Epidermal Growth Factor Induces Phosphorylation of Extracellular Signal-Regulated Kinase 2 via Multiple Pathways

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Expression of p21^{rasAsn-17}, a dominant negative mutant of p21^{ras} that blocks p21^{ras} activation by growth factors, inhibits activation of extracellular signal-regulated kinase 2 (ERK2) by insulin and platelet-derived growth factor in rat-1 cells [A. M. M. de Vries-Smits, B. M. T. Burgering, S. J. Leevers, C. J. Marshall, and J. L. Bos, Nature (London) 357:602–604, 1992]. Here we report that expression of p21^{rasAsn-17} does not abolish epidermal growth factor (EGF)-induced phosphorylation of ERK2 in fibroblasts. Since EGF activates p21^{ras} in these cells, this indicates that EGF induces a p21^{ras}-independent pathway for the phosphorylation of ERK2 as well. We investigated whether activation of protein kinase C (PKC) or increase in intracellular calcium could be involved in p21^{ras}-independent signaling. In rat-1 cells, inhibition of either PKC, by prolonged 12-*O*-tetradecanoylphorbol-13-acetate (TPA) pretreatment, or calcium influx, by ethylene glycol-bis(β-aminoethyl ether)-*N*,*N*,*N'*,*N'*-tetraacetic acid (EGTA) pretreatment, did not abolish EGF-induced ERK2 phosphorylation. However, a combined inhibition of both p21^{ras} and calcium influx, but not PKC, resulted in a complete inhibition of EGF-induced ERK2 phosphorylation. In contrast, in Swiss 3T3 cells, inhibition of both p21^{ras} activation and TPA-sensitive PKC, but not calcium influx, inhibited EGF-induced ERK2 phosphorylation. These results demonstrate that in fibroblasts, EGF induces alternative pathways of ERK2 phosphorylation in a cell-type-specific manner.

Activation of transmembrane signaling receptors by first messengers generates the formation of second messengers. These second messengers are implicated in the induction of a myriad of intracellular signals of various biochemical constitutions. A common factor of signaling by many mitogenic stimuli is the activation of microtubule-associated protein 2 (MAP2) kinase (for recent reviews, see references 10, 40, and 42). Two forms of MAP2 kinase have been purified from fibroblasts with apparent molecular weights of 42 and 44 kDa. The p42^{MAPK} and p44^{MAPK} have been cloned and are also named, respectively, extracellular signal-regulated kinase 2 (ERK2) and 1 (ERK1), to reflect the diversity of signals that can regulate them (3). Activation of both ERK1 and ERK2 requires phosphorylation at neighboring threonine and tyrosine residues and dephosphorylation of activated ERK2 by serine/threonine phosphatases or tyrosine phosphatases results in inactive ERK2 (1, 16). For ERK2, the sites of phosphorylation are located within one tryptic peptide and have been assigned to Thr-183 and Tyr-185 (35).

Since full activation of ERK2 requires both threonine and tyrosine phosphorylation, it has been suggested that integration of two signaling pathways is required for the activation of the kinase (1). However, it is now clear that ERK2 is part of a kinase/phosphatase cascade in which at least several serine/threonine kinases act upstream of ERK2. The MAP kinase kinase (MAPKK or MEK) that activates ERK2 and phosphorylates the threonine and tyrosine site of ERK2 has recently been cloned by several groups (11, 23). MAPKK activity itself is lost upon treatment with protein phos-

In this study we have investigated whether in fibroblasts receptor tyrosine kinases couple to ERK2 exclusively via p21^{ras} activation. In our previous study we showed that protein kinase C (PKC) activation induces ERK2 activity by a p21^{ras}-independent mechanism, indicating that at least two different pathways for ERK2 activation exist (12). Since growth factors like EGF and PDGF can stimulate PKC activity, the question is whether these growth factors indeed employ redundant signaling pathways towards ERK2. Here we demonstrate that independent pathways to regulate ERK2 activity coexist within one cell type, that the nature of these pathways is cell type specific, and that, depending on

phatase 2A but not following treatment with phosphotyrosine phosphatases (13). This suggests that receptors do not directly activate the MAPKK, and indeed putative MAPKK kinases have recently been identified. One of these (denoted as MEKK) has been implicated in conveying signals from serpentine receptors (26), whereas another, the Raf-1 kinase, is implicated in mediating signals from receptor tyrosine kinases towards ERK2 (17, 24). Raf-1 kinase acts downstream of p21^{ras} in mitogenic signaling (22, 38), and recent experiments show that in vitro Raf-1 kinase binds p21ras in a GTP-dependent manner (31, 44, 46). Also, at least in yeast cells, Raf-1 kinase and p21^{ras} are capable of interacting in vivo (46). We and others have shown that activation of p21^{ras} is essential for the induction of ERK2 activity by insulin and platelet-derived growth factor (PDGF) in rat-1 fibroblasts and for ERK2 and Raf-1 kinase activity by nerve growth factor (NGF) and epidermal growth factor (EGF) in PC12 cells (12, 43, 45). This suggests a model in which receptor tyrosine kinases are coupled to the Raf-1/ERK2 kinase cascade by a nonkinase and nonphosphatase intermediate, e.g., p21^{ras}

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the cell type, a stimulus can employ different combinations of these pathways.

MATERIALS AND METHODS

Materials and cell culture. Phosphate-free Dulbecco modified Eagle medium (DMEM), insulin, and EGF were obtained from Sigma. ³²P_i (3,000 Ci/mmol) was from Amersham, PDGF-BB homodimer was from AMGEN, and NGF, ionomycin (Ca²⁺ ionophore), and indo-1/AM {1-[2-amino-5-(6-carboxyindol-2-yl)phenoxy]-2-(2'-amino-5'-methylphenoxy) ethane-N,N,N',N'-tetraacetic acid, pentaacetoxymethyl ester} were from Boehringer Mannheim.

