

Extracellular Matrix Fibronectin Increases Prostaglandin E₂ Receptor Subtype EP4 in Lung Carcinoma Cells through Multiple Signaling Pathways

THE ROLE OF AP-2*

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We have previously demonstrated that fibronectin (Fn) stimulates the proliferation of non-small cell lung carcinoma (NSCLC) cell growth through the induction of cyclooxygenase-2 (COX-2) and prostaglandin E₂ secretion. Here, we demonstrate that NSCLC cells express mRNA and protein for the prostaglandin E₂ receptor EP4 and that Fn enhances its stimulatory effect by inducing the expression of EP4, but not of EP1, EP2, and EP3 receptor subtypes. The effect of Fn on EP4 was inhibited by an antibody against $\alpha 5 \beta 1$ integrin and by phosphoinositide 3-kinase (wortmannin) and extracellular signal-regulated kinase (PD98095), but not by protein kinase C (calphostin C), of protein kinase B (rapamycin), or mammalian target of rapamycin (rapamycin). siRNA was also inhibited by the binding activity in the presence of oligonucleotides. Using full-length and mutant constructs, we found that Fn stimulation increases EP4 expression when one AP-2 site (–100 to –150 bp) is present. Nuclear AP-2 α protein expression is increased in these pathways. Our results indicate that the effect of Fn on proliferation is mediated through EP4. Fn induces EP4 expression through integrin-dependent signals that include induction of phosphoinositide 3-kinase and phosphoinositide-dependent kinase as well as expression of COX-2. These events lead to activation of the transcription factor AP-2 α , which interacts with specific regions in the EP4 gene promoter, leading to transcription of the EP4 gene.

Extracellular matrix proteins are considered to play roles in the migration and differentiation of various cells, including car-

cino cells. Fibronectin (Fn),² a matrix glycoprotein highly expressed in tobacco and human lung diseases, has been shown to stimulate carcinoma cell growth, including lung carcinoma (1–3). We have previously demonstrated that Fn stimulated human lung carcinoma cell growth through the induction of cyclooxygenase-2 (COX-2) and prostaglandin E₂ secretion. Prostaglandin E₂ is considered important in the regulation of inflammation and inhibition of PGE₂ synthesis. The effect of PGE₂ on cell growth is mediated through its binding to its receptors. These receptors are involved in tumor growth and progression. For example, PGE₂ receptors are involved in tumor growth and progression in colorectal carcinoma (7, 8). In mice, PGE₂ receptors suggested a role for EP4 in tumor growth (7). PGE₂ and its signaling through EP4 have been shown to mediate non-small cell lung carcinoma (NSCLC) invasiveness (10). Taken together, these observations suggest that manipulation of prostaglandin E₂ signaling downstream from COX-2 produces more profound effects on carcinoma reduction than COX-2 inhibition alone and could be the basis for new approaches for the prevention of carcinoma.

At present, the mechanisms that link Fn and EP4 gene expression and how they might relate to lung carcinoma are unknown. Herein, we explore the relationship between these molecules and their role in lung carcinoma cell growth. Our results show that Fn stimulates lung carcinoma cell growth and that this inductive effect is partly dependent upon stimulation of PGE₂ production and induced PGE₂ receptor subtype EP4 gene expression, which is mediated through integrin-dependent signals, including the activation of phosphoinositide 3-kinase (PI3K) and extracellular signal-regulated kinase (ERK) signaling pathways.

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² The abbreviations used are: Fn, fibronectin; COX-2, cyclooxygenase-2; PGE₂, prostaglandin E₂; NSCLC, non-small cell lung carcinoma; PI3K, phosphoinositide 3-kinase; ERK, extracellular signal-regulated kinase; dmPGE₂, 16,16-dimethylprostaglandin E₂; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; RT, reverse transcription; siRNA, small interfering RNA; ODN, oligodeoxynucleotide(s); C/EBP, CCAAT/enhancer-binding protein; mTOR, mammalian target of rapamycin; PKA, cAMP-dependent protein kinase; PKC, protein kinase C.

EXPERIMENTAL PROCEDURES

Culture and Chemicals—NSCLC cell lines H1838 and H2106 were obtained from the American Type Culture Collection (Manassas, VA), and were grown in RPMI 1640 medium (H1838) supplemented with 10% heat-inactivated fetal bovine serum, HEPES buffer, 50 IU/ml penicillin/streptomycin, and 1 μ g of amphotericin (complete medium) or in Dulbecco's modified Eagle's medium/F-12 medium (H2106) supplemented with 10% heat-inactivated fetal bovine serum, 0.005 mg/ml insulin, 0.01 mg/ml transferrin, 30 nM sodium selenite, 10 nM hydrocortisone, 10 nM β -estradiol, 10 mM HEPES, as described previously (4). Afterward, cells were harvested and replaced in serum-free medium on Fn- or collagen type 1-coated culture plates for all experiments described later (the plates were coated with the matrix components diluted in buffer containing bovine serum albumin overnight at 4 °C. Afterward, the supernatants were removed, and the dishes were washed with phosphate-buffered saline three times before experiments were initiated). Mouse anti-human integrin α 5 β 1 (MAB1969) and anti-integrin α 2 β 1 antibodies (MAB1967) were purchased from Chemicon International Inc. (Temecula, CA). 16,16-dimethylprostaglandin E_2 (dmPGE $_2$), Polyclonal antibodies against COX-2, EP1, EP2, EP3, and EP4 were obtained from Cayman Chemical Co. [methyl- 3 H]Thymidine was purchased from Amersham Biosciences. The [γ - 32 P]ATP was purchased from PerkinElmer Life Sciences, Inc. Mammalian rapamycin (mTOR) inhibitor, rapamycin, and antibodies specific for Akt, ERK1, ERK2, and other forms ((AktS473) (ERK Thr 421 /Ser 424)) were purchased from Cell Signaling Inc. The ERK1/2 inhibitor, PD184352, and kinase C (PKC) inhibitor H89, and the PKA inhibitor H87, and clonal antibody against COX-2 were purchased from Calbiochem. The CellTiter-96 Aqueous One Solution Cell Assay kit, gel shift assay system, and RT-PCR kit were obtained from Promega. The DNA Master SYBR Green 1 kit was purchased from Applied Science. Antibodies against COX-2 (H-87), and AP-28 (H77) were purchased from Santa Cruz Biotechnology, Inc. All RT-PCR kit components were purchased from PerkinElmer Life Sciences. Human lung carcinoma cells (H1838 and H2106), collagen type 1, and other chemicals were purchased from Sigma unless otherwise indicated.

Reverse Transcriptase PCR—Total RNA was prepared from human lung carcinoma cells using TRIzol reagent (Invitrogen) according to the manufacturer's instructions. To amplify 465-bp EP4 and 200-bp GAPDH cDNA fragments, the sequences of PCR primers (Sigma) were 5'-TCGCGCAAGG-AGCAGAAGGAGAC-3' (for EP4 sense), 5'-GACGGTGGC-GAGAATGAGGAAGGA-3' (for EP4 antisense), 5'-CCATG-GAGAAGGCTGGGG-3' (for GAPDH sense), and 5'-CAA-AGTTGTCATGGATGACC-3' (for GAPDH antisense) according to published data (11, 12). The RT-PCR was carried out as previously described (12). The samples were first denatured at 95 °C for 30 s, followed by 32 PCR cycles, each with temperature variations as follows: 95 °C for 30 s, 60 °C for 30 s, and 72 °C for 30 s. The last cycle was followed by an additional

extension incubation of 7 min at 72 °C. Analysis of amplicons was accomplished on 1% agarose gel containing 0.2 μ g/ μ l ethidium bromide and visualized under UV transilluminator. The densitometric analysis of PCR products was performed by computer software (Bio-Rad Quantity One) and a GS-800 Imaging Densitometer (Bio-Rad) and standardized to the GAPDH product. EP4/GAPDH density bands in control groups were considered as 100%. Values of treatment group EP4/GAPDH ratios are given as percentage of controls. A 100-base pair ladder (Invitrogen) was used as a size standard.

Real Time RT-PCR—This procedure, which is based on the time point during cycling when amplification of the PCR product is first detected, rather than on the amount of PCR product accumulated after a fixed number of cycles, was described previously (12). Final results, which were expressed as *n*-fold differences in EP4 gene expression relative to the GAPDH gene, were calculated using a formula based on a doubling of the product after each cycle (13). The procedures for treatment and total RNA preparation were identical to those described for RT-PCR. All PCR reactions were performed using the LightCycler-FastStart DNA Master SYBR Green 1 kit (Roche Diagnostics, Mannheim, Germany) in the Cepheid SmartCycler (Bio-Rad, Hercules, CA). The cycling conditions were 95 °C for 10 min, 95 °C for 10 s, and 72 °C for 10 s. Each sample was run in triplicate for each data point. No template controls were included.

