The Fibroblast Growth Factor Receptor-1 Is Necessary for the Induction of Neurite Outgrowth in PC12 Cells by aFGF

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The PC12 subclone, fnr-PC12 cells, is defective in neurite outgrowth in response to acidic fibroblast growth factor (aFGF); however, its response to nerve growth factor (NGF) is normal. Examination of the expression of FGF receptors (FGFRs) revealed that although PC12 cells express FGFR-1, -3, and -4, fnr-PC12 cells have a reduced level of expression of FGFR-1 but not FGFR-3 and -4. Transfection of FGFR-1 into fnr-PC12 cells efficiently restored aFGF-induced neurite outgrowth, whereas transfection of FGFR-3 was much less efficient. Transfection of a chimeric receptor consisting of the extracellular domain of FGFR-3 fused to the transmembrane and intracellular domain of FGFR-1, termed FR31b, efficiently restored aFGF-induced neurite outgrowth. This demonstrates that the difference between these two receptors in their ability to induce neurite outgrowth is attributable to differences in the signaling capacity of their cytoplasmic domains. Activation of the chimeric receptor by aFGF induced a stronger and more persistent increase in the tyrosine phosphorylation of cellular proteins than did activation of FGFR-3 alone. In particular, the activation of MAP kinase by FR31b was more persistent than when activated by FGFR-3. This difference in signaling potential of FGFR-1 and -3 in fnr-PC12 cells may account for the difference in the potential for induction of neurite outgrowth. These results demonstrate that FGF-induced neurite outgrowth in PC12 cells occurs mainly via FGFR-1 and not via the other FGFRs expressed in these cells.

Key words: aFGF; FGFR-1; FGFR-3; neurite outgrowth; PC12 cells; fnr-PC12 cells; tyrosine phosphorylation; MAP kinase

The fibroblast growth factor (FGF) family of growth factors has diverse effects during embryonic development (for reviews, see Basilio and Moscatelli, 1992; Mason, 1994). In the nervous system, FGFs exhibit neurotrophic properties similar to those of the neurotrophins (for reviews, see Baird, 1994; Eckenstein, 1994), but their actions on nerve cells have been difficult to ascertain because they also affect glial cells (Engele and Bohn, 1991). Injection of FGF into the CNS of lesioned animals causes neurotrophic effects and stimulates the regeneration of lesioned axons (Anderson et al., 1988; Lipton et al., 1988; Cordeiro et al., 1989; Sasaki et al., 1992; MacMillan et al., 1993a,b; Nakata et al., 1993). In primary cell culture, FGFs mediate the survival, proliferation, and differentiation of both central and peripheral neurons and promote expression of neuronal characteristics (Morrison et al., 1986; Walicke et al., 1986; Schubert et al., 1987; Unsicker et al., 1987; Eckenstein et al., 1990; Sendtner et al., 1991; Ghosh and Greenberg, 1995; Vicario-Abejón et al., 1995). FGFs also stimulate the proliferation of peripheral sympathoadrenal precursors and induce responsiveness to nerve growth factor (NGF) (Birren and Anderson, 1990).

FGF actions are mediated by the binding and activation of FGF receptor (FGFR) tyrosine kinases. FGFRs are a gene family of four members (for review, see Jaye et al., 1992; Partanen et al., 1992; Johnson and Williams, 1993) termed FGFR-1 (flg), FGFR-2 (bek), FGFR-3, and FGFR-4. Expression patterns of FGFRs in the nervous system suggest that FGFs may have differential effects on distinct neuronal cell types throughout development. FGFR-1, -2, and -3 are expressed differentially in subsets of neurons within both the peripheral nervous system (PNS) and the CNS as well as in glial cells (Heuer et al., 1990; Wanaka et al., 1990; Thompson et al., 1991; Asai et al., 1993; Peters et al., 1993; Yazaki et al., 1994). During the development of the chick nervous system, FGFR-1 is transiently expressed on different neuronal cell types but in a manner reciprocal with that of the NGF receptor (Heuer et al., 1990). In some neuronal cell types, co-expression of FGFRs and neurotrophin receptors is observed (Claude et al., 1988; Stemple et al., 1988). Thus it has been suggested that the FGFs act coordinately with the neurotrophins to guide neuronal development. Localization of the four different FGFRs on neuronal cell types is complex (Heuer et al., 1990; Wanaka et al., 1990; Thompson et al., 1991; Asai et al., 1993; Peters et al., 1993; Yazaki et al., 1994), making it difficult to elucidate the physiological role for a specific FGF. FGF activity is complicated further by selective binding to different FGFRs (Jaye et al., 1992; Ornitz and Leder, 1992; Partanen et al., 1992; Johnson and Williams, 1993; Chellaiah et al., 1994). Consequently, the roles of each FGFR in the nervous system are still not known.

