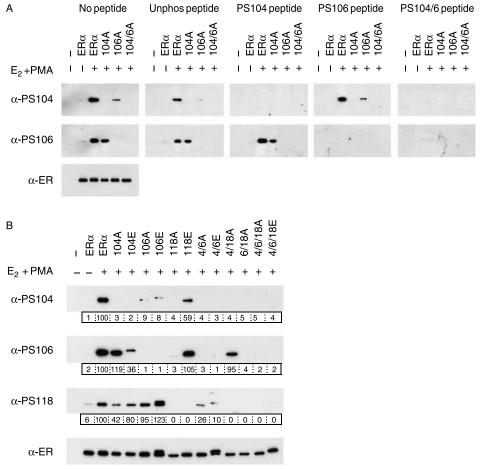


Figure 1 Characterization of phospho-specific antisera by peptide competition and phosphorylation site substitutions. Lysates prepared from COS-1 cells transiently transfected with an empty expression vector (-), or expression vectors for wild-type ERα or ERα in which S104, S106, and/or S118 had been substituted by alanine (A) or glutamic acid (E), as indicated, were immunoblotted using antibodies for total ERα (α -ER), or for ERα phosphorylated at S104 (α -PS104) or S106 (α -PS106). Cells were treated with ethanol solvent (-), or 17β-estradiol (E2; 10 nM) and 12-tetradecanoylphorbol-13-acetate (PMA; 100 nM), for 30 min prior to harvesting. (A) Replicate blots were incubated with primary antibody (no peptide), or antibody that had been pre-incubated with a 100-fold excess (10 μg/ml) of a peptide encompassing the ERα phosphorylation site; either unphosphorylated (unphos) or phosphorylated (PS104, PS106, or dual PS104/6) versions, as indicated. (B) Lysates were additionally immunoblotted using antibody for ERα phosphorylated at S118 (α -PS118). Levels of phospho-ERα were quantitated in relation to the respective total ERα level (boxed, below each immunoblot).

by Cdk7 (Chen *et al.* 2000, 2002). Further, S118 can be phosphorylated by MAPK in response to EGF and PMA (Kato *et al.* 1995, Bunone *et al.* 1996, Chen *et al.* 2002). Phosphorylation of S104 and S106 was also stimulated by E₂, OHT, ICI, and PMA (Fig. 2A). In the case of PMA, this phosphorylation was decreased by the MEK1/2 inhibitor U0126, and corresponds to a decrease in activated Erk1/2 MAPK. The phosphorylation induced by ligands was mostly unaffected by the addition of U0126, except that of S104 in response to OHT and ICI, which were reduced. Co-expression of ERα with constitutively active Ha-Ras (Ras-V12) or Raf1

(Raf-CAAX) resulted in enhanced phosphorylation of S104, S106, and S118, whereas the kinase-dead mutant of Raf1 (Raf-S621A) did not (Fig. 2B). Together, these data indicate that S104, S106, and S118 can be phosphorylated by Erk1/2 MAPK in response to activation of the upstream pathways, and also though MAPK-independent pathways upon ligand binding.

In agreement with the above findings, purified Erk2 readily phosphorylated GST-ERα at S104, S106, and S118 (Fig. 3A). Cdk2 also phosphorylated S104 and S118, but did not appear to phosphorylate S106. GSK3 also phosphorylated S104, with longer

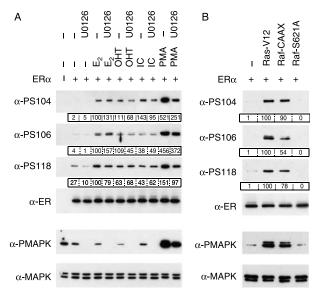


Figure 2 Phosphorylation of S104 and S106 is stimulated by ERα ligands and by activators of MAPK. Immunoblots of lysates prepared from COS-1 cells transfected with empty expression vector (—) or expression vector for ERα, were performed as for Fig. 1. Lysates were additionally immunoblotted using antibodies for total (α -MAPK) and phosphorylated (α -PMAPK) Erk1/2 MAPK. Levels of phospho-ERα were quantitated in relation to the respective total ERα level (boxed, below each immunoblot). (A) Cells were pre-incubated with U0126 (10 μM) for 1 h, followed by the addition of E₂ (10 nM), 4-hydroxytamoxifen (OHT, 100 nM), ICI 182 780 (ICI; 100 nM) or PMA (100 nM), as indicated, and cells harvested 30 min later. (B) Cells were transfected with expression vector for ERα, together with expression vectors for Ras-V12, Raf-CAAX or Raf-S621A, as indicated.