Rat-1, A14, and Swiss 3T3 cells were routinely grown in DMEM-10% fetal calf serum (FCS)-0.05% glutamine. PC12 cells were grown in DMEM (high glucose)-10% FCS (heat inactivated)-5% horse serum. For growth factor stimulation experiments, subconfluent cultures of cells were grown in DMEM-0.5% FCS (PC12 cells in DMEM [high glucose]-0.5% horse serum) for 18 h prior to stimulation with growth factors.

GTP and GDP binding to p21^{ras}. Serum-arrested cell cultures were washed two times with phosphate-free DMEM and labeled with 100 μCi of ³²P_i per ml for 4 h. After growth factor stimulation, cells were lysed and processed as previously described (7). Mature p21^{ras} was separated from nonprocessed p21^{ras} by a Triton X-114 phase split as described elsewhere (7, 15). p21^{ras} was precipitated from the detergent phase with monoclonal antibody Y13-259. As a control, immunoprecipitation with a nonrelated antibody directed against simian virus 40 large T (KT3) was used. Bound nucleotides were eluted from immunoprecipitates and analyzed by ascending thin-layer chromatography (TLC) as described elsewhere (7), and after exposure to Kodak XAR film spots corresponding to GDP and GTP were cut out. The amount of radioactivity (in counts per minute) was determined by liquid scintillation counting.

Determination of ERK2 phosphorylation and activity. Cells were lysed and 100 μg of protein was immunoprecipitated with a polyclonal rabbit antiserum (anti-ERK2) directed against bacterially produced, near-full-length ERK2 (a gift from Chris Marshall, London, England). The immune complexes were collected by using protein A-Sepharose beads, washed, and incubated in a volume of 30 µl with 0.4 µCi of $[\gamma^{-32}P]ATP$ in 30 mM Tris (pH 8.0), 20 mM MgCl₂, 2 mM MnCl₂, and 10 µM ATP for 30 min with 7.5 µg of myelin basic protein as a substrate. The mixture was electrophoresed on a 15% polyacrylamide gel, blotted to nitrocellulose, and autoradiographed. The labeled protein was counted by scintillation counting. To measure the mobility shift of the ERK2 protein, 50 µg of total lysate was electrophoresed on a 10% polyacrylamide gel, blotted to nitrocellulose, and incubated with anti-ERK2. Immune complexes were detected by horseradish peroxidase second antibodies and then by enhanced chemiluminescence (Amersham). All experiments analyzing ERK2 phosphorylation and/or activation were repeated at least three times.

Measurement of [Ca²⁺]_i. Nearly confluent rat-1 cells grown on glass coverslips were loaded with 10 mM indo-1/AM at 37°C. Measurements were performed in N-2-hydroxyethylpiperazine-N'-2-ethanesulfonic acid (HEPES)-buffered saline of the following composition: KCl, 5 mM; NaCl, 140 mM; CaCl₂, 2 mM; MgCl₂, 1 mM; HEPES, 10 mM; and glucose, 10 mM. With a fluorescence spectrometer (Perkin-Elmer 3000), [Ca²⁺]_i was determined, with an excitation wavelength of 355 nm (5-nm slit) and an emission

wavelength of 405 nm (10-nm slit) according to standard procedures (14). PDGF-BB, stored in 1 M acetic acid plus 0.01% bovine serum albumin, was Speed-Vac dried before use and resuspended in HEPES-buffered saline.

RESULTS

Activation of p21^{ras} and ERK2 by PDGF and EGF in rat-1 cells. Previously, we have investigated growth factor-induced p21ras activation in NIH 3T3 fibroblasts. Both PDGF and EGF hardly induce p21^{ras}-GTP formation in these cells (7). Here we analyzed growth factor-induced p21^{ras} activation in rat-1 cells. The rat-1 cell line used is highly responsive to growth factors with respect to the induction of [3H]thymidine incorporation and growth in soft agar (8, and data not shown), confirming the original characterization by Kaplan et al. (19). Rat-1 fibroblasts were stimulated with either PDGF and EGF for 5 min, and GTP and GDP binding to p21^{ras} was determined (Fig. 1A). Both PDGF and EGF were found to stimulate p21^{ras}-GTP formation, although EGF gave rise to significantly higher relative levels of p21^{ras}-GTP compared with those induced by PDGF (basal, 17% GTP; EGF, 55% GTP; and PDGF, 25% GTP [n = 4]). EGF receptor numbers are somewhat lower than those of PDGF receptors on rat-1 cells, but they are comparable (PDGF receptor, ~80,000, and EGF receptor, ~30,000 [27a]), so in these cells, EGF appears to be a more efficient activator of p21^{ras} than PDGF.

Next, we investigated the induction of ERK2 activity following either PDGF or EGF stimulation of rat-1 fibroblasts. Activation of ERK2 can be measured by an immune complex kinase assay, with myelin basic protein as a substrate, or alternatively, activation can be monitored by the appearance of a more slowly migrating form in gel electrophoresis caused by phosphorylation (27). Mutation of both Thr-183 and Tyr-185 results in a protein that can no longer be phosphorylated and lacks the mobility shift (37). However, formally it is not proven that the appearance of phosphorylated ERK2 is identical to the activation of ERK2 per se. Stimulation of rat-1 fibroblasts with EGF and PDGF induced a complete shift in ERK2 mobility, indicating full activation of ERK2 (Fig. 1B).