Small Interfering RNA (siRNA)—siRNAs were synthesized using the siRNA oligodeoxynucleotide (ODN) synthesis protocol. The siRNA sequences in this study were 5'-GCCGCCGCC-ATGGCGGCGGCGGCGGC-3' (for COX-2), and the corresponding siRNA for AP-2 antisense was 5'-CGTCAATTC-ATGGATCGG-3' (−13 to +18 of the AP-2 ODN). The control oligonucleotide, 5'-CAAAGTCT-ATTCGGTCATAATGGCC-3', consisted of similar sequences with scrambled sequences. They were synthesized by Sigma (The Woodlands, TX) according to published data (13, 14). The COX-2 siRNA (catalog ID number 116912) was purchased from Ambion. EP4 siRNA (catalog number M-005714-00) and control nonspecific siRNA oligonucleotides (catalog number D-001206-13-05) were purchased from Dharmacon, Inc. (Lafayette, CO), as described previously (15). For the transfection of ERK and AP-2 ODN, cells were grown to 70% confluence, and a 1 μ M concentration of ERK and 4 μ M AP-2 phosphorothioate ODN mixed with FuGENE 6 transfection reagent per well of serum-free medium was added to the cells for 24 h at 37 °C. COX-2 and EP4 siRNA or control siRNA were transfected using the oligofectamine reagent (Invitrogen) according to the manufacturer's instructions. Briefly, oligofectamine reagent was incubated with serum-free medium for 10 min. Subsequently, a mixture of respective antisense or sense ODN or siRNA was added. After incubation for 15 min at room temperature, the mixture was diluted with medium and added to each well. The final concentration of siRNAs in each well was 100 nM. After culturing for 30 h, cells were washed and resuspended in new culture medium in the presence

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or absence of Fn for an additional 24 h for Western blot analysis, cell growth, and gel mobility shift assays.

Western Blot Analysis—The procedure was performed as previously described (16). Protein concentrations were determined by the Bio-Rad protein assay. Equal amounts of protein from whole cell lysates were solubilized in 2× SDS-sample buffer (125 mM Tris-HCl, pH 6.8, 4% SDS, 20% glycerol, 5–10% 2-mercaptoethanol, and 0.004% bromophenol blue) and separated on SDS-8–10% polyacrylamide gels. The separated proteins were transferred onto nitrocellulose using a Bio-Rad Trans Blot semidry transfer apparatus for 1 h at 25 V, blocked with Blotto (1× TBS (10 mM Tris-HCl, pH 8.0, 150 mM NaCl)) with or without 5% bovine serum albumin, 5% nonfat dry milk, and 0.1% Tween 20 overnight at 4 °C, and washed twice for 5 min with wash buffer (1× TBS and 0.1% Tween 20). Blots were incubated with polyclonal antibodies against COX-2, EP1, EP2, EP3, EP4, Akt, ERK1, ERK2, and their phosphorylated forms and for AP-2, AP-2 α , AP-2 β , and AP-2 δ (1:1000) overnight at 37 °C, washed three times for 5 min with wash buffer, and incubated with a secondary antibody raised against rabbit IgG conjugated to horseradish peroxidase (1:1000; Sigma) for 1 h at room temperature. The blots were washed four times in wash buffer, transferred to freshly made ECL solution (Amersham Biosciences) for 1 min, and exposed to x-ray film. Protein bands were quantified by densitometric scanning using a Bio-Rad GS-800 calibrated densitometer. In controls, antibodies were omitted or replaced by serum IgG.

Cell Viability Assay—NSCLC cells (1.5 × 10⁵) were transfected with EP4 siRNA for 30 h before being seeded into (20 μ g/ml)-coated culture plates. After an additional 48 h in 96-well plates, cells were harvested, and cell viability was determined using a fluorescent cell viability assay (ATP, an indicator of metabolic activity) according to the manufacturer's instructions.

[methyl-³H]Thymidine Incorporation Assay—Cells were transfected with EP4 or control siRNA and then seeded into (20 μ g/ml)-coated culture plates or dmFBS-coated plates for an additional 24 h. Medium was removed, and cells were washed with 1× PBS. Afterward, the attached cells were treated with ice-cold 6% trichloroacetic acid at 4 °C for 20 min and washed once with 6% trichloroacetic acid. Cells were solubilized with 0.1 N NaOH and counted in a liquid scintillation counter in 4 ml of scintillation fluid.

Site-directed Mutagenesis—To prepare site-directed mutants of the promoter, the following oligonucleotides were synthesized: mutated AP-2 (–1529 bp), 5'-GGTTTAAATTGCCCTTGGTGTTCCTCCGATC; mutated AP-2 (–1133 bp), 5'-GCTCGCCTTCCTCCCTCCGCTTTGG; mutated AP-2 (–1000 bp), 5'-GCCTCTGCCAAGTCTACCCGGAGCTCTCG. The lowercase letters indicate mutation, and the underlined letters indicate the AP-2 binding site. The EP4 plasmid constructs containing site-directed mutations of AP-2 cis-acting elements were generated by oligonucleotide-directed muta-

tion using the GeneEditor *in vitro* site-directed mutagenesis system according to recommendations by the manufacturer (Promega). Briefly, double-stranded EP4 promoter plasmid was alkaline-denatured, precipitated, washed, and resuspended in Tris-EDTA buffer. Mutated AP-2 oligonucleotides and selection oligonucleotides were annealed; mutant strands were synthesized, ligated, and transformed into BMH 71-18 *mutS* competent cells. The mutated AP-2 EP4 plasmid was isolated and transformed into JM109 competent cells. Colonies (10–15) were selected and screened for mutants by sequencing using an Applied Biosystems ABI Prism 377 DNA sequencer.

Transient Transfection Assay—The human EP4 wild-type and deletion promoter constructs (pGlep4-1 to -5) ligated to the luciferase reporter gene have been reported previously (18). The EP4 promoter construct contains ~4200 bp of the 5'-flanking region of the mouse EP4 receptor gene connected to the pGL3 basic luciferase reporter vector (Promega). NSCLC cells were seeded at a density of 5 × 10⁵ cells/well in 6-well dishes and grown to 50–60% confluence. For each well, 2 μ g of the above plasmid DNA and 0.2 μ g of the internal control pRL-TK-luciferase reporter vector were cotransfected using the EUGENE 6 lipofection reagent as described in our earlier work (18). Cells were treated with or without (20 μ g/ml) for 1 h before exposure to (20 μ g/ml each)-coated culture plates and measured using the dual luciferase assay. The luciferase activities were normalized sequentially with changes in cell number and luciferase activity were normalized with changes in cell number within the same sample.

Gel Mobility Shift Assay—Electrophoretic mobility shift experiments were performed as described before (18). The oligonucleotides used as probes were as follows: wild type Sp1 (5'-CTCCCCGCCCAAGCCTGG-3'), mutant Sp1 (5'-CTCCCTCCCAAGCCTGG-3'), wild type C/EBP (5'-GATAATTAAGAAATGAT-3'), mutant C/EBP (5'-GATCTTAAGAAATGA-3'); wild-type AP-2 (5'-TCCTCCCCGCCCTCCGC-3'), and mutant AP-2 (5'-TCCTCTTGCCTCCGC-3'), which is based on the EP4 promoter sequences (18) and consensus AP-2 α binding motif (5'-GATCGAACTGACGCCCGCGGCCCGT-3'). The complementary oligonucleotides were annealed and purified following the manufacturer's protocol. The Sp1, C/EBP, and AP-2 oligonucleotides were end-labeled with [γ -³²P]ATP using T4 polynucleotide kinase as recommended by the manufacturer. Nuclear proteins (5 μ g) were first incubated under binding conditions (10 mM HEPES, 10 mM Tris-HCl (pH 7.9), 50 mM KCl, 0.1 mM EDTA, 1 mM dithiothreitol, 12% (v/v) glycerol, and 2 μ g of poly(dI-dC)) for 10 min, followed by the addition of [γ -³²P]ATP probe for another 20 min at room temperature in a final volume of 20 μ l in the presence or absence of AP-2 antibodies (2 μ g/ μ l). For cold competition, a 100-fold excess of the respective unlabeled consensus oligonucleotides was incubated for 15 min before adding the

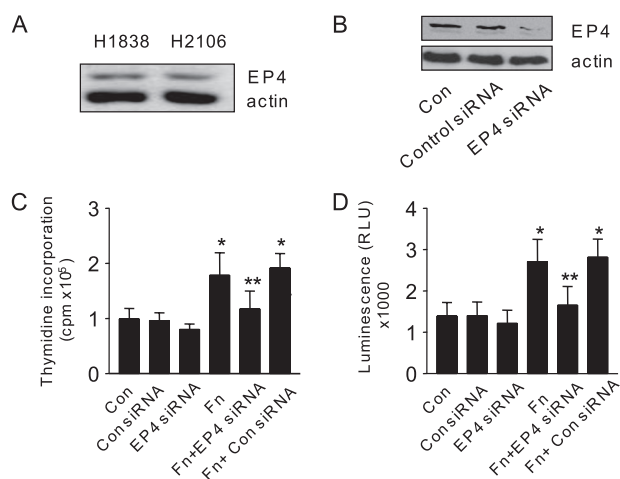


FIGURE 1. EP4 siRNA blocks Fn-induced lung carcinoma cell growth. A, expression of EP4 in NSCLC cells. Cellular proteins were isolated from H1838 and H2106 cells, followed by Western blot analysis for EP4 protein. B, blockade of EP4 production by EP4 siRNA. Cellular protein was isolated from H1838 cells transfected with control or EP4 siRNA pool (100 nM each) for 30 h and then subjected to Western blot analysis for EP4 protein. C, effect of EP4 siRNA on Fn-induced NSCLC cell growth. H1838 cells were transfected with control or EP4 siRNA pool for 30 h before exposing the cells to culture plates coated with Fn (20 μ g/ml) and incubation with 1 μ Ci/ml [methyl-³H]thymidine for an additional 24 h as indicated. Data are expressed as mean \pm S.D. of at least three independent experiments. D, effect of EP4 siRNA on Fn-induced NSCLC cell growth. H1838 cells were transfected with control or EP4 siRNA pool for 30 h before exposing the cells to culture plates coated with Fn (20 μ g/ml) for an additional 48 h. Afterward, viable cell number was determined by the CellTiter-Glo Luminescent cell viability assay. Data are presented as means \pm S.D. *, significant difference from control; **, significant difference from combination treatment as compared with Fn alone. Con indicates untreated control cells.