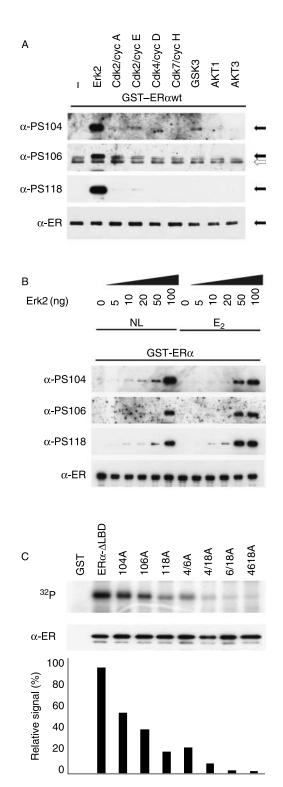
exposures showing phosphorylation at S106, but not S118. However, Cdk2 and GSK3-mediated ERa phosphorylation appeared to be less significant than that mediated by Erk2. Similar results were obtained using recombinant ERα purified from SF9 insect cells and transiently transfected ERa in crude COS-1 extracts (data not shown). The presence of estrogen provided a moderate enhancement of Erk2-dependent phosphorylation (Fig. 3B), but confirmed that ERα can be phosphorylated by MAPK in the absence of ligand. To investigate the relative levels of phosphorylation at these sites, GST-ERα-ΔLBD and alanine substitution mutants were incubated with Erk2 in the presence of ³²P-γATP (Fig. 3C). Substitution of S104, S106, or S118 by alanine resulted in significant reductions in Erk2 phosphorylation, to ~ 40 , 60, and 80% respectively, relative to wild-type $ER\alpha$, with negligible detectable phosphorylation when all three sites were mutated, suggesting that S104, S106, and S118 are the only significant Erk2 phosphorylation sites within amino acids 1-281 of ERa. These data further support the possibility that phosphorylation at each of these sites influences phosphorylation at the other sites.

S104, S106, and S118 are important for ERα activity

In reporter gene assays, substitution of S104, S106, and S118 by alanine (4/6/18A) resulted in a reduction in ER α activity in the presence of E₂ by nearly 50%, whilst substitution with glutamic or aspartic acid, to 'mimic' phosphorylation at these sites, stimulated ERα activity two- to threefold (Fig. 4A). ERa activity for the alanine mutant in the presence of OHT was reduced by > 80%, and was two- to threefold higher for the 4/6/18E and 4/6/18D mutants. Significant ERα activation (sevenfold relative to wild-type ERα) was also observed in the absence of ligand for 4/6/18E and 4/6/18D. These data suggest that phosphorylation at one or more of these residues is critical for AF-1 activity and for the agonist activity of OHT. Western analysis of total ERa levels indicated relatively equal levels of expression across the mutants and across different ligand treatments. ICI is generally thought to decrease $ER\alpha$ levels, but in many cases a change in solubility is mistaken for degradation (Lipfert et al. 2005); the assays here have measured total ERα levels.

To examine whether acidic substitutions in ER α resulted in ligand hypersensitivity, we examined ER α activity by reporter assay in the presence of increasing concentrations of estrogen and OHT (Fig. 4B). The acidic ER α mutant displayed an enhanced activity, of similar magnitude compared with wild-type ER α , at all ligand concentrations. Hence, although the acidic mutant could yield equivalent levels of activity at lower ligand concentrations compared with that of wild-type ER α at higher ligand concentrations (compare 4/6/18E at 0·1 nM E₂ with wt at 1 nM E₂, and 4/6/18E at 1 mM OHT with wt at 100 mM OHT), there was no evidence for enhanced ligand sensitivity.