p21^{ras} involvement in PDGF and EGF signaling. Previously, we demonstrated the use of a vaccinia virus vector expressing the dominant negative p21^{ras} mutant p21^{rasAsn-17} (vv-p21^{rasAsn-17}) in dissecting growth factor pathways mediated by p21^{ras} (12). These vectors, however, are not suitable for studying EGF signaling, since they express the gene encoding vaccinia virus growth factor (VGF). VGF can bind to the EGF receptor and stimulate EGF receptor activity (6). Therefore, we cloned p21^{rasAsn-17} into a vaccinia virus vector lacking VGF gene expression (vv-152) (30). Expression of p21^{rasAsn-17} was tested by infecting insulin receptor overexpressing NIH 3T3 cells (A14 cells) with various amounts of recombinant virus (Fig. 1C). Compared with the endogenous p21^{ras}, p21^{ras}Asn-17 was expressed at an approximately 10-fold-higher level (50 µl [20 PFU] of recombinant virus) or at an equal level (5 µl). Inhibition of insulin-induced ERK2 phosphorylation could already be observed at the lowest level of p21^{rasAsn-17} expression and was complete when p21^{rasAsn-17} was slightly overexpressed compared with endogenous p21^{ras} (Fig. 1C). Therefore, in all further experiments using vv-152-p21^{ras} Asn-17, cells were infected with 50 µl of the virus stock. We also tested whether p21^{ras} Asn-17 expression blocks not only ERK2 phosphorylation but also activation, as measured by an immune complex kinase assay

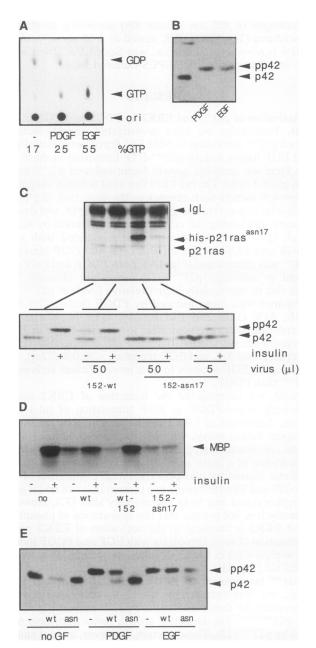


FIG. 1. PDGF-induced, but not EGF-induced, ERK2 phosphorylation in rat-1 cells is blocked by p21^{rasAsn-17}. (A) Activation of p21^{ras} by PDGF and EGF in rat-1 cells. Serum-arrested rat-1 cells were treated with PDGF-BB homodimer (50 ng/ml) and EGF (20 ng/ml) for 5 min. p21^{ras} activation was measured by determining the nucleotide content of p21^{ras}. Shown is an autoradiogram of a TLC separation of guanine nucleotides. Indicated are the positions of the GTP and GDP standards, and under each lane, the average amount (n = 4) of p21^{ras}-GTP (as a percentage of total nucleotide bound to p21^{ras} [GTP plus GDP]) is indicated. (B) Phosphorylation of ERK2 by PDGF and EGF in rat-1 cells. Serum-arrested rat-1 cells were treated with PDGF-BB homodimer (50 ng/ml) and EGF (20 ng/ml) for 5 min, total protein lysates were prepared by scraping cells into sample buffer, and phosphorylation of ERK2 was determined by the occurrence of a mobility shift. The position of the unphosphorylated (inactive) ERK2 is indicated by p42, and the phosphorylated ERK2 is indicated by pp42. (C) Characterization of recombinant non-VGF-producing (vv-152) p21^{ras Asn-17} expressing vaccinia virus. Construction of viruses has been described previously (12, 30). Cells overexpressing the insulin receptor (A14) were infected with various

(Fig. 1D). In agreement with our previous results (12), this construct was also found to inhibit insulin-induced ERK2 activation in A14 cells (Fig. 1D). In addition, the increase in basal ERK2 activity observed following infection with wild-type virus, presumably due to VGF expression, was reduced by using wild-type vv-152 (Fig. 1D, compare lanes wt – and wt + with lanes 152 – and 152 +).

Next, we compared PDGF- and EGF-induced ERK2 phosphorylation in rat-1 cells following infection with vv-152-p21^{rasAsn-17} (Fig. 1E). In contrast to PDGF, EGF-induced ERK2 phosphorylation was not inhibited by the expression of p21^{rasAsn-17}.

Activation of ERK2 by EGF occurs also through a p21^{ras}-independent pathway. As shown in Fig. 1A, the levels of p21^{ras}-GTP induced by EGF compared with those induced by PDGF are much higher. It would therefore be possible that the level of p21^{ras}-Asn-17 expression is sufficient to block p21^{ras} activation completely following PDGF stimulation but not following EGF stimulation. A residual increase in p21^{ras}-GTP might then account for ERK2 phosphorylation. In NIH 3T3 cells overexpressing the insulin receptor (A14 cells), insulin and EGF induce similar levels of p21^{ras}-GTP as well as similar kinetics of ERK2 activation (Fig. 2A). Despite the fact that both insulin and EGF induce up to 60 to 70% p21^{ras}-GTP, only insulin-induced ERK2 phosphorylation is inhibited significantly by p21^{ras}-Asn-17 expression (Fig. 2B).