probe. The same amount of probe was used with the probe was used with the same binding conditions. The protein-DNA complex was resolved on a 4.5% polyacrylamide gel and transferred to a nitrocellulose membrane (10 \times Tris/glycine: Tris-base, 3.92 g), and H₂O added up to 100 ml. The membrane was dried and subjected to autoradiography.

Statistical Analysis—All experiments were performed in triplicate. All data from Western blot analysis, luciferase activity assays, RT-PCR, and Western blot analysis were expressed as mean \pm S.D. The data presented in some figures are from representative experiments, which was qualitatively similar in the replicate experiments. Statistical significance was determined with Student's *t* test (two-tailed) comparison between two groups of data sets. The asterisks shown in the figures indicate significant differences of experimental groups in comparison with the corresponding control condition (*p* < 0.05; see figure legends).

RESULTS

Effect of Fn on EP4 Gene Expression in Human Lung Carcinoma Cells—We previously showed that mRNAs encoding for the four PGE₂ receptor subtypes are present in human NSCLC cells (17). Consistent with this, we show that the PGE₂ receptor subtype EP4 protein is expressed in the two NSCLC cell lines studied (Fig. 1A). We also demonstrated that Fn stimulated NSCLC cell growth (4). Here, we examined whether blockade

of EP4 could influence the effects of Fn on cell growth. We first depleted EP4 from cells in culture using siRNA approaches. Treatment of H1838 cells with EP4 siRNA blocked EP4 production. Levels of EP4 were unchanged in cells transfected with control siRNA oligonucleotides (Fig. 1B). To determine if EP4 siRNA can block Fn-induced lung carcinoma cell growth in our system, H1838 cells were transfected with EP4 siRNA duplexes. Afterward, the cells were plated onto Fn-coated culture plates for an additional 24 h. As shown in Fig. 1C, the EP4 siRNA duplexes inhibited Fn-induced H1838 cell proliferation, whereas the control siRNA had no effects as determined by the [³H]thymidine incorporation assay. Similar results were also found by cell viability assays (Fig. 1D).

Since EP4 has been shown to be involved in human lung carcinoma biology, we tested if Fn can affect its expression. H1838 cells exposed to Fn showed increased EP4 protein levels in a time- and dose-dependent manner with maximal increases noted in 24 h at concentrations of 20 μ g/ml (Fig. 2, A and C). Similar results were also observed in an additional NSCLC cell line (H2106) (Fig. 1, B and C). Fn did not significantly stimulate EP4 mRNA levels in a time- or dose-dependent manner, with maximal increases noted in 24 h at concentrations of 20 μ g/ml Fn as determined by RT-PCR (Fig. 2B). Fn also stimulated collagen type 1, another marker for cell growth (Fig. 2F). This result was consistent with our previous findings (Fig. 2G). Of note, cells transfected with EP4 siRNA did not adhere well to Fn-coated plates. In order to determine if the effect of Fn on EP4, we examined the effect of Fn on other EP receptors. We found that Fn did not affect on other EP receptors. We then tested the effect of Fn on EP4 expression in H2106 cells (not shown). We found that Fn stimulated EP4 expression, and COX-2, but Not PKC and PKA (Fig. 2, D and E). To this end, we treated H1838 cells with or without anti-integrin α 5 β 1 (MAB1969) or anti-integrin α 2 β 1 (MAB1967; 25 μ g/ml each) for 2 h before exposing the cells to culture plates coated with Fn. We found that Fn-induced EP4 protein was eliminated in the presence of anti- α 5 β 1 antibodies, whereas the anti- α 2 β 1 antibodies had no effect (Fig. 3A). This suggests that α 5 β 1 integrin mediates Fn-induced EP4 expression.

Fn has been shown to affect kinase signaling pathways in several studies (21–23). We previously demonstrated that Fn activated ERK and PI3K/Akt signaling pathways in NSCLC cells (4, 23, 24). Here, we examined if inhibition of these kinase signal pathways diminished or abrogated the effects of Fn on EP4. The specific inhibitors of ERK (PD98095 (25 μ M)) and of PI3K (wortmannin (100 nM)) significantly blocked Fn-induced EP4 protein levels in H1838 cells (Fig. 3B). We previously demonstrated that mTOR signals were mediating some of the effects of Fn on NSCLC cell growth (23). However, we found that rapamycin, an inhibitor of mTOR, had no effect on inhibition of Fn-induced expression of EP4 protein (Fig. 3C). Also, the inhibitor of PKA (H89 (10 μ M)) or of PKC (calphostin C (*Cal*; 0.5 μ M)) had no effects (Fig. 3, D and E). In addition, we showed