ERα AF-1 is only weakly active in HeLa cells, with the majority of ERα activity in HeLa cells being due to AF-2 (Bocquel et al. 1989). Further, the lack of AF-1 activity correlates with lack of agonist activity for OHT in HeLa cells (Bocquel et al. 1989, Berry et al. 1990). Triple acidic substitution stimulated ERa activity in the absence of ligand, as well as in the presence of E₂ (Fig. 4C). Activity in the presence of OHT was also increased, but was not significantly greater than that obtained in the absence of ligand. Hence, acidic substitutions in AF-1 enhanced ERα activity in HeLa cells in a similar manner to COS-1 cells, but did not activate an OHT response. Finally, in the estrogen responsive and ERα-positive MCF7 breast cancer cell line, transfection of ERα stimulated reporter gene activity over and above the activity observed for endogenous ERα (Fig. 4D). Reporter gene activity for the glutamic and aspartic acid mutants was significantly greater in the absence of ligand or in the presence of E₂ or OHT, compared with the activities obtained for wild-type ERα, generally following what was seen using COS-1 cells.



Phosphorylation at S104, S106, and S118 contributes to $ER\alpha$ activity

The above data demonstrate that serines 104, 106, and 118 are critical for AF-1 activity, and likely for AF-1 cooperativity with AF-2. In order to define the relative importance of these sites, single and double mutants were examined for their effect on ER α activity (Fig. 5). Substitution of S118 by alanine reduced OHT-stimulated ERa activity. Alanine substitution of 104A or 106A had little effect on OHT stimulation of ERα, although ERα-104A/106A was significantly less active than wild-type ERa. Interestingly, S104 or S106 mutations together with mutation of S118 reduced ERα activity more potently than the S118 substitution alone. On the basis of these findings, the order of importance of the three serines appears to be S118> S104>S106. Although mutation of S106 to alanine had very little effect on ER α activity, its substitution by glutamic acid resulted in the greatest stimulation in ERa activity, with the order of activity for the three mutants being S106>S104>S118, opposite to the relative activities observed with the alanine substitutions. Nevertheless, in agreement with the findings from the alanine mutants, multiple substitutions increased ERα activity correspondingly, again suggesting that all three phosphorylation sites are important for AF-1 activity.

Activated Ras/Raf enhances ERa activity

In order to confirm the role of MAPK in the phosphorylation of S104 and S106, NIH-3T3 cells were co-transfected with reporter constructs and ER α , together with expression vectors encoding Ras or Raf (Fig. 6A). ER α activity was significantly enhanced by constitutively active Ha-Ras (Ras-V12) and Raf1 (Raf-CAAX), and was inhibited by a kinase-dead mutant of Raf1 (Raf-S621A). By contrast, ER α -104A/106A/118A activity was not stimulated by Ras-V12 or Raf-CAAX. U0126 did not significantly inhibit the activities of

Figure 3 S104 and S106 are phosphorylated by Erk2 in vitro. (A) Purified GST-ERα was incubated with a panel of purified kinases, as indicated, according to manufacturer's instructions, followed by immunoblotting with phospho-specific antibodies (α -PS104, α -PS106, and α -PS118) or α -ER. The filled arrows indicate the position of GST-ERa, and the open arrow indicates a non-specific product seen with α-PS106 antisera. (B) Purified GST-ERα was incubated with increasing amounts of purified Erk2 MAPK (0, 5, 10, 20, 50, and 100 ng) in the absence of ligand (NL) or in the presence of E2 (10 nM), followed by immunoblotting as before. (C) Purified GST, or wild-type GSαT-ERα-ΔLBD (ERα-ΔLBD) or GST-ERα-ΔLBD in which S104, S106, and/or S118 had been substituted by alanine (A), as indicated, were incubated with Erk2 in the presence of ³²P-γATP, followed by SDS-PAGE and autoradiography of the dried gel. Immunoblotting a duplicate gel with α -ER was used to determine the relative levels of each mutant. The bar chart shows quantification of each 32P signal relative to the respective total GST-ER α - Δ LBD level.