The PC12 cell line has provided a useful model for studying the actions of neurotrophins (Greene and Tischler, 1982). These cells respond to FGFs and NGF by the elaboration of a sympathetic neuron-like phenotype (Togari et al., 1985; Rydel and Greene,
1987). The mechanisms of NGF and FGF action on PC12 cells seem to use similar signal transduction pathways for neurite outgrowth, beginning with the activation of a receptor tyrosine kinase and using a signaling cascade involving src, ras, raf, and the MAP kinases (Kremer et al., 1991; Thomas et al., 1992; D’Arcangelo and Hagegoua, 1993; Wood et al., 1993). The FGF Rs that mediate FGF actions on PC12 cells, however, have not been well characterized. Here we analyze FGF expression and function in PC12 cells. Using a PC12 subline that is nonresponsive to FGF, we demonstrate that although PC12 cells normally express FGFR-1, -3, and -4, the FGF-nonresponsive cells no longer express high levels of FGFR-1. The levels of FGFR-3 and -4 are similar in both cell types. By transfection of FGFR-1 or -3 into the FGF-nonresponsive subline, we determined that only activation of FGFR-1 leads to high-efficiency neurite outgrowth. These data indicate that FGF-induced neurite outgrowth in PC12 cells occurs primarily via FGFR-1 and not via the other FGF Rs expressed in PC12 cells. These studies provide a first step toward elucidating the actions of FGF through specific FGF Rs expressed in neurons.

MATERIALS AND METHODS

Cell culture and growth factors. fnr-PC12 (also named GR-5) cells are a subclone of PC12 cells (Greene and Tischler, 1976) isolated and kindly provided by Rae Nishi (Oregon Health Sciences University, Portland, OR). PC12, fnr-PC12, and fnr-PC12-derived transfected lines were grown on tissue culture dishes in DMEM (Life Technologies, Grand Island, NY) supplemented with 10% horse serum (JRH Biosciences, Lenexa, KS), 5% fetal bovine serum (JRH Biosciences), and 1% penicillin/streptomycin (Life Technologies) in an atmosphere of 10% CO2; 95% air; 5% O2. This medium was supplemented with 10 ng/ml NGF (provided by Rae Nishi (Oregon Health Sciences University, Portland, CA) and centrifuged for 15 min at 4°C in an Eppendorf centrifuge. Cell pellets were washed in PBS and then resuspended in PBS and collected by centrifugation in an Eppendorf centrifuge. Cell pellets were washed in PBS and then resuspended in PBS and collected by centrifugation in an Eppendorf centrifuge. Cell pellets were washed in PBS and then resuspended in PBS and collected by centrifugation in an Eppendorf centrifuge.

Cell line generation. Using a pc12 subline that is nonresponsive to FGF, we determined that only activation of FGFR-1 leads to high-efficiency neurite outgrowth. These data indicate that FGF-induced neurite outgrowth in PC12 cells occurs primarily via FGFR-1 and not via the other FGF Rs expressed in PC12 cells. These studies provide a first step toward elucidating the actions of FGF through specific FGF Rs expressed in neurons.

Plasmids and transfection. The human FGFR-1 expression vector pFlgFL24 was kindly provided by Michael Jaye (Rhöne-Poulenc Rorer Central Research). This vector consists of full-length human FGFR-1 cDNA (Dionne et al., 1990) in the eukaryotic expression vector pM300 (Jaye et al., 1988). The human FGFR-3 expression vector pLhK3-5 consists of full-length human FGFR-3 cDNA in the retroviral vector pLNCX (Miller and Rosman, 1989). Mo/FRIII/ires expression vector consists of cDNA encoding a chimeric receptor with the extracellular domain of mouse FGFR-3 IIIb (Chelliah et al., 1994) and the transmembrane domain and intracellular domain of mouse FGFR-1 (Yayon et al., 1991). Mo/FRMIIIb/ires expression vector consists of cDNA encoding the full-length mouse FGFR-3 IIIb. In these latter two expression vectors, expression of FGF Rs were driven by Moloney murine leukemia virus LTR, and expression of neomycin-resistant gene was driven by encephalomyocarditis virus internal ribosomal entry site (EMCV IRES) (Ghattas et al., 1991).