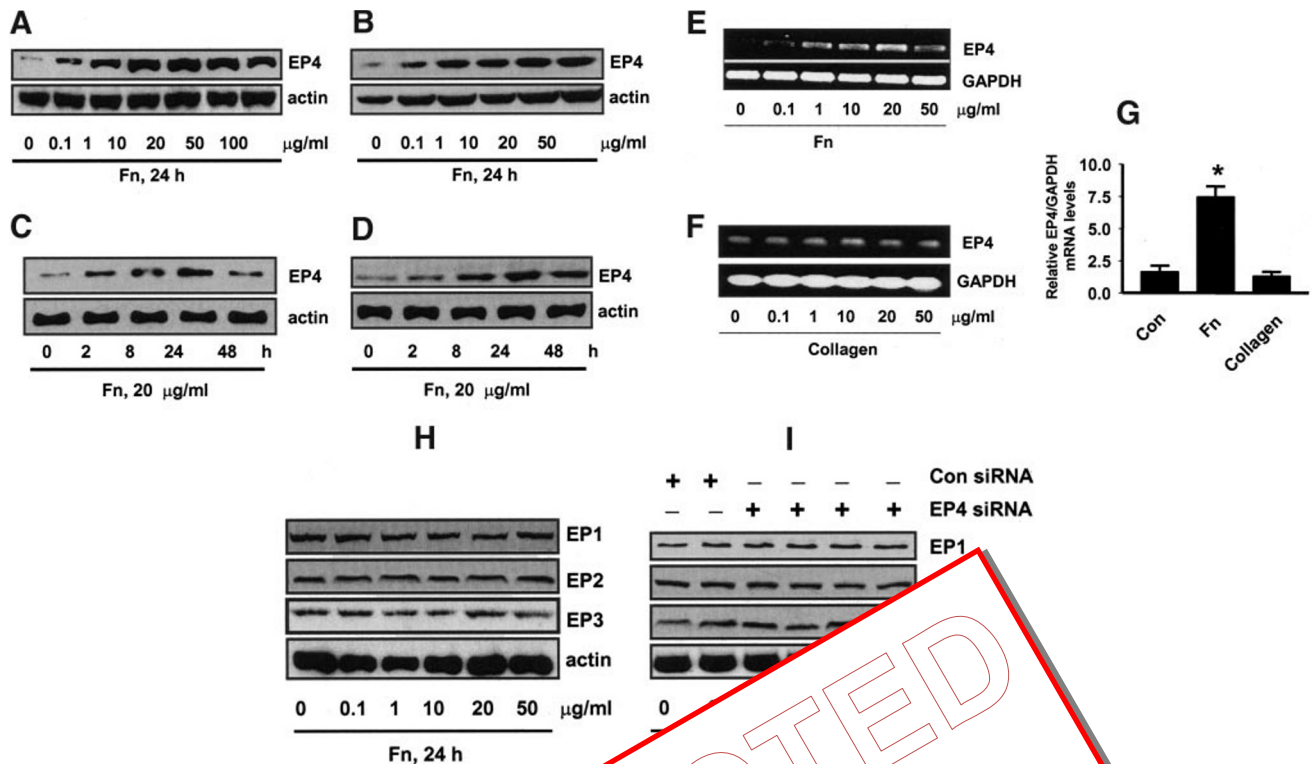


FIGURE 2. The effect of Fn on EP4 gene expression in human lung cancer cells. H1838 (A) and H2106 (B) cells were isolated from human lung cancer and cultured on plates coated with increasing concentrations of Fn (20 $\mu\text{g/ml}$) for the indicated time period. Afterward, Western blots were performed to determine the effect of Fn on EP4 expression. C and D, time-dependent effect of Fn on EP4 expression. Cells were treated with 20 $\mu\text{g/ml}$ of Fn for the indicated time period. Afterward, Western blots were performed to determine the effect of Fn on EP4 expression. E and F, dose-dependent effect of Fn and collagen type 1 on EP4 expression. Cells were treated with increasing concentrations of Fn or collagen type 1 (20 $\mu\text{g/ml}$) for the indicated time period. Afterward, Western blots were performed to determine the effect of Fn on EP4 expression. G, relative EP4/GAPDH mRNA levels in H1838 cells treated with control (Con), Fn (20 $\mu\text{g/ml}$), or collagen type 1 (20 $\mu\text{g/ml}$) for 24 h. * indicates significant difference from control ($p < 0.05$). Con, control; Fn, fibronectin; collagen, collagen type 1. H, time-dependent effect of Fn on EP4 expression. Cells were treated with 20 $\mu\text{g/ml}$ of Fn for the indicated time period. Afterward, Western blots were performed to determine the effect of Fn on EP4 expression. I, effect of EP4 siRNA on EP4 expression. Cells were treated with control (Con) or EP4 siRNA, with or without Fn (20 $\mu\text{g/ml}$) for 24 h. Afterward, Western blots were performed to determine the effect of Fn on EP4 expression. Actin indicates untreated control cells. GAPDH indicates untreated control cells. EP1, EP2, and EP3 expression levels were determined by real time PCR analysis. Actin served as internal control for protein expression. Cellular proteins were determined by Western blot for EP4, EP1, EP2, and EP3. GAPDH was used as a loading control for normalization purposes. Data represent the mean \pm SD of three independent experiments. * indicates significant difference from control ($p < 0.05$).

that the inhibitor of PI3K, wortmannin, blocked the phosphorylation of ERK1/2 by Fn and found that, as expected, the inhibitors of ERK1/2, PD98095, had no effect on the phosphorylation of Akt (Fig. 3G). This suggests that the PI3K/Akt pathway was involved in the regulation of EP4 transcriptional activity. Similar results were obtained with H2106 cells (not shown).

Fn Increased EP4 Promoter Activity. To further examine whether the effects of Fn on EP4 expression occur at the transcriptional level, the EP4 promoter contains multiple transcription factor binding sites, including NF- κ B, NF-IL6 (C/EBP), Sp1, and AP-2, among others (Fig. 4A). These sites have been shown to be differentially responsive to various stimuli (15, 18, 25, 26). We found that H1838 cells, transfected with the full-length wild-type EP4 promoter (−4200/−116 bp) luciferase reporter construct, exposed to Fn showed increased promoter activity. Collagen type 1 had no effect on the wild-type promoter (Fig. 4B). The Fn-induced EP4 promoter activity was slightly reduced in one EP4 deletion reporter construct (−1555/−116 bp). There was no response to Fn with another EP4 deletion reporter construct (−992/−116 bp) (Fig. 4B), indicating that the region between −1555 and −992 bp in the EP4 promoter played an important role in stimulation of EP4 gene expression in response to Fn. We also tested whether

ERK1/2 and PI3K pathways were involved in EP4 transcriptional activity. We tested the effect of Fn and found that, as expected, the inhibitors of ERK1/2, PD98095, and PI3K partially reduced Fn-induced EP4 promoter activities (Fig. 4C). Similar results were obtained with H2106 cells (not shown).

Regulation of AP-2, Sp1, and C/EBP in the EP4 Promoter by Fn in Human Lung Carcinoma Cells. To further explore the role of Fn in regulation of EP4 promoter activity, electrophoretic mobility shift assays were performed to identify the transcription factors involved. We found that H1838 cells treated with Fn for 24 h showed a significant increase in AP-2 (Fig. 5C), but we observed no effect in C/EBP (Fig. 5A) and a slight increase in Sp1 (Fig. 5B) nuclear protein binding activities as compared with solvent controls. In contrast, collagen type 1 (20 $\mu\text{g/ml}$) had no effect. We also tested if the activation of ERK signals by Fn was involved in the induction of AP-2 binding activity. ERK1/2 antisense oligonucleotide (1 μM) completely blocked ERK1 and reduced ERK2 production. The levels of ERK1, ERK2, and actin were unaffected in untransfected cells and in cells treated with ERK sense oligonucleotides (Fig. 5D, top). The antisense ERK ODN prevented the Fn-induced AP-2 binding activity (Fig. 5D, bottom). Similarly, the inhibitors of ERK (PD98095) and of PI3K (wortmannin) also blocked Fn-