We have previously shown that both insulin and EGF activate p21^{ras} by increasing nucleotide exchange (30). This mechanism of p21^{ras} activation is blocked by p21^{rasAsn-17} expression (5, 30), and p21^{rasAsn-17} has been shown to compete with endogenous p21^{ras} for interaction with mSos, a putative p21^{ras} nucleotide exchange factor, thus providing a rationale for the inhibition of p21^{ras} activation (4). These results strongly suggest that p21^{rasAsn-17} will inhibit EGF-induced p21^{ras} activation as efficiently as insulin-induced p21^{ras} activation, since p21^{rasAsn-17} acts between activation of the exchange factor and p21^{ras}. However, the possibility that for some unknown reason EGF-induced p21^{ras} activation activation is refractory to our approach of p21^{rasAsn-17} expression, by using vaccinia virus, still remains. For PC12

amounts of wild-type virus or recombinant virus and either left unstimulated or stimulated with insulin (10^{-6} M). Expression of p21^{rasAsn-17} was determined by immunoprecipitation with the p21^{ras}specific monoclonal antibody Y13-238 and immunoblotting with Y13-259 to detect p21^{ras} protein. Owing to the presence of six histidine residues at the N terminus, the p21^{rasAsn-17} protein runs at a slightly slower mobility than endogenous p21^{ras}. Of the lysates, 50 μg was used to determine phosphorylation of ERK2. The position of the unphosphorylated (inactive) ERK2 is indicated by p42, and the phosphorylated ERK2 is indicated by pp42. IgL, immunoglobulin light chain. (D) Activation of ERK2 is blocked by non-VGF-producing (vv-152) p21^{rasAsn-17} expressing vaccinia virus. A14 cells were either left uninfected or infected with viruses (50 µl) as indicated, prior to stimulation with insulin (10^{-6} M). ERK2 was immunoprecipitated, and ERK2 activity of immunoprecipitates was determined as described in Materials and Methods. MBP, myelin basic protein. (E) Phosphorylation of ERK2 in rat-1 cells by PDGF, but not EGF, is blocked by p21^{rasAsn-17} expression. Rat-1 cells either were left uninfected (-) or were infected with control virus (wt) or virus (asn). Following infection cells were either left untreated (no GF) or treated with PDGF (50 ng/ml) or EGF (20 ng/ml) as indicated. The phosphorylation of ERK2 was determined by the occurrence of a mobility shift. The position of the unphosphorylated (inactive) ERK2 is indicated by p42, and the phosphorylated ERK2 is indicated by pp42.

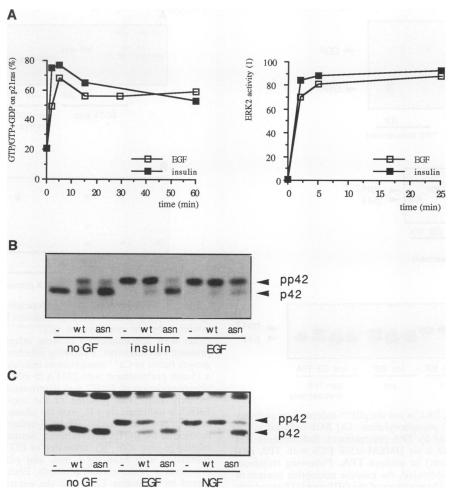


FIG. 2. Phosphorylation of ERK2 by EGF in A14 cells occurs through p21^{ras}-independent pathways. (A) Time course of EGF- and insulin-induced p21^{ras} activation and ERK2 activation in insulin receptor-overexpressing cells (A14 cells). A14 cells were serum arrested and treated for various times with either EGF (20 ng/ml) or insulin (10⁻⁶ M). Activation of p21^{ras} and ERK2 was measured as described in the legend to Fig. 1A and D, respectively. The percentage of GTP is expressed as relative to the total amount of labeled nucleotide (GTP + GDP). The activation of ERK2 is expressed as an apparent enzyme activity (1 U = 1 pmol of PO₄ per min per 100 μg of total protein). (B) Phosphorylation of ERK2 in A14 cells by insulin, but not EGF, is blocked by p21^{rasAsn-17} expression. A14 cells either were left uninfected (–) or were infected with control virus (wt) or with p21^{rasAsn-17} virus (asn). Following infection, cells either were left untreated (no GF) or were treated with insulin (10⁻⁶ M) or EGF (20 ng/ml) as indicated. The phosphorylation of ERK2 was determined by the occurrence of a mobility shift. The position of the unphosphorylated (inactive) ERK2 is indicated by p42, and the phosphorylated ERK2 is indicated by p42. (C) Effects of p21^{rasAsn-17} expression on EGF- and NGF-induced ERK2 phosphorylation in PC12 cells. PC12 cells either were left uninfected (–) or were infected with control virus (wt) or with p21^{rasAsn-17} virus (asn). Following infection, cells were either left untreated (no GF) or treated with EGF (20 ng/ml) or NGF (20 ng/ml) as indicated. The phosphorylation of ERK2 was determined by the occurrence of a mobility shift. The position of the unphosphorylated (inactive) ERK2 is indicated by p42, and the phosphorylated ERK2 is indicated by p942.

cells, it has been shown that induction of p21^{rasAsn-17} expression inhibits EGF-induced ERK2 activity (42, 45). Therefore, we tested the ability of vv-152-p21^{rasAsn-17} to inhibit EGF signaling in PC12 cells. As shown in Fig. 2C, infection of PC12 cells with vv-152-p21^{rasAsn-17} efficiently inhibits EGF-induced ERK2 phosphorylation. This demonstrates that the observed lack of inhibition of EGF-induced ERK2 phosphorylation in rat-1 and A14 cells is not inherent to the use of the vv-152-p21^{rasAsn-17} construct to express p21^{rasAsn-17}.

In conclusion, these results suggest that in rat-1 and A14 cells, ERK2 phosphorylation by EGF occurs through the activation of p21^{ras} but that EGF activates a pathway that is fully capable of stimulating ERK2 phosphorylation in the absence of p21^{ras} activation.