Transfection of the expression vectors to fnr-PC12 cells was performed as described previously (D’Arcangelo and Hagegoua, 1993). The transfection was carried out by electroporation, using a Gene Pulser (Bio-Rad, Richmond, CA). Approximately 4 × 106 cells were electroporated in 0.4 ml of DMEM containing either 20 μg of pFlgFL24 plus 2 μg of SV2hs (Hartman and Mulligan, 1988) or 20 μg of the other expression vectors described above, using settings of 250 V and 500 μF. Two days after transfection, cells were incubated in histidine-free RPMI 1640 medium (Life Technologies) containing 1 mM histidine for pFlgFL24-transfected cells or 800 μg of geneticin (Life Technologies) for the other transfected cells. After ~3 weeks, histidinol-resistant or neomycin-resistant clones were isolated and then screened for FGF-induced neurite outgrowth and the level of expression of the transfected receptors.

Covariant cross-linking of [125I]aFGF to intact cells. Covariant cross-linking of [125I]aFGF to intact cells was performed as described previously (Dionne et al., 1990). Cells were grown on 60-mm diameter tissue culture dishes coated with human fibronectin (Collaborative Research, Bedford, MA). At 80% confluency, cells were washed twice with binding buffer (DMEM containing 0.2% BSA, 5 μM heparin, and 20 mM HEPES, pH 7.5) and incubated for 90 min on ice with binding buffer containing 25 ng/ml [125I]aFGF. After cells were washed once with binding buffer and once with PBS (137 mM NaCl, 2.7 mM KCl, 4.3 mM Na2HPO4, 7 mM H2O, 1.4 mM KH2PO4, pH 7.3), cells were incubated further with PBS containing 0.3 mM disuccinimidyl suberate (DSS) (Pierce, Rockford, IL) as cross-linker (prepared as 30 mM stock solution in DMSO). Cells were then washed once with 10 mM HEPES, pH 7.5, 200 mM glycine, 2 mM EDTA, once with PBS, and then scraped in PBS and collected in an Eppendorf centrifuge. Cell pellets were lysed in 50 μl of lysis buffer (20 mM HEPES, pH 7.5, 150 mM NaCl, 1% Triton X-100, 10% glycerol, 1.5 mM MgCl2, 1 mM EDTA, 1 μg/ml aprotinin, 1 μg/ml leupeptin, and 1 mM PMSF) incubated for 15 min on ice, and centrifuged for 15 min at 4°C in an Eppendorf centrifuge. The protein concentrations in the clarified lysates were determined by using BCA protein assay reagent (Pierce). Aliquots of the lysates (500 μg) of each sample were mixed with Laemmli sample buffer (Laemmli, 1970), boiled for 4 min, and fractionated on a SDS-7.5% polyacrylamide gel. The gels were stained with Coomassie brilliant blue to verify that all samples contained equal amounts of protein. Autoradiograms of the dried gels were made on Fuji medical x-ray film RX.

Anti-FGFR antisera. The anti-FGFR-1 antisera flg-2B were raised against the peptide SSQEDSVGHEPLPEEPEPTV (human FGFR-1 C-terminal penultimate sequence). The two different anti-FGFR-2 anti-sera Bek-78B and Bek-1A were raised against the peptides RPSFSEVSQQPPPPP (human FGFR-2 N-terminal peptide) and YEPFLCQYPYPHNSVKT (human FGFR-2 C-terminal sequence), respectively. These anti-FGFR-1 and anti-FGFR-2 antisera was generously provided by Michael Jaye (Rhöne-Poulenc Rorer Central Research). The anti-FGFR-3 monoclonal antibody 8.34 was raised against a bacterially expressed polyepitopes encompassing the amino acids 94–255 of the extracellular domain of human FGFR-3. The anti-FGFR-4 polyclonal antisera was raised against the peptide CGGGFGGGSGQT, the C-terminal 11 amino acids of murine FGFR-4 tagged with CGG for conjugation to carrier. These anti-FGFR-4 antisera was generously provided by Jen-Kuei Wang and Mitchell Goldfarb (Regeneron Pharmaceuticals, Tarrytown, NY).

Immunoprecipitation and Western blot. For immunoprecipitation, to detect aFGF-stimulated autophosphorylation of the transfected FGF Rs, cells were treated with aFGF for 5 min, lysed in lysis buffer (20 mM Tris, pH 7.6, 150 mM NaCl, 50 mM NaF, 1 mM Na2VO4, 5 mM benzamidine, 1 mM EDTA, 10 μg/ml leupeptin, 10 μg/ml aprotinin, 1 mM phenylmethylsulfonyl fluoride, and 1% Nonidet P-40). The lysates were clarified by the addition of Protein A-Sepharose-4B (Zymed, South San Francisco, CA) and centrifugation. The protein concentrations in the supernatant were determined as described previously (Dionne et al., 1990). Aliquots (10 μg of each sample were incubated with anti-phosphotyrosine polyclonal antisera SB56 for 1 hr on ice. Protein A-Sepharose-4B was added for an additional 0.5 hr. Pellets were collected by centrifugation, washed, boiled in sample buffer, and subjected to SDS-PAGE. Western blotting was performed, and membranes were probed with anti-FGFR-3 antibody.