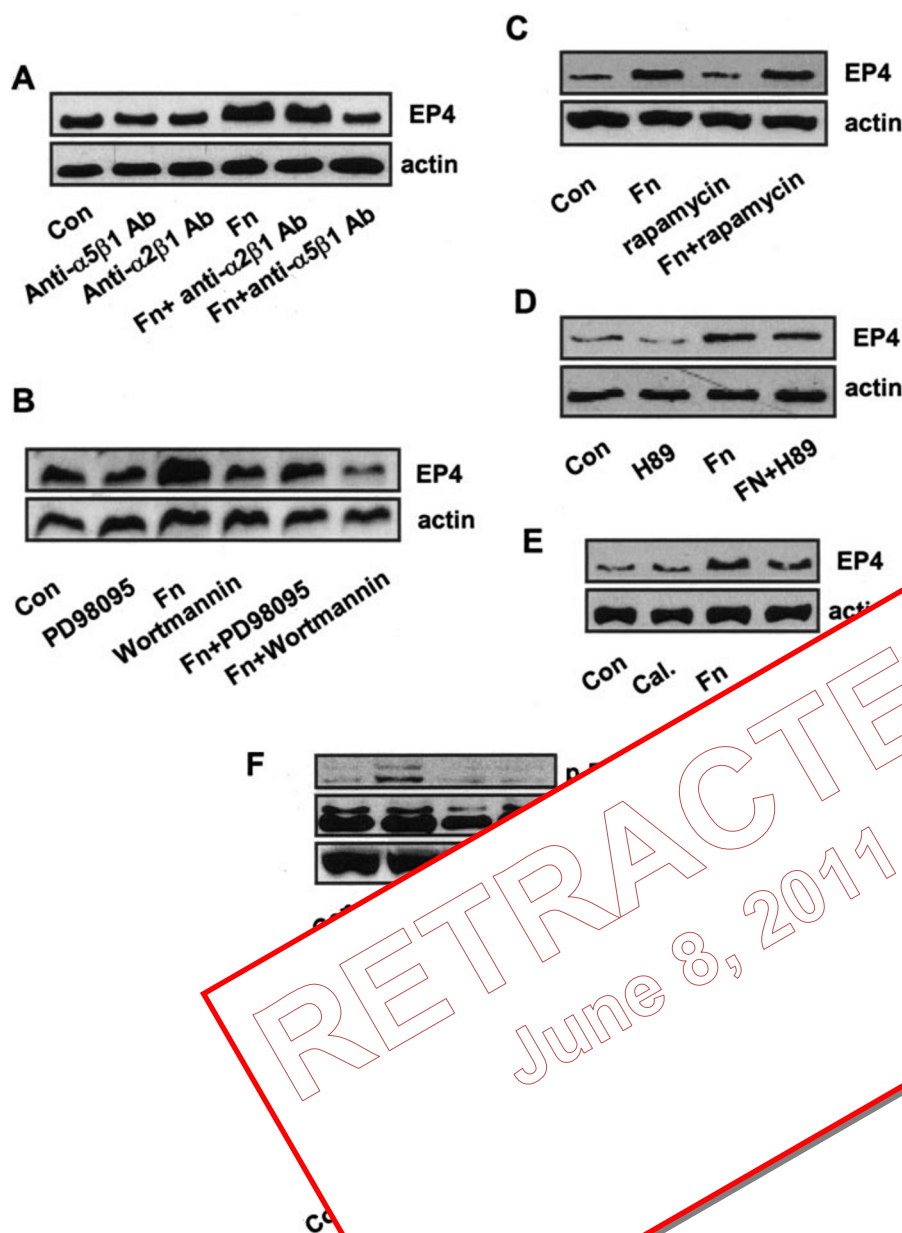


FIGURE 3. Involvement of $\alpha 5 \beta 1$ integrin, PI3K, and PKA pathways in the induction of EP4 by Fn. A, effect of anti- $\alpha 5 \beta 1$ antibodies on Fn-induced EP4 protein levels. Cellular protein was isolated from H1838 cells cultured for up to 2 h in the presence or absence of anti- $\alpha 5 \beta 1$ and anti- $\alpha 2 \beta 1$ antibodies (25 μ g/ml each) before exposing the cells to culture plates coated with Fn for an additional 24 h and then subjected to Western blot analysis for EP4. B, effect of inhibitors of ERK1/2 and PI3K on Fn-induced EP4 protein levels. Cellular protein was isolated from H1838 cells cultured for up to 2 h in the presence or absence of PD98095 (25 μ M) or wortmannin (100 nM) before exposing the cells to Fn-coated culture plates for an additional 24 h and then subjected to Western blot analysis for EP4. C, effect of mTOR inhibitor on Fn-induced EP4 protein levels. Cellular protein was isolated from H1838 cells cultured for up to 4 h in the presence or absence of rapamycin (10 nM) before exposing the cells to Fn-coated culture plates for an additional 24 h and then subjected to Western blot analysis for EP4 protein. D, effect of PKA inhibitors on Fn-induced EP4 protein levels. Cellular protein was isolated from H1838 cells cultured for up to 2 h in the presence or absence of H89 (10 μ M) before exposing the cells to Fn-coated culture plates for an additional 24 h and then subjected to Western blot analysis. E, effect of PKC inhibitors on Fn-induced EP4 protein levels. Cellular protein was isolated from H1838 cells cultured for up to 2 h in the presence or absence of calphostin C (0.5 μ M) before exposing the cells to Fn-coated culture plates for an additional 24 h, then subjected to Western blot analysis. F, effect of PI3K inhibitor on ERK1/2. Cellular protein was isolated from H1838 cells cultured for 1 h in the presence or absence of wortmannin (100 nM) before exposing the cells to Fn (20 μ g/ml) coated onto the culture plates for an additional 1 h. Afterward, Western blot was performed to detect phosphorylated ERK1/2 and total ERK1 and ERK2 proteins. G, effect of ERK inhibitor on Akt. Cellular protein was isolated from H1838 cells cultured for 1 h in the presence or absence of PD98095 (25 μ M) before exposing the cells to Fn (20 μ g/ml) for an additional 1 h. Afterward, Western blot was performed to detect phosphorylated Akt and total Akt protein. Actin served as internal control for normalization purposes. Con, untreated control cells.

induced AP-2 binding activity (Fig. 5E). The addition of an AP-2 α antibody induced a super-shift band, whereas AP-2 β and AP-2 γ antibodies had no effects (Fig. 5F). The specific bands for AP-2, Sp1, and C/EBP were attenuated by a 100-fold molar excess of unlabeled wild-type oligonucleotides but were not inhibited by the mutated unlabeled oligonucleotides (Fig. 5, *Mut*). Oligonucleotides containing a mutated AP-2 (*Mut AP-2*), Sp1 (*Mut Sp1*), or C/EBP (*Mut C/EBP*) site were end-labeled with [γ - 32 P]ATP and used as another control to confirm the binding specificity. Similar results were obtained with H2106 cells (not shown).

Role of Transcription Factor AP-2 in Fn Induction of EP4 and Cell Growth—We further tested the role of AP-2 in mediating Fn-induced EP4 expression in human H1838 cells by using the antisense approach. We showed that the antisense ODN completely inhibited the production of EP4 protein, whereas the control sense ODN had no effect (Fig. 6A). Consistent with these findings, we found that cells transfected with AP-2 antisense ODN resulted in inhibition of Fn-induced cell growth as determined by [3 H]thymidine incorporation assay (Fig. 6B) and inhibition of Fn-stimulated effect on EP4 promoter activities (Fig. 6C). The control sense ODN had no effect. By using site-directed mutagenesis, we found that the stimulatory effect of Fn on EP4 promoter activity was lost with EP4 promoter constructs in which one AP-2 site was mutated (–1000 bp) (Fig. 6D).

PI3K, ERK, and $\alpha 5 \beta 1$ Integrin Signaling Are Involved in Fn-induced AP-2 Protein Expression—We also examined the effects of Fn on AP-2 protein expression, and, consistent with the gel shift experiment results,

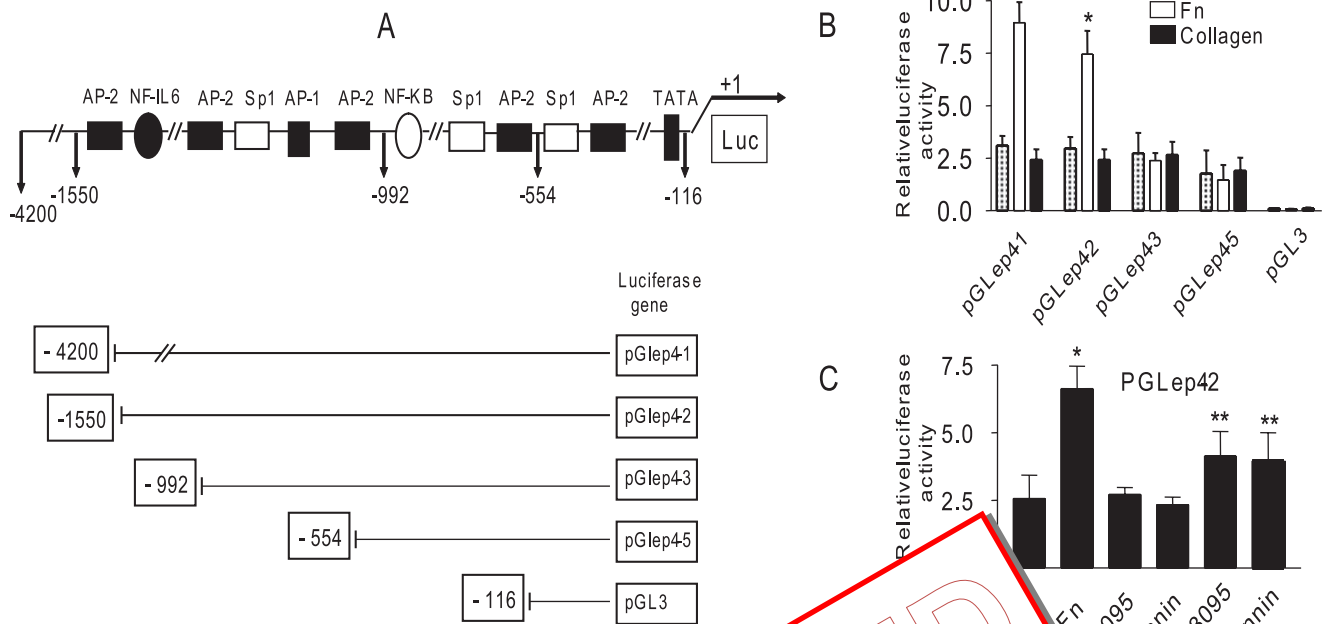


FIGURE 4. Fn stimulates EP4 promoter activity. A, promoter map and schematics are presented. These regions contain several transcription factor binding sites that are known to regulate EP4 promoter activity. H1838 lung cancer cells (1×10^5 cells) were transfected with the indicated promoter reporter constructs and treated with "Experimental Procedures" for 24 h and then treated as indicated. B, effects of Fn on EP4 promoter activity. H1838 cells were transfected with the indicated promoter reporter constructs and treated with "Experimental Procedures" for 24 h and then treated with Fn (10 μ M) for an additional 24 h. The ratio of firefly luciferase to Renilla luciferase activity was determined as the mean \pm S.D. of at least four independent experiments. C, effects of inhibitors of ERK1/2 or PI3K on EP4 promoter activity. H1838 cells were transfected with the indicated promoter reporter constructs and treated with "Experimental Procedures" for 24 h and then treated with PD98095 (25 μ M) and wortmannin (100 nM) for an additional 24 h. The ratio of firefly luciferase to Renilla luciferase activity was determined as the mean \pm S.D. of at least four independent experiments. * indicates $p < 0.05$ compared with Fn alone; ** indicates $p < 0.01$ compared with Fn alone.

we found that Fn induced AP-2 δ protein production and AP-2 δ protein production was inhibited by PD98095 and AP-2 δ protein production was inhibited by PD98095 (Fig. 7C) and the anti- α 5 β 1 antibody (Fig. 7D). These results suggest that AP-2 α protein expression is involved in the regulation of EP4 expression. Similar results were obtained with H2106 cells (not shown).