TPA-sensitive PKC is not involved in EGF-induced p21^{ras}-independent ERK2 phosphorylation in A14 cells. If p21^{ras} activation is not the only determinant in EGF signaling towards ERK2, what, then, can be the other(s)? Activation of PKC by short-term treatment with 12-O-tetrade-canoylphorbol-13-acetate (TPA) induces ERK2 activation (10) (Fig. 3B, lane 4). However, identical treatment does not result in a significant activation of p21^{ras} (29). In addition, TPA-induced ERK2 activation is also not inhibited by p21^{rasAsn-17} expression (12) (see Fig. 6C). These results suggest the possibility of growth factor-induced activation of TPA-sensitive PKC as a pathway of p21^{ras}-independent activation of ERK2. To study a role for this pathway in the induction of ERK2 phosphorylation by EGF, A14 cells were depleted of TPA-sensitive PKC by prolonged treatment with

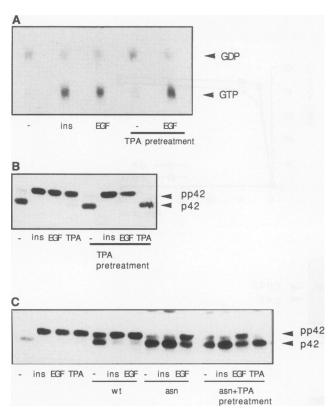


FIG. 3. In A14 cells, PKC is not the p21^{ras}-independent pathway in EGF-induced ERK2 phosphorylation. (A) EGF-induced p21^{ras} activation is not inhibited by TPA pretreatment. Subconfluent A14 cells were grown for 18 h on DMEM-0.5% FCS with TPA (200 ng/ml) (TPA pretreatment) or without TPA. Following treatment with growth factors as indicated, the guanine nucleotide content of p21^{ras} was determined. The positions of the GDP and GTP standards are indicated. (B) EGF-induced ERK2 phosphorylation is not inhibited by TPA pretreatment. A14 cells were treated and stimulated as described for panel A; TPA stimulation was with 100 ng/ml. ERK2 phosphorylation was determined by the occurrence of a mobility shift. The position of the unphosphorylated (inactive) ERK2 is indicated by p42, and the phosphorylated ERK2 is indicated by pp42. (C) EGF induces ERK2 phosphorylation by a p21^{ras}-independent and PKC-independent pathway. A14 cells either were left uninfected or were infected with control virus (wt) or with p21^{rasAsn-17} virus (asn) either alone or in combination with TPA (100 ng/ml) pretreatment (asn+TPA pretreatment). Following infection, cells either were left untreated (-) or were stimulated with EGF (20 ng/ml), insulin (ins; 10⁻⁶ M), or TPA (100 ng/ml) as indicated. The phosphorylation of ERK2 was determined by the occurrence of a mobility shift. The position of the unphosphorylated (inactive) ERK2 is indicated by p42, and the phosphorylated ERK2 is indicated by pp42.

TPA. This resulted in the loss of immunodetectable PKC, TPA-inducible myristylated arginine-rich C kinase substrate (MARCKS) phosphorylation, and TPA-induced ERK2 activation (Fig. 3B) (data not shown) (29). Under these conditions, both EGF-induced p21^{ras} activation (Fig. 3A) and EGF-induced ERK2 phosphorylation (Fig. 3B) are not affected. In the case of EGF it can be proposed that neither p21^{ras} nor PKC signaling is essential but that both are sufficient. Therefore, we analyzed the possibility that both pathways are employed in EGF signaling towards ERK2 phosphorylation. To this end, EGF-induced ERK2 phosphorylation was analyzed in PKC depleted, vv-152-p21^{ras}Asn-17

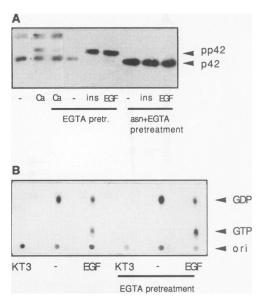
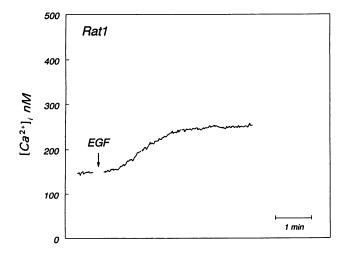


FIG. 4. EGF-induced p21^{ras}-independent signaling is inhibited by EGTA pretreatment. (A) EGF-induced ERK2 phosphorylation is inhibited by the combination p21^{ras}-Asn-17 and EGTA. A14 cells either were left uninfected or were infected with p21^{ras}-Asn-17 expressing vaccinia virus. Following infection, cells were treated with growth factor or Ca²⁺ ionophore as indicated, either with or without a 15-min pretreatment with EGTA (5 mM final concentration). The phosphorylation of ERK2 was determined by the occurrence of a mobility shift. The position of the unphosphorylated (inactive) ERK2 is indicated by p42, and the phosphorylated ERK2 is indicated by p42. ins, insulin. (B) EGF-induced p21^{ras} activation is not modulated by EGTA pretreatment. Serum-starved A14 cells were labeled in vivo with ³²P_i, untreated or EGTA pretreated cells were stimulated with EGF for 5 min, and p21^{ras} was precipitated as described in Materials and Methods. Eluted nucleotides were separated by ascending TLC, and the autoradiogram of the TLC is shown.

infected A14 cells (Fig. 3C). As shown in Fig. 3C, this did not result in complete inhibition of EGF-induced ERK2 phosphorylation. For rat-1 cells, essentially identical results were obtained (not shown). Thus, it appears that activation of a TPA-sensitive PKC is not the p21^{ras}-independent pathway in EGF signaling towards ERK2 activation in rat-1 and A14 cells.