For Western blotting, (1) to compare the levels of FGF Rs in PC12, fnr-PC12, and fnr-PC12-derived transfected lines, the cells were lysed in lysis buffer and clarified, and the protein concentrations in the lysates were determined. Aliquots of the lysates (500 μg and 75 μg for the results shown in Figs. 1 and 6B, respectively) of each sample were mixed with sample buffer, boiled, and fractionated on an SDS-7.5% polyacrylamide gel. Protein was transferred to nitrocellulose membranes (Schleicher & Schuell, Keene, NH) by electroblotting. Non-specific binding sites were blocked with blocking buffer (3% BSA in PBS, 1 mM EDTA, 1 mM sodium vanadate) for 1 hr at room temperature. Membranes were then incubated with the appropriate anti-FGFR antisera diluted in blocking buffer for 1 hr at room temperature. Membranes were washed with PBST (PBS, 0.1% Tween 20) and subsequently incubated with anti-mouse Ig- or anti-rabbit-5-ε-linked swine anti-rabbit IgG (Afollow et al., Arlington Heights, IL) at a dilution of 1:2500 in PBS plus 5% milk, and reactivity was determined by the ECL chemiluminescence reaction (Amersham). (2) To detect aFGF-stimulated tyrosine phosphorylation of cellular proteins, cells were treated with 50 ng/ml of aFGF at 37°C for the time indicated in the figure. Aliquots of the lysates (75 μg) of each sample were fractionated on an SDS-7.5% polyacrylamide gel. Western blotting was performed, and membranes were probed with a solution consisting of
the anti-phosphotyrosine monoclonal antibody 4G10 (generously provided by Brian J. Druker, Oregon Health Sciences University) at a concentration of 1 μg/ml in blocking buffer. (3) To detect aFGF-stimulated activation of MAP kinase, cells were treated with 50 ng/ml aFGF at 37°C for the time indicated in the text. Aliquots of the lysates (75 μg) of each sample were fractionated on an SDS-12.5% polyacrylamide gel (Leevers and Marshall, 1993). Western blotting was performed, and membranes were probed with a solution consisting of an anti-ERK2 monoclonal antibody (generously provided by Roger Davis, University of Massachusetts, Worcester, MA) in blocking buffer. Activation of MAP kinase was determined by mobility shift. All Western blots were performed in antibody excess.

RESULTS
Neurite outgrowth of fnr-PC12 cells
fnr-PC12 cells are a subclone of PC12 cells that show a greatly reduced response in neurite outgrowth after FGF treatment when compared with the parental PC12 cells. For example, when these two cell types were treated with aFGF for 5 d, PC12 cells changed morphology (data not shown) and extended neurites (Fig. 1D), whereas fnr-PC12 cells remained morphologically unchanged (Fig. 1A,B). In contrast, NGF stimulated neurite outgrowth of both PC12 cells (data not shown) and fnr-PC12 cells (Fig. 1C). When the percentages of neurite-bearing cells were scored, both PC12 cells and fnr-PC12 cells showed a similar level of response to NGF (Fig. 1D). These results indicate that fnr-PC12 cells are defective in some aspect of the FGF-specific signaling pathway.

Chemical cross-linking of [125I]aFGF to fnr-PC12 cells
To examine whether the FGF-nonresponsiveness of the fnr-PC12 cells was attributable to a lack of cell surface receptors, we determined the level of aFGF binding sites on PC12 and fnr-PC12 cells by chemical cross-linking of [125I]aFGF. In PC12 cells, a major band of 180 kDa was cross-linked to [125I]aFGF. This band, however, was barely seen in fnr-PC12 cells (Fig. 2A). With a longer exposure of the gel, two bands of 180 and 160 kDa were found to be cross-linked to [125I]aFGF in both cells. The relative intensity of these two bands was different in these two cells. In PC12 cells, the band of 180 kDa was much stronger than the band of 160 kDa. In fnr-PC12 cells, however, the band of 160 kDa was slightly stronger than the band of 180 kDa (Fig. 2B). These results indicate that fnr-PC12 cells have endogenous FGF receptors, but at a significantly reduced level in comparison with PC12 cells.