COX-2 Signaling Is Involved in the Regulation of EP4 Protein Expression—Having established that Fn stimulates NSCLC cell proliferation through COX-2 and via EP4 (this study), we explored whether COX-2 signals could also affect EP4 expression. We found that COX-2 siRNA blocks endogenous COX-2 protein production; no changes were noted in cells transfected with control siRNA (Fig. 8A). COX-2 siRNA significantly abrogated Fn-induced EP4 expression; the control siRNA had no effect (Fig. 8A). We also found that, under the same conditions, COX-2 siRNA abrogated the effect of Fn on AP-2 α protein expression (Fig. 8B), whereas the control siRNA had no effect. These studies suggest that COX-2 activation not only promotes PGE₂ secretion but may also amplify its effects by stimulating EP4 expression. Therefore, we tested whether PGE₂ can stimulate EP4 expression. We observed that dmPGE₂ (0.1 μ M) also enhanced AP-2 α and EP4 protein expression, with greater stimulation seen when Fn and dmPGE₂ were used concomitantly (Fig. 8, C and D). Finally, similar to the effect of Fn

DISCUSSION

Fn is a heterodimeric extracellular matrix glycoprotein implicated in a number of physiological events during embryogenesis, angiogenesis, thrombosis, and inflammation (27–29). Fn expression is increased in lung carcinomas, particularly in non-small cell lung carcinoma (3, 29–31). Also, the adhesion of lung carcinoma cells to Fn enhances tumorigenicity and confers resistance to apoptosis induced by standard chemotherapeutic agents (32). Previously, we found that Fn stimulates human lung carcinoma cell growth *in vitro* by increasing expression of COX-2 and PGE₂ biosynthesis (4). We also demonstrated that all four PGE₂ receptors are expressed in NSCLC cells studied (17). Based on these data, we predicted that Fn stimulated NSCLC growth through one or more of these four EP receptors capable of recognizing PGE₂. The development of aberrant crypt foci and putative preneoplastic lesions in the colon was decreased in the EP4 knock-out mice (7). Blockade of EP4 production also mediated inhibition of NSCLC cell inva-

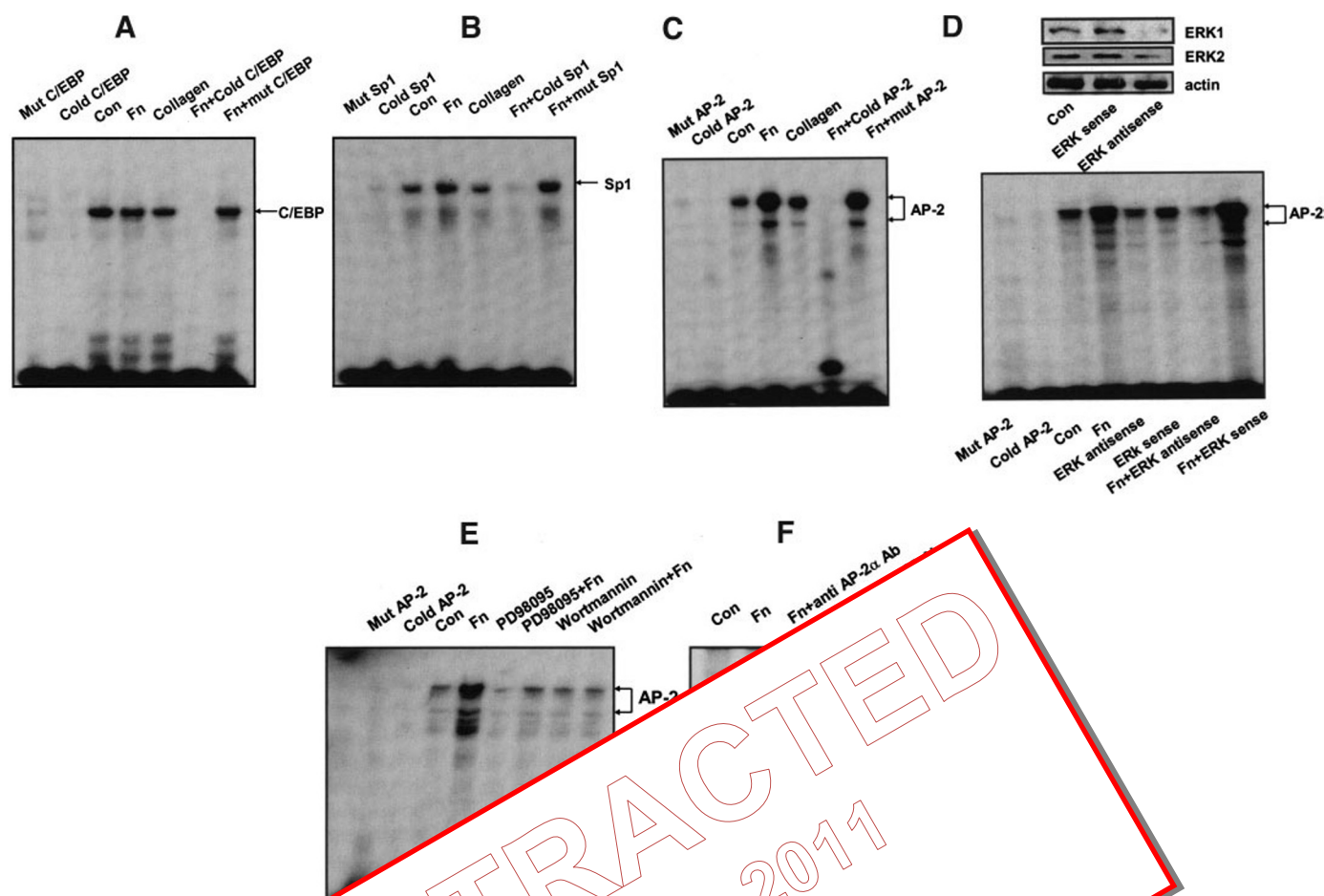


FIGURE 5. Electrophoretic mobility shift assays (EMSA) showing the effect of Fn on transcription factor binding. A–C, effect of Fn on C/EBP (A), Sp1 (B), and AP-2 (C) sites were end-labeled with [γ - 32 P]ATP and incubated with nuclear extracts from H1838 cells treated with PD98095 (5 μ M) for an additional 24 h. D, ERK1/2 phosphorylation in H1838 cells transfected with 1 μ M ERK antisense or sense oligonucleotide for 24 h before exposure to Fn (20 μ g/ml) for an additional 24 h. E, the inhibitors of ERK1/2 and PI3K abrogated the Fn-induced AP-2 binding activity. F, anti-AP-2 antibody (5 μ g) abrogated the Fn-induced AP-2 binding activity. AP-2 (Cold AP-2) oligonucleotide (1 μ M) was added to the reaction mixture. AP-2 (Mut AP-2) site that were end-labeled with [γ - 32 P]ATP and incubated with nuclear extracts from H1838 cells treated with PD98095 (5 μ M) for an additional 24 h. Con, untreated control cells.

siveness (10). Here, we reported that Fn induced the expression of EP4, whereas it had no effect on other EP receptors. A similar lack of changes in the other EP receptors was observed when EP4 was knocked down by EP4 siRNA. This finding indicates that Fn selectively targets EP4 receptor subtypes in NSCLC cells. In view of the above, we focused on EP4 and explored the mechanisms involved.

First, we demonstrated that EP4 siRNA antagonized Fn-induced lung carcinoma cell growth, suggesting a direct role for EP4 in mediating this process in our system. This is consistent with data from others who reported that EP4 antisense oligonucleotides diminished EP4 protein expression and abolished the PGE₂-stimulated production of cAMP and blocked the ability of PGE₂ to augment release of immunoreactive substance P and calcitonin gene-related peptide in sensory neurons (33). The EP4 antagonist, ONO-AE3-208, and the EP4 siRNA have

been shown to inhibit extracellular matrix-induced metalloproteinase-9 expression in macrophages (34). More recently, EP4 antagonists inhibited breast cancer cell growth and reduced breast, lung, and colon cancer metastasis, suggesting that blockade of EP4 may be an alternative approach to the use of COX-2 inhibitors to prevent tumor metastasis (35, 36).

In this study, we confirm that Fn increased EP4 gene expression in NSCLC cells, whereas collagen type 1 had no effect. This suggested that Fn may induce NSCLC growth not only by stimulation of PGE₂ but may also enhance this process by inducing the expression of EP4. The connection of Fn and EP4 expression has never been reported in lung cancer cells, although an antagonist of the PGE₂ receptor EP1 has been shown to decrease Fn synthesis in a rat diabetic nephropathy model (37). Also, PGE₂-accelerated ProNectin F(TM) (a proteolytic fragment of Fn)-dependent adhesion was mediated through coop-

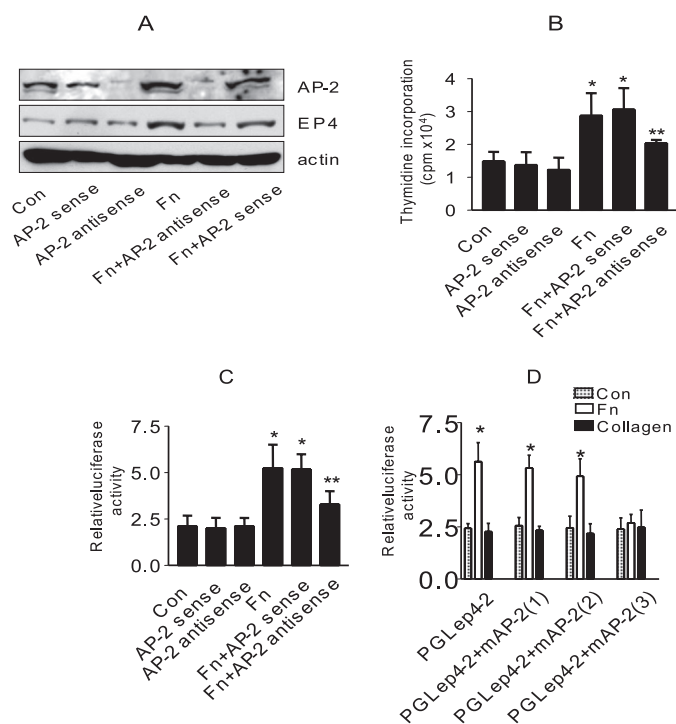


FIGURE 6. The role of transcription factor AP-2 in Fn induction of EP4.