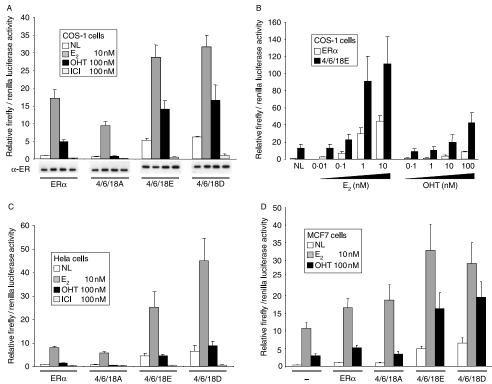


Figure 4 Effect of phosphorylation site mutations on ERα activity. (A and B) COS-1, (C) HeLa, and (D) MCF7 cells were co-transfected with the ERE-3-TATA-firefly luciferase reporter gene, a renilla luciferase control reporter gene, and expression vectors encoding wild-type ERα (ERα) or ERα in which S104, S106 and S118 were substituted by alanine (4/6/18A), glutamic acid (4/6/18E) or aspartic acid (4/6/18D), as indicated. Cells were treated with ethanol solvent (no ligand; NL), E₂ (10 nM), OHT (100 nM) or ICI (100 nM), as indicated, except (B) where E₂ was added at 0.01, 0.1, 1, and 10 nM and OHT at 0.1, 1, 10, or 10 nM. The cells were harvested for luciferase assays 20 h later. Results are presented as relative ratios of firefly to renilla control luciferase activities, as described in Materials and methods. (A) A parallel transfection series was assayed for ERα expression by immunoblot (α -ER). (D) MCF7 cells were additionally transfected with an empty vector control (-) in order to determine the contribution of endogenous ERα.

ERα-104A/106A/118A or 104E/106E/118E, but did inhibit the activities of wt ERα and those encoding individual alanine mutants (Fig. 6B). Similar results were obtained by treatment with another MEK1/2 inhibitor, PD98059 (data not shown). These data suggest that the activation of the Ras/Raf/MEK/MAPK signal transduction pathway results in ERα activation through phosphorylation at S104, S106, and S118.

Discussion

Phosphorylation at serine 118 in the AF-1 domain of ER α has been demonstrated by us and other researchers to be important for ER α activity (Ali *et al.* 1993, Le Goff *et al.* 1994, Kato *et al.* 1995, Bunone *et al.* 1996, Chen *et al.* 2000, 2002). Phosphorylation of serines 104

and 106 has also been shown to augment $ER\alpha$ activity (Le Goff *et al.* 1994), but their phosphorylation and function have not been investigated thoroughly.

In order to investigate phosphorylation of these residues in greater detail, we generated phosphorylation site-specific antisera. The antisera demonstrated appropriate specificity for ERα phosphorylated at S104 or S106, determined using multiple point mutants of S104 and S106 and peptide competition of antibody binding, and confirms that both S104 and S106 of ERα can be phosphorylated. Interestingly, phosphorylation at S104 appeared to be substantially reduced by substitutions at S106 and S118, and S106 phosphorylation reduced by S104E and S118A. S118 phosphorylation was also somewhat reduced by S104 substitutions. The epitopes for the α-PS104 and α-PS106 antisera may overlap, but the peptide competition studies of antibody specificity go some way to suggest that these antisera are not influenced by the

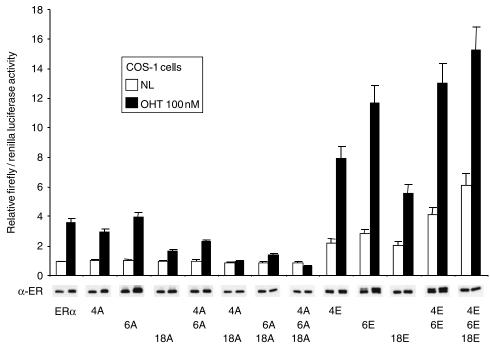


Figure 5 Effect of individual site mutations on ER α activity. COS-1 cells were co-transfected with ERE-3-TATA-luc, pRL-TK, and the wild-type ER α expression vector (ER α) or versions with alanine or glutamic acid substitutions, as indicated. Cells were treated and luciferase assays carried out as for Fig. 4. A parallel transfection series was assayed for ER α expression by immunoblot (α -ER).