Figure 1. aFGF- and NGF-stimulated neurite outgrowth. PC12 cells and fnr-PC12 cells were untreated or treated with 50 ng/ml aFGF or 50 ng/ml NGF. The photographs show the morphology of fnr-PC12 cells, (A) untreated, (B) treated with aFGF, and (C) treated with NGF for 5 d (D). The percentage of neurite-bearing cells was counted from 2 to 5 d after treatment with aFGF or NGF.

Figure 2. Cross-linking of [125I]aFGF to PC12 and fnr-PC12 cells. PC12 cells or fnr-PC12 cells were incubated with 25 ng/ml [125I]aFGF for 90 min, followed by cross-linking with DSS. The lysates were fractionated by SDS-PAGE, and the gel was exposed for autoradiography. A. Four day exposure of the gel; B, 33 d exposure of the gel.
Analysis of FGF receptors expressed in PC12 cells and fnr-PC12 cells

To determine which specific FGF receptor was no longer expressed at high level in fnr-PC12 cells, the expression of the four known FGF receptors was analyzed by Western blot probing with receptor-specific antibodies (Fig. 3). The endogenous level of FGFR-1 was found to be much lower in fnr-PC12 cells when compared with PC12 cells, whereas the endogenous level of FGFR-3 or -4 essentially remained the same. By using two different anti-FGFR-2 antibodies for this analysis, we failed to detect the expression of FGFR-2 in these cells (data not shown). On the basis of the known molecular weights of FGFR-1, -3 and -4, together with these expression studies, this would indicate that the 180 kDa receptor detected by ligand binding in Figure 2B is FGFR-1 and the 160 kDa receptor is FGFR-3. No ligand-binding activity was detected, even on the longer exposures, that would correspond to FGFR-4. Furthermore, these results also suggest that the endogenous FGFR-3 and -4 are unable to induce neurite outgrowth and that the reduced expression of FGFR-1 may be responsible for the inability of fnr-PC12 cells to undergo efficient aFGF-induced neurite outgrowth.

Transfection of fnr-PC12 cells with FGFR-1 or -3 restores aFGF-stimulated neurite outgrowth of fnr-PC12 cells with different efficiency

Western blot analysis of endogenous FGF receptors in PC12 cells and fnr-PC12 cells revealed that fnr-PC12 cells expressed greatly reduced levels of FGFR-1 but essentially the same levels of FGFR-3 and -4 (Fig. 3). We hypothesized that if this reduced level of FGFR-1 expression caused the lack of neurite outgrowth, then transfection of FGFR-1 into these cells should restore aFGF-stimulated neurite outgrowth. Because the functional role of different FGFRs in PC12 cells was not known and at least three different FGFRs were coexpressed in these cells, it would be possible that transfection of any of the three receptors into fnr-PC12 cells and selection of high-level expressors would also restore FGF-induced neurite outgrowth in these cells. To address these hypotheses, human FGFR-1 or -3 cDNAs was transfected into fnr-PC12 cells. Cell lines stably expressing FGFR-1 or -3 were selected and screened by aFGF-stimulated neurite outgrowth and cross-linking of the cells with \[^{125}\text{I}]aFGF. Among the 20 FGFR-1-transfected clones and 20 FGFR-3-transfected clones tested, FGFR-1-transfected clones elaborated neurites more efficiently than FGFR-3-transfected clones when treated with aFGF. This result indicates that FGFR-1 and -3 may have different potentials to restore aFGF-stimulated neurite outgrowth in fnr-PC12 cells. To compare quantitatively the functional difference between these two receptors, the percentage of neurite-bearing cells was scored for clones expressing approximately equal levels of transfected FGFR-1 or -3. Analysis of one clone of FGFR-1, hR1-3B2, and two clones of FGFR-3 expressing receptors, hR3–1D1 and hR3–1D2, demonstrates the different efficiency of aFGF-stimulated neurite outgrowth by these two receptors (Fig. 4). As determined by cross-linking of \[^{125}\text{I}]aFGF to these cells, all of these three transfected clones showed increased levels of cell surface aFGF binding sites when compared with the parental line fnr-PC12 cells (Fig. 4A). Among these clones, hR3–1D2 and hR1–3B2 expressed equally high levels of transfected receptors (Fig. 4A), yet the efficiency of aFGF-induced neurite outgrowth was dramatically different (Fig. 4B). After the 5 d treatment with aFGF, the percentage of neurite-bearing cells in hR1–3B2 cells was significantly higher than that in PC12 cells, whereas the percentage of neurite-bearing cells in hR3–1D2 was significantly lower than that in PC12 cells or hR1–3B2 cells. Therefore, expression of human FGFR-1 to a high level in fnr-PC12 cells efficiently restored aFGF-induced neurite outgrowth, and expression of human FGFR-3 in these cells did not. These results indicate that FGFR-1 and -3 may have different potentials to induce neurite outgrowth in PC12 cells.