Involvement of Ca²⁺-dependent pathways in EGF-induced **ERK2 activation.** The role of Ca²⁺ in EGF signaling has been well documented (reference 36 and references therein). Therefore, in search of other signaling pathways involved in EGF-induced ERK2 phosphorylation, we analyzed the involvement of Ca²⁺-dependent pathways. First, we tested whether Ca²⁺ by itself could induce ERK2 phosphorylation in A14 cells. The Ca²⁺ ionophore ionomycin rapidly induced ERK2 phosphorylation (Fig. 4A). This induction was dependent on Ca²⁺, since pretreatment of cells for 5 min with EGTA to sequester extracellular Ca2+ did inhibit ionomycininduced ERK2 phosphorylation. This also indicates that the EGTA concentration used is sufficient to inhibit signaling induced by a large increase in intracellular Ca²⁺, as in the case of ionomycin. In contrast, EGTA pretreatment did not significantly affect EGF-, insulin (Fig. 4A)- or PDGF (not shown)-induced ERK2 phosphorylation. Again, we tested whether a combination of p21^{rasAsn-17} expression and EGTA pretreatment could block EGF-induced ERK2 phosphoryla-



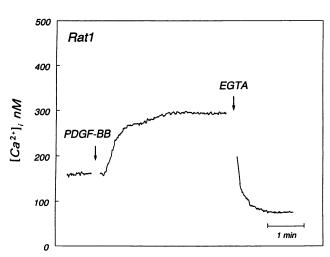


FIG. 5. PDGF- and EGF-induced Ca²⁺ fluxes. Serum arrested rat-1 cells were loaded with indo-1/AM as described in Materials and Methods. Following treatment with growth factor, fluorescence was monitored in time. A trace is shown from a representative experiment following EGF and PDGF treatment.

tion (Fig. 4A). Interestingly, the phosphorylation of ERK2 by EGF was now completely blocked. EGF-induced phosphorylation of ERK2 in EGTA-pretreated, p21^{ras}Asn-17-infected rat-1 cells was also completely blocked (not shown). As shown in Fig. 4B, EGTA pretreatment did not significantly alter EGF-induced p21^{ras} activation. This suggests that EGTA pretreatment defines a pathway separate from that of p21^{ras}. We conclude from these results that in A14 and rat-1 cells, both the p21^{ras} and the Ca²⁺-mediated pathways for EGF signaling towards ERK2 appear sufficient, but neither one is essential.

PDGF- and EGF-induced Ca²⁺ response. If the second pathway towards ERK2 phosphorylation employed by EGF is induced by Ca²⁺ entry, then it is to be expected that insulin and PDGF would differ from EGF in this respect. To test whether there is a difference between PDGF and EGF signaling with respect to their ability to induce Ca²⁺ fluxes, rat-1 cells were loaded with indo-1/AM and stimulated with

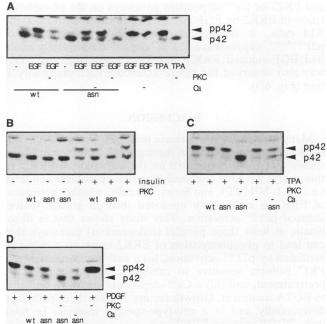


FIG. 6. p21^{ras}-dependent and -independent signaling in Swiss 3T3 cells. Serum-arrested Swiss 3T3 cells either were left uninfected or were infected with wild-type vaccinia virus (wt) or p21^{rasAsn-17} expressing virus (asn). In some cases, uninfected or infected cells were also pretreated with TPA (300 ng/ml for 16 h) to downmodulate PKC expression (indicated as PKC-) or pretreated with EGTA (indicated by Ca -) before growth factor stimulation. Shown are representative examples of the analysis of induction of ERK2 phosphorylation by EGF (A), insulin (B), TPA (C), and PDGF (D). As before, the phosphorylation of ERK2 was determined by the occurrence of a mobility shift. The position of the unphosphorylated (inactive) ERK2 is indicated by p42, and the phosphorylated ERK2 is indicated by pp42.

PDGF or EGF, and Ca²⁺ traces were recorded. As shown in Fig. 5, it is clear that the Ca²⁺ transients induced by EGF and PDGF are very similar. This indicates that Ca²⁺ mobilization by itself is not the difference between EGF and PDGF signaling in rat-1 cells. Apparently, EGTA pretreatment defines a pathway towards ERK2 phosphorylation that is not Ca²⁺ induced but is Ca²⁺ dependent, yet specifically induced by EGF and not by PDGF.

Growth factor-induced ERK2 activation in Swiss 3T3 cells. Next, we addressed the question whether in other fibroblast cell lines EGF employs a similar redundant mechanism of ERK2 phosphorylation as in rat-1 and A14 cells. We chose to study Swiss 3T3 cells, since in these cells a role for TPA-sensitive PKC has been described in the activation of ERK2 by oncogenic p21^{ras} (27). Also, in Swiss 3T3 cells, expression of p21^{rasAsn-17} did not inhibit EGF- or TPAinduced ERK2 phosphorylation (Fig. 6A and C). However, insulin-induced phosphorylation of ERK2 was completely blocked by p21^{rasAsn-17} expression, suggesting that insulin again activates only the p21ras-mediated branch of ERK2 phosphorylation (Fig. 6B). As for the other fibroblast cell lines, TPA induced ERK2 phosphorylation by a p21rasindependent pathway fully blocked by prolonged TPA pretreatment (Fig. 6C). Surprisingly, in contrast to rat-1 cells, PDGF-induced ERK2 phosphorylation was not inhibited by p21^{rasAsn-17} expression (Fig. 6D). As described above for the other cell lines, we tested the effect of inhibiting both p21^{ras}

and PKC- or Ca²⁺-dependent pathways on the phosphorylation of ERK2 by EGF and PDGF. In contrast to rat-1 and A14 cells, it was found that now a combination of p21^{rasAsn-17} expression and PKC depletion completely inhibited EGF-induced ERK2 phosphorylation (Fig. 6A). This was also observed for PDGF-induced ERK2 phosphorylation (Fig. 6D).