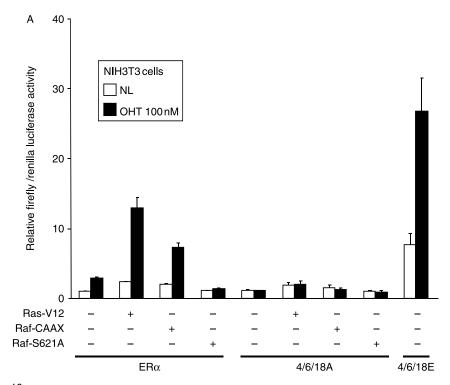
phosphorylation status at the respective adjacent site. Thus, in general, phosphorylation at these sites appears to be partly interdependent. We cannot, of course, rule out the possibility that these substitutions simply affect an $ER\alpha$ conformation required for efficient phosphorylation.

Previous studies have indicated that the agonist activity of tamoxifen (OHT) is due to activation of AF-1, whilst AF-2 function is inhibited (Tora et al. 1989b, Berry et al. 1990). In reporter gene assays using OHT, alanine substitution of S104, S106, or S118 inhibited ERα activity, with the greatest inhibition being observed with substitution of all three residues. Similarly, substitution of the individual sites by glutamic acid augmented ERα activity, with the greatest activity being seen for the triple mutant. Additionally, these sites appear to be important for estrogen-induced and ligandindependent ER α activities. These data suggest that phosphorylation at some or all three sites is critical for AF-1 activity and for the agonist action of OHT. Substitution of S118 to alanine resulted in a greater loss of function compared with that of S104 and S106, but glutamic acid substitutions suggested that achieving a high level of phosphorylation at S104 and S106 might have a greater impact upon ER\alpha function than that at S118. These apparently contradictory results may be explained by functionally stronger S104 and S106 sites being phosphorylated at lower levels in COS-1 cells, relative to that at S118. It is additionally difficult to assign

levels of importance to these sites, since immunoblot analysis of ER α substitution mutants suggested that S104, S106, and S118 phosphorylation is partly interdependent. Our data instead suggest that these three sites, together, comprise a complex phosphorylated domain, involved in augmenting ER α activity.

Reporter gene assays indicated that the activity of the ER α -4/6/18E mutant (and 4/6/18D mutant) in the absence of ligand was at a level similar to that seen for wildtype ERα in the presence of OHT, suggesting that OHTinduced phosphorylation at these residues may be sufficient for AF-1 activity. However, the further stimulation of ER α -4/6/18E activity by tamoxifen in COS-1 cells, suggests that the agonist activity of OHT is not solely due to stimulation of phosphorylation at these sites. Indeed, acidic substitutions did not lead to a restoration of OHT-induced ERα activity in HeLa cells, in which ERα AF-1 activity is known to be low and associated with little or no stimulation by OHT (Tora et al. 1989b, Berry et al. 1990). These data further support the view that the agonist activity of OHT is not due simply to stimulation of AF-1 phosphorylation. Thus, although OHT agonist activity is dependent upon (and induces) phosphorylation, other factors - possibly cell context-dependent expression of coactivators and corepressors (Shang & Brown 2002) – are also likely to be involved.

Previous reports have shown that Cdk2 (Trowbridge et al. 1997, Rogatsky et al. 1999) and GSK3 (Medunjanin



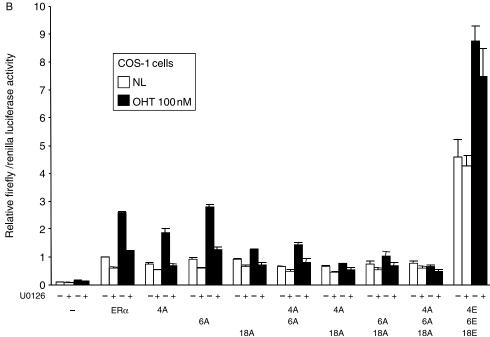


Figure 6 Effect of MAPK signaling activators and inhibitors on ERα activity. (A) NIH3T3 cells were co-transfected with ERE-3-TATA-luc, pRL-TK, and the wild-type ERα expression vector (ERα), or versions in which S104, S106, and S118 were substituted by alanine (4/6/18A) or glutamic acid (4/6/18E), as indicated. Cells were additionally co-transfected with empty expression vector (-) or expression vectors for Ras-V12, Raf-CAAX, or Raf-S621A, as shown. Cells were treated and luciferase assays carried out as for Fig. 4. (B) COS-1 cells were transfected with ERE-3-TATA-luc, pRL-TK, and the wild-type ERα expression vector (ERα) or versions with alanine or glutamic acid substitutions, as indicated. Cells were pre-incubated with U0126 (10 μM), 16 h following transfection, as indicated, for 1 h prior to addition of OHT, and harvested for luciferase assays 7 h later.