To rule out the possibility that the difference between these two receptors reflected differences in their abilities to bind to aFGF, cDNAs encoding either the full-length mouse FGFR-3 or a chimeric receptor, FR31b, with the extracellular domain of mouse FGFR-3 fused to the transmembrane domain and intracellular domain of FGFR-1 (Fig. 5) were transfected into fnr-PC12 cells. Fnr-PC12 cells transfected with either the mouse FGFR-3 or the chimeric receptor were selected for neo-resistance and screened for aFGF-stimulated neurite outgrowth and by Western blot analysis for receptor expression. At least 95 clones of each receptor-transfected cell were tested for their response to aFGF. Among the FR31b-transfected clones, 19% of the clones showed outgrowth of long and dense neurites after only an overnight incubation of the cells with aFGF, whereas no significant level of neurites was found in any of the mouse FGFR-3-transfected clones (data not shown). The Western blot analysis indicated that the FR31b clones that elaborated neurites when aFGF was added
were those that expressed the FR31b receptor. In contrast, although none of the FGFR-3 transfected clones efficiently elaborated neurites, several clones were identified that expressed high levels of FGFR-3. This indicates that the transmembrane domain and intracellular domain of FGFR-1 is sufficient to confer the high efficiency of induction of neurite outgrowth to fnr-PC12 cells. To quantitatively compare the difference between FR31b and FGFR-3, we identified several clones expressing equivalent levels of transfected FR31b or FGFR-3 and compared them for the percentage of neurite-bearing cells when aFGF was added. Representative data from two such clones is shown in Figure 6. Comparison of an FR31b-transfected clone, FR31b-1D1, and an FGFR-3-transfected clone, FR3IIIb-1C4, clearly demonstrates the difference in the efficiency of aFGF-stimulated neurite outgrowth by these two receptors (Fig. 6). As determined by Western blot analysis using an antibody against the extracellular domain, the level of expression of mouse FGFR-3 in FR3IIIb-1C4 was slightly higher than the level of FR31b in FR31b-1D1 (Fig. 6A). The efficiency of aFGF-stimulated neurite outgrowth of FR3IIIb-1C4, however, was lower than the efficiency of FR31b-1D1 (Fig. 6B). When the cells were incubated with aFGF for more than 2 d, FR3IIIb-1C4 cells had significant neurite outgrowth, yet the neurites were less dense and shorter than the neurites of FR31b-1D1 (data not shown). The higher efficiency of neurite outgrowth in FR3IIIb-1C4 (Fig. 6B) compared with the poor efficiency in hR3–1D2 (Fig. 4B) could be attributed to the higher level of receptor expression in FR3IIIb-1C4 cells or species-specific difference. During the course of this analysis, we have analyzed >100 individual clones expressing various FGFR constructs. We have noticed that FGFR-1, or FR31b clones, expressing high levels of receptor respond very quickly to aFGF addition, whereas those expressing intermediate levels respond more slowly, and the clones with low level expression respond the slowest to aFGF addition. This observation indicates that the expression level of the receptor is an important factor in determining the responsiveness of the cells to aFGF. Cells expressing very high levels of FGFR-3, however, still respond only poorly to aFGF addition. In summary, the results with the chimeric receptor FR31b further support the notion that FGFR-1 has a significantly higher potential than FGFR-3 to induce neurite outgrowth.

Comparison of signaling potentials of the FR31b chimera and FGFR-3 in fnr-PC12 cells

It is likely that differences in the signaling potential of FGFR-1 and -3 may be responsible for the differences in the efficiency of induction of neurite outgrowth. To dissect the molecular signaling events required for efficient induction of neurite outgrowth, fnr-PC12, FR31b-1D1, and FR3IIIb-1C4 cells were treated with aFGF for different time periods, and the aFGF-stimulated increase in tyrosine phosphorylation was analyzed. After stimulation with aFGF for 2–60 min, both FR31b-1D1 and FR3IIIb-1C4 cells showed an increase in tyrosine phosphorylation of a similar array of proteins (Fig. 7A). The most prominent increase in
tyrosine phosphorylation seen in FR31b-1D1 cells was in a 90 kDa protein (Fig. 7A). The signal was seen within 2 min and persisted for 60 min, whereas in FR3IIIb-1C4 cells, the signal was not as strong as seen in FR31b-1D1 cells and was already maximum at 2 min and proceeded to decrease. Interestingly, the decline seemed to be maximal after 30 min, and then there was a slight increase in the phosphorylation of several proteins at the 60 min time point. This is most easily seen for the two proteins of apparent molecular weights 44,000 and 42,000, Figure 7A. The reason for this transient decline is unknown, but it is a reproducible phenomenon, having been seen in six separate experiments. In addition to the 90 kDa protein, several other proteins seemed to be phosphorylated to higher levels after FR31b activation than they were after FGFR-3 activation. Results similar to those obtained with FR31b were seen when FGFR-1 was used (data not shown).