DISCUSSION

Many growth factors activate both p21ras and ERK2. For some cell types and some of these growth factors, it has been shown that p21^{ras} is essential for the activation of ERK2. In this study, we have analyzed in various fibroblast cell lines i.e., rat-1, NIH 3T3, and Swiss 3T3 cells, whether activation of ERK2 is exclusively mediated through growth factorinduced p21^{ras} activation. This study shows that in fibroblasts, at least three parallel (independent) pathways that can lead to phosphorylation of ERK2 exist: (i) a pathway mediated by p21^{ras} activation, (ii) a pathway mediated by a PKC isoform sensitive to inhibition by prolonged TPA pretreatment, and (iii) a Ca²⁺-dependent pathway sensitive to EGTA treatment. Growth factors employ these pathways differentially and in a cell-type-specific manner. In rat-1 cells, PDGF-induced ERK2 activation appears to be exclusively mediated by p21^{ras}, since expression of p21^{rasAsn-17} completely inhibits ERK2 activation, and PKC depletion or EGTA treatment has no effect. In contrast, ERK2 activation by TPA is not mediated by p21ras and requires a TPAsensitive PKC. The activation of ERK2 by EGF appears to be mediated by two pathways acting simultaneously. The first pathway involves p21^{ras}, whereas the second is dependent on extracellular Ca²⁺. This conclusion is based upon the following observations. First, EGF induces p21^{ras} activation and a transient increase in Ca²⁺. Second, blocking only Ca2+ increase or p21ras activation does not inhibit ERK2 phosphorylation significantly. Third, this failure to fully block EGF-induced ERK2 phosphorylation was not due to incomplete inhibition of p21^{ras} activation by p21^{rasAsn-17} (Fig. 1C and 2) or due to incomplete inhibition of the Ca²⁺ increase (Fig. 4A). Fourth, only combined inhibition of Ca²⁺ influx and p21^{ras} activation resulted in full inhibition of ERK2 phosphorylation.

In Swiss 3T3 cells, growth factor induction of ERK2 activity occurs along similar pathways, yet compared with rat-1 and A14 cells, the ability of growth factors to activate these pathways is different. In this cell line, PDGF and EGF activate both the p21^{ras}- and the (TPA-sensitive) PKC-mediated pathways. However, other stimuli activate only either one pathway, e.g., insulin activates the p21^{ras}-mediated pathway and TPA activates the PKC-mediated pathway. Other studies analyzing the effects of PKC inhibition have already suggested that in Swiss 3T3 cells and derivatives (3T3-TNR9), activation of ERK2 occurs through PKC-dependent and -independent pathways (20, 25). From our experiments it is apparent that the p21^{ras}-mediated pathway is this PKC-independent pathway. In addition, this corroborates the notion that, also in the Swiss 3T3 cells, lack of inhibition by p21^{ras}Asn-17 is not due to insufficient expression of p21^{ras}Asn-17.

Several observations argue that the above-mentioned pathways are indeed independent pathways. PKC-mediated activation of ERK2 in fibroblasts is different from p21^{ras}-mediated ERK2 activation, since stimulation of fibroblasts with TPA does not result in p21^{ras} activation. TPA-induced ERK2 activation is not blocked by p21^{ras/Asn-17} expression,

and TPA pretreatment does not (rat-1 and A14) or only modestly (Swiss 3T3) attenuates tyrosine kinase receptorinduced ERK2 activation. This conclusion appears to contrast with some other reports studying the relationship of PKC and p21^{ras} in which PKC is positioned either upstream or downstream of p21^{ras}. It should be noted, however, that some of these studies make use of the properties of oncogenic p21^{ras}, which is not necessarily identical to growth factor-stimulated increase in p21^{ras}-GTP (for a discussion, see reference 12). Furthermore, in these studies p21^{ras}Asn-17 is introduced by scrape loading or transient transfection, and the expression levels of p21^{rasAsn-17} we achieve may be lower than those obtained after transient transfection or scrape loading of p21^{rasAsn-17} protein. In PC12 cells, a differential effect of p21^{rasAsn-17} expression has already been noted. Low levels of p21^{rasAsn-17} block the NGF-induced differentiation response but not NGF-induced expression of early response genes (c-fos and c-jun); high levels of p21^{rasAsn-17} also block NGF-induced early response gene expression (41). Whether high levels of p21^{rasAsn-17} expression inhibit only p21ras-mediated events or also result in aspecific inhibition of receptor signalling remains to be established. Irrespective of this, the level of p21^{rasAsn-17} expression we obtain (Fig. 1C) is sufficient to inhibit the induction of high levels of p21^{ras}-GTP by insulin in A14 cells (60 to 70% GTP) to such an extent that insulin-induced ERK2 phosphorylation and activation is inhibited not only in these cells but also in Swiss 3T3 cells (Fig. 1C, 1D, 2B, and 6B) (30). Finally, recent results show that certain PKC isozymes can directly activate the Raf-1 kinase and thereby ERK2 (21, 39). This would indicate that the PKC-mediated pathway and the p21ras-mediated pathway converge at the level of Raf-1 kinase activation.