A, AP-2 antisense blocked Fn-induced EP4 protein expression. Cellular protein was isolated from H1838 cells transfected with AP-2 antisense sense ODN (4 μ M) for 30 h before exposing the cells to the platelet-derived growth factor (PDGF) for an additional 24 h. Afterward, Western blot analysis was performed to examine for AP-2 and EP4 protein. B, AP-2 antisense ODN blocked Fn-induced cell proliferation. H1838 cells transfected with AP-2 antisense ODN (4 μ M) for 30 h before exposing the cells to PDGF with 20 μ g/ml Fn and incubation with 1 μ Ci [³H]thymidine for an additional 24 h. The bars represent the mean \pm SD of three independent experiments for each condition. C, AP-2 antisense ODN blocked Fn-induced cell proliferation. H1838 cells transfected with AP-2 antisense ODN (4 μ M) for 30 h before exposing the cells to PDGF with 20 μ g/ml Fn and incubation with 1 μ Ci [³H]thymidine for an additional 24 h. The bars represent the mean \pm SD of three independent experiments for each condition. D, effects of AP-2 promoter constructs on cell proliferation. H1838 cells were transfected with various AP-2 promoter constructs (–1529, –1133, and –1011) together with the firefly luciferase reporter construct (–1529, –1133, and –1011) under “Experimental Procedures.” The cells were then exposed to PDGF for an additional 24 h. The bars represent the mean \pm SD of three independent experiments for each condition. **, significance compared with controls. Con, untreated.

erative activation of EP3 and EP4 in human lung carcinoma P-815 cells (38).

We previously demonstrated that Fn stimulated lung carcinoma cell growth through its receptor $\alpha 5 \beta 1$, since anti- $\alpha 5 \beta 1$ antibodies eliminated the mitogenic response (4). Here, the Fn effect on EP4 protein levels was blocked by anti- $\alpha 5 \beta 1$ antibodies but not by anti- $\alpha 2 \beta 1$ antibodies, indicating that the integrin receptor $\alpha 5 \beta 1$ mediated this regulation. In order to elucidate the mechanism(s) involved in Fn induction of EP4, we attempted to delineate the signaling pathways involved in induction of EP4 expression in lung carcinoma cells in response to Fn treatment. Data from our laboratory (4, 21, 24, 39, 40) and data of others (3, 41) have demonstrated that adhesion to Fn activates several kinase signaling pathways, including ERK, PI3K, PKC, and PKA. We found that inhibitors of the PI3K kinase and ERK prevented Fn-induced EP4 protein expression, suggesting that the activation of these dual kinase signaling

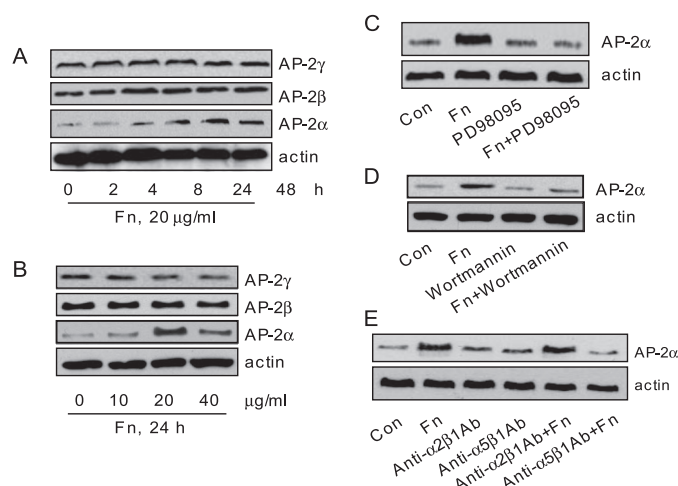


FIGURE 7. The role of PI3K, ERK, and $\alpha 5 \beta 1$ integrin in Fn-induced AP-2 α expression. A, time-dependent effect of Fn on AP-2 protein expression. Cellular protein was isolated from H1838 cells exposed to culture plates coated with Fn (20 μ g/ml) at the indicated time periods. Afterward, Western blot analysis was performed using polyclonal antibodies against AP-2 α , AP-2 β , and AP-2 δ . B, dose-dependent effect of Fn on AP-2 protein expression. Cellular proteins were isolated from H1838 cells exposed to different concentrations of Fn (20 μ g/ml) for the indicated time periods. Afterward, Western blot analysis was performed using polyclonal antibodies against AP-2 α and AP-2 δ . C, effect of ERK inhibitor PD98095 on Fn-induced AP-2 α expression. Cellular protein was isolated from H1838 cells exposed to 20 μ g/ml Fn for 2 h before exposing the cells to PD98095 (10 μ M) for an additional 24 h. Afterward, Western blot analysis was performed using polyclonal antibodies against AP-2 α and actin. D, effect of PI3K inhibitor wortmannin on Fn-induced AP-2 α expression. Cellular protein was isolated from H1838 cells exposed to 20 μ g/ml Fn for 2 h before exposing the cells to wortmannin (0.1 μ M) for an additional 24 h. Afterward, Western blot analysis was performed using polyclonal antibodies against AP-2 α and actin. E, effect of anti- $\alpha 5 \beta 1$ antibody on Fn-induced AP-2 α expression. Cellular protein was isolated from H1838 cells treated with anti- $\alpha 5 \beta 1$ antibody (1 μ g/ml) for 2 h before exposing the cells to 20 μ g/ml Fn for an additional 24 h. Afterward, Western blot analysis was performed using polyclonal antibodies against AP-2 α and actin. Con, untreated.

pathways were involved in the up-regulation of EP4 expression. We previously demonstrated that Fn up-regulated COX-2 expression in lung carcinoma cells (4). We found that blockade of the ERK signal pathway and that blockade of PI3K abrogated Fn-induced COX-2 expression (4). Together, these results suggested a strong connection between COX-2, ERK, and PI3K signaling pathways. However, ERK played no role in the increase of EP4 expression induced by peroxisome proliferator-activated receptor β / δ activation (15), suggesting the existence of independent pathways that differ according to the stimulus. Fn activates MMP-9 via the ERK and PI3K/Akt signaling pathways in NSCLC and ovarian cancer cells (24, 42, 43). The inhibitor of PI3K, wortmannin, blocked the effect of Fn on stimulation of ERK phosphorylation, indicating cross-talk between the PI3K and ERK1/2 pathways in NSCLC cells. The cross-talk between these kinases has been reported in other cell systems as well (44, 45). In contrast, our data indicated that mTOR, PKC, and PKA signaling pathways are not involved in the up-regulation of the EP4 gene induced by Fn, although the latter two kinases were involved in prostaglandin E receptor expression in other studies (46, 47). This might suggest distinct effects, depending on the stimulant and cells studied.

EP4 has been shown to be regulated at the level of gene transcription in different cell types (21, 47). We found that Fn, not collagen type 1, increased EP4 promoter activity. Furthermore, the region between –1555 and –992 was demonstrated to play