It is also interesting to note that certain proteins were phosphorylated more significantly after FGFR-3 activation than after FR31b activation. For example, see those at 42 and 35 kDa (Fig. 7A). This indicates that it is not just that FGFR-3 activation is somehow weaker than FR31b activation; Figure 7B shows that both receptors were autophosphorylated in response to aFGF stimulation. Higher basal levels of FGFR-3 autophosphorylation without aFGF treatment of the cells were observed reproducibly. The reason for this is unknown. These data indicate that in general, FR31b and FGFR-1 may phosphorylate and activate the key component for neurite outgrowth more efficiently than FGFR-3.

Kinetics of MAP kinase phosphorylation by FR31b and FGFR-3

Previous studies have indicated that activation of MAP kinase is involved in neurite outgrowth. Furthermore, for efficient neurite outgrowth, the activation of MAP kinase has to be persistent. Therefore we examined the abilities of both FR31b and FGFR-3 to activate MAP kinase. Activation of MAP kinase has been shown to result in a reduced mobility of the MAP kinase protein in SDS-PAGE. We examined aFGF-stimulated activation of MAP kinase by FR31b or FGFR-3 using this assay (Fig. 8). The parental line fnr-PC12 still contains some endogenous FGF receptors, and addition of aFGF to these cells showed a small but detectable activation of MAP kinase that was maximum at 5 min and transient in nature. Overexpression of FR31b or FGFR-3 greatly enhanced MAP kinase activation, but the activation is more prolonged in cells expressing FR31b; again similar results were obtained with FGFR-1 (data not shown). At least 50% of the activated form of the enzyme was still present after 60 min of treatment, whereas the signal by FGFR-3 was much more transient compared to the signal by FR31b. Within the first 5 min, the levels of activation by FR31b and FGFR-3 were similar; however, by 10 min the MAP kinase was returning to its inactive form in FGFR-3-expressing cells, whereas a significant portion of the enzyme was still active in FR31b-expressing cells (Fig. 8). On the basis of previous studies, this difference in the ability to induce prolonged activation of MAP kinase may be sufficient to explain the differing abilities of these two FGF receptors to induce neurite outgrowth.

DISCUSSION

Analysis of an FGF-nonresponsive PC12 subclone, named fnr-PC12, has allowed us to determine the role of the different FGFRs in FGF-stimulated neurite outgrowth. By cross-linking with [125I]aFGF, these cells expressed a greatly reduced level of FGFRs when compared with the level in PC12 cells. By Western blot analysis, the major difference of endogenous FGFRs was the greatly reduced level of FGFR-1 in fnr-PC12 cells. Previous studies by overexpression of some other growth factor receptors, such as EGF receptor (Traverse et al., 1994), insulin receptor (Dikic et al., 1994), or PDGF receptor (Heasley and Johnson, 1992), or selection of PC12 variants expressing reduced levels of NGF receptor (Schlessinger and Bar-Sagi, 1994), suggest that the number of receptors affects the biological outcome: differentiation versus mitogenesis. Ligand activation of PC12 cells that highly overexpressed any of these receptor tyrosine kinases led to sustained activation of MAP kinase, cessation of cell division, and stimulated neurite outgrowth. In contrast, PC12 cells that expressed these receptors below a threshold level no longer stimulated neurite outgrowth. These experiments raised two possibilities for the defect in neurite outgrowth seen in fnr-PC12 cells. One possibility is that the number of endogenous FGFRs is not high enough to give rise to FGF-stimulated neurite outgrowth. If the different FGFRs in PC12 cells are functionally redundant, transfection of any FGF receptor should restore FGF-stimulated neurite outgrowth with the same efficiency. The other possibility is that fnr-PC12 cells lack the receptor that is mainly responsible for neurite outgrowth. Our results support this second possibility.
FGFR-1 was significantly more active than FGFR-3 in the induction of neurite outgrowth when both receptors are overexpressed at high level. This suggests that these two receptors are not functionally redundant for induction of neurite outgrowth in PC12 cells. Even though overexpression of FGFR-3 in fnr-PC12 cells may restore some aFGF-stimulated neurite outgrowth in these cells, this may not represent the physiological role of FGFR-3 in PC12 cells, because in the absence of FGFR-1 the physiological level of FGFR-3 and -4 found in fnr-PC12 cells cannot support aFGF-stimulated neurite outgrowth. Therefore, FGFR-1 seems to be the FGFR primarily responsible for induction of neurite outgrowth. Additional experiments will be necessary to explore the role of the other FGFRs in the differentiation of PC12 cells.