The pathway defined by EGTA pretreatment appears to be different from the p21^{ras}-mediated pathway. Pretreating cells with EGTA does not affect EGF-induced p21^{ras} activation, suggesting that at least EGF-induced Ca²⁺ mobilization does not play an essential role in EGF-induced p21^{ras} activation. Still, Ca2+-dependent signaling may be involved in modulating p21ras-GTP levels. For example, in situ activation of p21^{ras} in permeabilized T cells is slightly lowered in Ca²⁺free medium (18). Also, we observed a small increase in p21^{ras}-GTP levels, although not reproducibly, after treatment with Ca2+ ionophore. Therefore, we cannot fully exclude the possibility that the Ca²⁺-dependent signal converges towards the ERK2 pathway at the level of p21^{ras} activation. If so, this would indicate that the mechanism by which the Ca2+-dependent signal contributes to p21ras activation is independent of the recently proposed mSos-mediated pathway (reviewed in reference 28), since our experiments with insulin suggest that p21rasAsn-17 expression completely blocks this pathway. Inactivation of p21^{ras} GT-Pase-activating proteins by the Ca²⁺-dependent pathway may be such a mSos-independent p21^{ras}-activating pathway.

At present the nature of the signal that is blocked by EGTA pretreatment is unknown. It is important to stress that apparently a Ca²⁺-dependent signal and not a Ca²⁺-induced signal is involved. The experiments using the Ca²⁺ ionophore ionomycin show that Ca²⁺ influx by itself can induce ERK2 phosphorylation, as was also reported by Chao et al. for human foreskin cells (9). However, the increase in intracellular Ca²⁺ after ionophore treatment is more than 10-fold higher than obtained with any natural agonist that increases intracellular Ca²⁺. For instance, treatment of rat-1 cells with endothelin, inducing a strong Ca²⁺ response, results in only a very small increase of ERK2

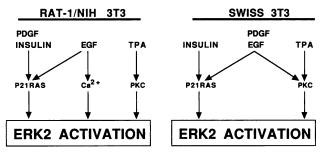


FIG. 7. ERK2 signaling pathways in fibroblasts. Different pathways leading to operational ERK2 phosphorylation and activation in rat-1, NIH 3T3, and Swiss 3T3 fibroblasts as identified in this study. p21^{ras} is the pathway inhibited by p21^{rasAsn-17} expression, Ca²⁺ is the Ca²⁺-dependent pathway defined by EGTA pretreatment, and PKC is the pathway involving those PKC isozymes that can be inhibited by a prolonged pretreatment with TPA. For a detailed discussion, see the text.

activity. In addition, when analyzing a dose response for endothelin, the induction of ERK2 activity does not appear to correlate with the ability of endothelin to stimulate an increase in Ca²⁺ (32) (data not shown). Finally, we compared EGF and PDGF with respect to the ability to induce an increase in intracellular Ca²⁺, and no differences were observed (Fig. 5). Yet EGF, and not PDGF, induced a redundant, Ca²⁺-dependent, signaling pathway towards ERK2. Thus, it appears that EGF can induce an EGF-specific, Ca²⁺-dependent enzyme activity involved in regulating ERK2 activity.

Since in Swiss 3T3 cells a PKC isozyme is involved, it would be possible that in rat-1 and A14 cells an EGTAsensitive, but not TPA-sensitive, PKC isozyme is involved in EGF signaling. Molecular cloning analysis of PKC has revealed a family of multiple isoforms (for a review, see reference 34). A marked difference between these isoforms is their sensitivity to Ca^{2+} . cPKC α , - β I, - β II, and - γ are dependent on Ca^{2+} for their activity, whereas nPKC ϵ , - δ , - ζ , and -η/L are not. The EGTA pathway in EGF signaling towards ERK2 as defined in this study could thus involve the isoforms cPKC α , - β I, - β II, and - γ . The ability to downregulate certain PKC isoforms by prolonged treatment with TPA differs between cell lines. Immunoblotting with monoclonal antibody MC5 (Amersham), detecting cPKCα, showed complete loss of immunodetectable PKC in A14 cells (data not shown). This leaves the possibility that cPKCβI, -βII, or -y is involved in Ca²⁺-dependent ERK2 activation. However, EGTA pretreatment by itself or in combination with p21^{rasAsn-17} expression did not block TPAinduced ERK2 phosphorylation (Fig. 6C) (data not shown). Furthermore, in human foreskin fibroblasts, Ca²⁺ does appear to activate ERK2 without activating PKC (9). Finally, Buday and Downward, using our clone of rat-1 cells, have obtained no evidence of PKC activation following EGF treatment (5). From these data, we conclude that in these cells activation of PKC is not involved in EGF-induced phosphorylation of ERK2 mediated by p21ras and also that the pathway which is blocked by EGTA is unlikely to involve PKC

The results described in this report suggest a model for cell-type-specific signaling towards ERK2 that is depicted in Fig. 7. In fibroblasts we can discriminate three different pathways that can coexist within one cell type: a p21^{ras}-mediated pathway inhibited by p21^{rasAsn-17} expression, a

PKC-mediated pathway inhibited by TPA pretreatment, and a Ca²⁺-dependent pathway inhibited by EGTA pretreatment. Depending on the cell type, growth factors may connect to one or more of these pathways. The cellular context, i.e., lack of mediators or presence of inhibitors, will determine which of these connections is made. It is clear that our results do not provide a quantitative measurement of the relative percentages at which these different pathways contribute to ERK2 activation; rather, they provide a qualitative measurement of the type of pathways involved in ERK2 activation. Also, at present we do not know whether these pathways converge at the level of ERK2 activation or at some level preceding ERK2 activation. For the PKC-mediated pathway, it is clear that it may converge at the level of Raf-1 kinase activation (21, 39). Whether the Ca²⁺-dependent signal also converges at the Raf-1 kinase level is currently under investigation. Our studies thus far suggest that for all genuine growth factors (not TPA), the p21^{ras}mediated pathway is a common factor. In this respect it is noteworthy that in all insulin-responsive cells tested thus far, insulin will activate ERK2 only by the p21ras-dependent pathway. This is in agreement with earlier findings (2, 33) that led to the suggestion that growth factor signaling could be divided in an insulin-like branch and a growth factorspecific branch.

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