et al. 2005) can phosphorylate S104 and/or S106. However, whilst investigating the phosphorylation of S118, we had previously observed that mutation of S104 and S106 reduced the levels of ERa phosphorylation by MAPK in vitro (Chen et al. 2002). In agreement with this, we show here that S104 and S106 phosphorylation was stimulated by PMA and inhibited by the MEK1/2 inhibitor U0126, in transfected COS-1 cells. Further, co-transfection with activated Ras-V12 or Raf-CAAX, but not dominant-negative Raf-S621A, stimulated S104 and S106 phosphorylation. Together, these results suggest that serines 104 and 106 are phosphorylated by Erk1/2 MAPK. In vitro kinase experiments confirmed that, in addition to phosphorylating S118, Erk2 could also directly phosphorylate S104 and S106. Of the other kinases tested, Cdk2 was able to phosphorylate S104 and S118, and GSK3 able to phosphorylate S104, but to levels considerably lower than that achieved by Erk2. However, Cdk2 and/or GSK3 may phosphorylate S104 and/or S106 in vivo. Indeed, ligand-stimulated phosphorylation was in most cases insensitive to U0126, suggesting the involvement of other kinases. However, OHT- and ICI-induced phosphorylation of S104 did appear to partly involve MAPK, raising the possibility that multiple kinases are differentially involved, depending on the specific ligand.

In reporter gene assays, constitutively active Ras and Raf stimulated ERa activity in the absence of ligand or in the presence of OHT. Substitution of S104/S106/S118 by alanine prevented ERa activation by Ras and Raf, indicating that these sites are required for MAPK stimulation of ERa activity. Furthermore, U0126 inhibited ERa activity, as did dominant-negative Raf, but had little effect on triple S104, S106, and S118 mutants. These results suggest that the inhibitory effect of Raf-S621A and U0126 on ERα activity was due to the inhibition of phosphorylation at some or all of the three sites, and further implicates MAPK in the phosphorylation of S104, S106, and S118. Phosphorylation of S104, S106, and S118 could also be induced by ERα ligands; E₂, OHT, and ICI in COS-1 cells. The in vitro kinase assays indicated that ligand-binding results in marginally more efficient phosphorylation of ERa by MAPK, perhaps due to the altered conformation of ERa and/or unmasking of potential MAPK docking site(s) (Obenauer et al. 2003). However, ligand-stimulated phosphorylation of ERα in vivo was largely insensitive to U0126, and may be mediated by Cdk2 and/or GSK3 (Trowbridge et al. 1997, Rogatsky et al. 1999, Medunjanin et al. 2005).

In conclusion, phosphorylation of S104, S106, and S118 is important for ER α AF-1 activity, as displayed by enhanced ligand-independent, and E₂- and OHT-dependent, activities. This enhanced activity is not due to ligand hypersensitivity. No one site is critical, but lack of phosphorylation at all of the sites together results in near complete loss of AF-1 activity and prevents the

agonist action of OHT. Additionally, phosphorylation of these sites occurs in a partially interdependent manner and phosphorylation at each site appears to act via a similar mechanism to enhance ERα activity, suggesting that this region constitutes a phospho-regulated domain of cooperative MAPK phosphorylation sites. Activation of the EGF receptor and ErbB2 pathways, which signal through MAPK, has been associated with more aggressive breast cancer phenotypes and poor patient prognosis (Ross & Fletcher 1998, Arteaga 2001). These pathways have additionally been linked to the tamoxifen resistance phenotype (Benz et al. 1993, Kurokawa et al. 2000, Gee et al. 2001, Kurokawa & Arteaga 2003, Shou et al. 2004). The evidence presented here suggests that modulation of ER α phosphorylation can determine whether or not tamoxifen acts as an ER α agonist or antagonist, and that hyperphosphorylation may result in tamoxifen-induced activities at levels high enough to support the growth of cells that depend upon ERa activity, such as those found in the majority of breast cancers.

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