Recent reports suggest that activation of multiple signal transduction pathways and the duration of activation are essential for growth factor-stimulated neurite outgrowth in PC12 cells (Peng et al., 1995; Vaillancourt et al., 1995). In this study, we demonstrate that neurite outgrowth correlates with the differing ability of the different FGF receptors to activate MAP kinase. FGFR-1 activates MAP kinase in a more sustained manner than FGFR-3. This is consistent with previous observations for other growth factor receptors expressed in PC12 cells. In these studies, it was demonstrated that sustained activation of MAP kinase correlates with efficient induction of neurite outgrowth (Heasley and Johnson, 1992; Hempstead et al., 1992; Dikic et al., 1994; Traverse et al., 1994). In addition to MAP kinase, the higher phosphorylation of an array of cellular proteins that is induced by FGFR-1 (Fig. 6A) may also play a role in the more efficient induction of neurite outgrowth by this receptor. Among these proteins, the higher and more sustained phosphorylation of the 90 kDa protein (Fig. 6A) and Shc (H. Lin and M. Hayman, unpublished data) further suggests that FGFR-1 activates ras signaling pathway more efficiently than FGFR-3 (Klint et al., 1995). Activation of the proteins PLCγ, Src, and SNT has been implicated in the signaling pathway for neurite outgrowth in PC12 cells (Peng et al., 1995; Vaillancourt et al., 1995). Whether these two receptors differentially activate other signaling pathways involving these molecules remains to be determined. Interestingly, in other cell types both

Figure 7. Activation of FGFR-1 kinase in FR31b-transfected fnr-PC12 cells induces stronger and more persistent protein tyrosine phosphorylation than activation of FGFR-3 kinase in FR3IIIb-transfected fnr-PC12 cells. Fnr-PC12 cells, FR31b-1D1 cells, and FR3IIIb-1C4 cells were untreated or treated with aFGF for 2, 5, 10, 30, or 60 min. The cells were lysed and subjected to SDS-PAGE. A, The increase in tyrosine phosphorylation of cellular substrates was monitored with an anti-phosphotyrosine antibody, 4G10. Proteins phosphorylated on addition of aFGF are indicated by arrowheads. The position of the 90 kDa protein discussed in the text is indicated by double asterisks. B, IP: anti-phosphotyrosine; blot: anti-FGFR-3. Autophosphorylation of the receptor molecules was assayed by immunoprecipitation of the lysates with an anti-phosphotyrosine antibody, SB56, followed by Western blotting and probing with the anti-human FGFR-3 monoclonal antibody.

Figure 8. Kinetics of aFGF-induced activation of MAP kinase in fnr-PC12, FR31b-1D1, and FR3IIIb-1C4 cells. The cells were untreated or treated with aFGF for 2, 5, 10, 30, or 60 min. The lysates were subjected to SDS-PAGE followed by Western blotting using an anti-ERK2 antibody. The activation of MAP kinase was detected by mobility shift in the SDS-PAGE. MAPK, Inactive form of MAP kinase; MAPK*, activated form of MAP kinase.
FGFR-3 and -4 have weaker functional potentials when compared with FGFR-1 (Ornitz and Leder, 1992; Wang et al., 1994; Shaoul et al., 1995). The underlying mechanism for this weaker potential remains unclear.

FGFR-1, -2, and -3 are distributed differentially throughout the nervous system among various neuronal as well as glial cell types (Heuer et al., 1990; Wanaka et al., 1990; Thompson et al., 1991; Asai et al., 1993; Peters et al., 1993; Yazaki et al., 1994). These distributions suggest that the FGFRs have diverse functions and that each receptor may regulate unique phenotypic characteristics of neuronal and glial cells. Our data support these ideas in that two different FGF receptors have been shown to confer different functionalities in the same neuronal background. FGFR-1 is much more effective in regulating characteristics associated with neurite outgrowth than is FGFR-3. This may be consistent with the more prevalent distribution of FGFR-1 in neuronal cell types (Heuer et al., 1990; Wanaka et al., 1990; Thompson et al., 1991; Asai et al., 1993; Peters et al., 1993; Yazaki et al., 1994). Because FGFR-3 is found in some neuronal cell types, and occasionally in cells with overlapping FGFR-1 expression, it will be of interest to determine whether any neuronal phenotypes other than neurite outgrowth may be elicited through FGFR-3.

REFERENCES