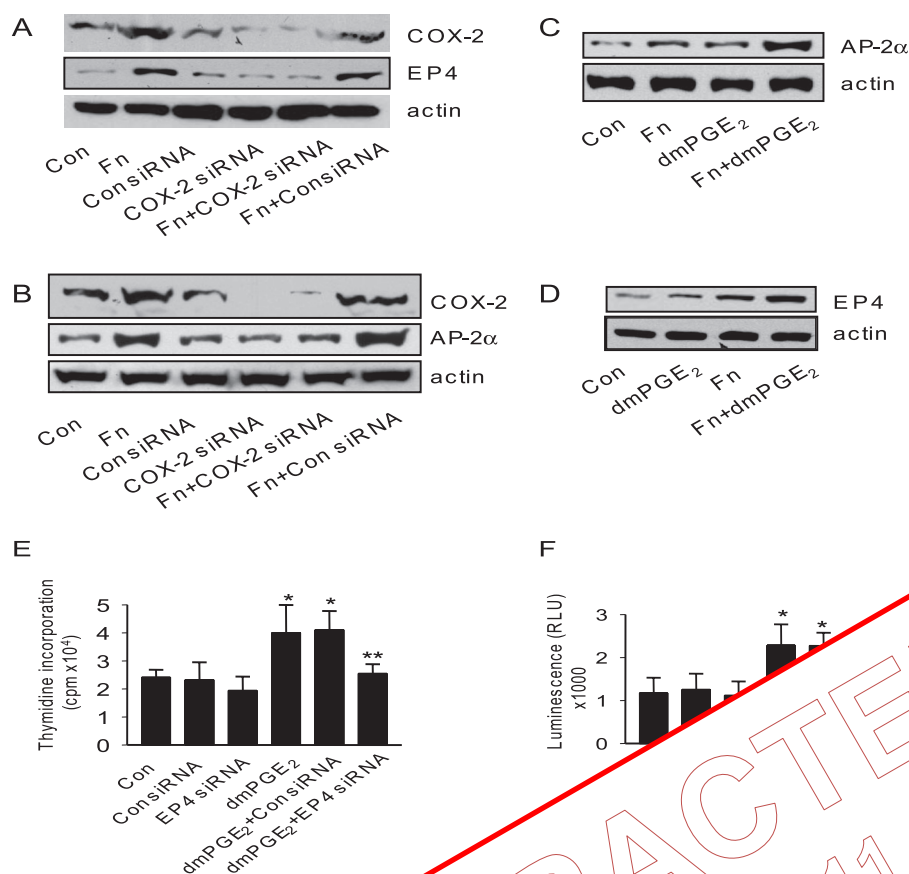


FIGURE 8. The role of COX-2 signaling in mediating fibronectin-induced EP4 expression and inducing lung carcinoma cell growth. H1838 cells were transfected with control or COX-2 siRNA (100 nm each) for 30 h before exposing the cells to 1 μ M Fn for an additional 24 h and then subjected to Western blot analysis. **A**, effect of Fn on COX-2 and EP4 protein expression. **B**, effect of COX-2 siRNA on COX-2 and AP-2 α protein expression. **C**, effect of PGE₂ on AP-2 α protein expression. **D**, effect of PGE₂ on EP4 protein expression. **E**, effect of PGE₂ on cell growth. **F**, effect of PGE₂ on EP4 promoter activity. Cells were transfected with control or EP4 siRNA (100 nm) for 30 h before exposing the cells to 1 μ M Fn for an additional 24 h. Afterward, Western blot analysis was performed for COX-2, EP4, AP-2 α , and actin. **E** and **F**, effect of PGE₂ on cell growth and EP4 promoter activity, respectively. Data are expressed as mean \pm S.D. of at least three independent experiments. **p* < 0.05, significant difference from control; ***p* < 0.01, significant difference from control. Con, untreated control cells.

a major role. The results showing that both PD98095 and wortmannin partially prevented Fn-stimulated EP4 promoter activity further suggested a role for PI3K and ERK kinase signaling pathways in mediating Fn up-regulation of EP4 gene expression. Several transcription factor binding sites within regions of the EP4 promoter have been characterized, including regulatory elements for AP-2, C/EBP, Sp1, and others (15, 18, 25). We showed that treatment of H1838 cells with Fn significantly increased protein binding activities of AP-2 in the EP4 promoter, whereas it had little effect on C/EBP and Sp1. This, together with the supershift assay results, indicated that AP-2 α binding to the AP-2 site was necessary for the up-regulation of EP4 gene transcription in response to Fn. There are three AP-2

binding sites in this region. By mutations of each of these sites, we found that only one site (\sim 1000 bp) was involved in the Fn-induced EP4 promoter activity. To our knowledge, a role for the AP-2 site in regulation of EP4 expression has never been reported. The transcription factor AP-2 regulates genes involved in a spectrum of important biological functions. Data obtained from different experimental models *in vitro* and *in vivo* indicate that AP-2 proteins function as important regulators of c-Myc targets in cell cycle progression and apoptosis (48). AP-2 α overexpression leads to anchorage-independent growth and malignant transformation *in vitro*. It has therefore been suggested that AP-2 is involved in malignant transformation of breast cancer cells. Immunohistochemical analysis of AP-2 α - and AP-2 γ -specific promoters confirmed up-regulation of AP-2 proteins in human breast cancer (50). Other data indicate that AP-2 α might be involved in cell proliferation by regulating cell cycle progression and apoptosis with retardation (51). There are a few studies that explore the transcriptional regulation of EP4 and transcription factor interactions in its promoter region. Studies show that a GC-rich/Sp1 binding site located within the first 80 bases of the transcription start site in the EP4 promoter region is important in transcription initiation of the EP4 gene (24), and several negative, positive and lipopolysaccharide/serum-responsive regions are

located at different areas in the mouse EP4 promoter (16). We confirmed that AP-2 sites were involved in Fn-induced EP4 gene expression using mutated EP4 constructs.

Fn increased nuclear AP-2 α protein levels, and inhibition of ERK signals prevented Fn-induced AP-2 expression. The connection between the AP-2 and ERK signal has been reported in other studies. For example, the ability of estradiol to increase AP-2 protein expression and AP-2 DNA binding activity was reversed by PD98059 (52). ERK activation is necessary for induction of the binding activities of AP-2 in T cells (53). Also, increased ERK signal has been reported in the absence of AP-2 α in mouse epidermis (54). In addition, we found that blockade of AP-2 by AP-2 antisense approaches had no effect on ERK activation.

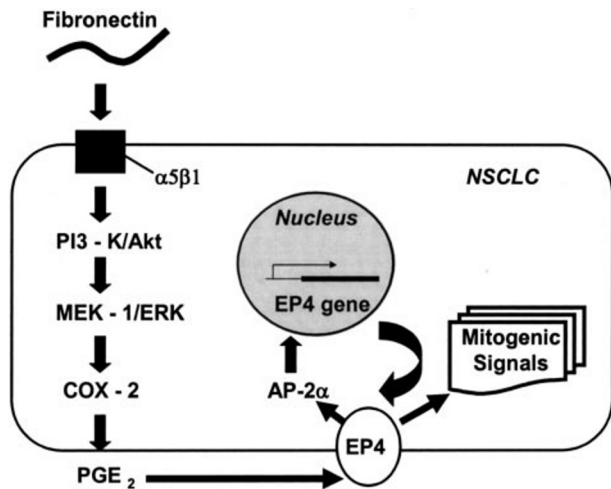


FIGURE 9. Schematic representation of signal pathways triggered in NSCLC in response to Fn. By binding to its $\alpha 5 \beta 1$ integrin receptor, Fn stimulates PI3K/Akt, followed by the induction of MEK-1/ERK and COX-2 signal pathways. These events stimulate the production of PGE_2 and expression of the PGE_2 subtype EP4 receptor gene through activation of the transcription factor AP-2 α . These two events, induction of PGE_2 and EP4, serve to amplify the mitogenic effects of Fn on lung carcinoma cells.

ity in the presence or absence of PGE_2 (not shown), suggesting that ERK was not a downstream signal of AP-2. Furthermore, we showed that inhibition of PI3K and blockade of $\alpha 5 \beta 1$ integrin signals abrogated Fn-induced AP-2 α protein expression, indicating that multiple signals were involved in this process. Showing that AP-2 antisense oligonucleotides abrogated Fn-induced cell growth, our data suggest that AP-2 is a critical component in this pathway. Thus, Fn is at least partly responsible for the stimulation of proliferation.

We previously demonstrated that siRNA abrogated Fn-induced expression and stimulation of AP-2 α protein expression, indicating that COX-2 signaling was involved in this process. We showed that siRNA abrogated Fn-induced AP-2 α protein expression in NSCLC cells (15). Since COX-2 is a key enzyme in PGE_2 production, the effect of Fn on AP-2 α protein expression is through stimulation of PGE_2 production. The literature contains data that link PGE_2 production to AP-2 signaling (55, 56). PGE_2 not only stimulates AP-2 α expression but also enhanced Fn-induced AP-2 α protein expression and cell proliferation, suggesting that stimulation of PGE_2 by Fn might account, at least partially, for the production of AP-2 α that resulted in EP4 expression and, subsequently, increased cell proliferation. PGE_2 has been shown to enhance the binding activity of AP-2 in different gene promoters in several studies (57, 58). $dmPGE_2$ up-regulated expression of the receptor subtype EP4, suggesting that PGE_2 might combine with EP4 receptors in autocrine or paracrine modes. Thus, blockade of the EP4 gene by siRNA approaches diminished the stimulatory effect of PGE_2 on lung carcinoma cell growth (this study). The dose of PGE_2 used in this study was based on our previous results showing Fn-induced PGE_2 production in NSCLC cells *in vitro* (4). Similar or even higher doses of exogenous PGE_2 have been shown to suppress both Th1- and Th2-polarized antigen-specific human T-cell responses and to reduce radiation-induced apoptosis in human colon cancer cells without toxicity (59, 60). Previously, we reported

that PGE_2 enhanced NSCLC cell growth in the presence of the EP2 agonist, Butaprost, suggesting that EP2 also plays a role in mediating NSCLC cell growth (17). Whether EP2 is directly involved in the effect of PGE_2 on NSCLC cell proliferation and how this pathway is affected by Fn needs to be determined.

In summary, our studies show that Fn stimulates human lung carcinoma cell proliferation through the PGE_2 receptor subtype EP4. This effect is enhanced by Fn-induced EP4 expression. Control of EP4 gene expression by Fn is dependent on $\alpha 5 \beta 1$ integrin-mediated signals that include activation of PI3K/Akt, ERK, and COX-2. These signals stimulate PGE_2 production, which induces the transcription factor AP-2 α and stimulates AP-2 α interactions with critical DNA regions within the EP4 gene promoter (−1555 to −992 bp). Thus, Fn, by increased PGE_2 production and EP4 gene expression, induces mitogenic signals in NSCLC cells (Fig. 9). This study reveals a novel molecular mechanism for Fn regulation of human lung carcinoma cell growth and provides further evidence for the role of E prostanoïd receptors in lung carcinoma biology.

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