

Characterization of Insulin-Like Growth Factor-I and Its Receptor and Binding Proteins in Transected Nerves and Cultured Schwann Cells

Hsin-Lin Cheng, Ann Randolph, *Douglas Yee, †Patrick Delafontaine, Gihan Tennekoon, and Eva L. Feldman

University of Michigan Department of Neurology and Neuroscience Program, Ann Arbor, Michigan; *University of Texas Health Science Center, San Antonio, Texas; and †Emory University, Atlanta, Georgia, U.S.A.

Abstract: The insulin-like growth factors (IGFs) are trophic factors whose growth-promoting actions are mediated via the IGF-I receptor and modulated by six IGF binding proteins (IGFBPs). In this study, we observed increased transcripts of both IGF-I and IGF-I receptor after rat sciatic nerve transection. Schwann cells (SCs) were the main source of IGF-I and IGFBP-5 immunoreactivity until 7 days after nerve transection, when invading macrophages in the distal nerve stumps were strongly IGF-I positive. In vitro, IGF-I promoted SC mitogenesis. Northern analysis revealed that SCs expressed IGF-I receptor and IGFBP-5. IGF-I treatment increased the intensity of IGFBP-5 without affecting gene expression. Des(1-3)IGF-I, an IGF-I analogue with low affinity for IGFBP, had no such effect. Incubation of recombinant human IGFBP-5 with SC conditioned media revealed IGF-I protection of IGFBP-5 from proteolysis, implying the presence of an IGFBP-5 protease in SC conditioned media. Collectively, these data support the concept that, in response to nerve injury, invading macrophages produce IGF-I and SC express the IGF-I receptor, to facilitate regeneration. This regenerative process may be augmented further by the ability of SC to secrete IGFBPs, which in turn may increase local IGF-I bioavailability. **Key Words:** Insulin-like growth factor—Receptor—Binding protein—Schwann cells—Nerve transection. *J. Neurochem.* **66**, 525–536 (1996).

The insulin-like growth factors (IGF-I and IGF-II) are polypeptides with both growth-promoting and insulin-like metabolic activities (Krywicky and Yee, 1992). The mitogenic actions of both IGFs are mediated via the IGF-I receptor, a tyrosine kinase receptor whose activation results in both auto- and intracellular substrate protein phosphorylation (Chou et al., 1987). The actions of the IGFs are modulated by six structurally related proteins, the IGF binding proteins (IGFBPs) (Clemmons, 1991; Drop, 1992). The expression of IGFBPs varies by cell and tissue type

(Clemmons, 1991; Drop, 1992) and is regulated by hormones (Torrington et al., 1991; Unterman et al., 1991) and growth factors (Martin and Baxter, 1991; Barreca et al., 1992), including IGF-I (Clemmons, 1991; Conover, 1991; Martin et al., 1992). The IGFBPs transport IGFs to target tissues (Binoux et al., 1991) and alter IGF binding to IGF receptors (McCusker et al., 1991). Secreted IGFBPs can also bind to extracellular surfaces and target IGFs to mediate growth in specific cell types. For example, in fibroblasts, IGFBP-5 associated with the extracellular matrix (ECM) augments IGF-I action, implying IGFBP-5 increases local access of IGF-I to the IGF-I receptor (Jones et al., 1993).

In the CNS, IGF, IGF receptor, and IGFBP expression is high during the fetal period (Beck et al., 1988; Clemmons, 1989; Daughaday and Rotwein, 1989; Sara and Hall, 1990; Liu et al., 1991), suggesting a role for these factors in normal development. During rat embryogenesis, IGF-I mRNA is detected in regions of active nerve sprouting, spinal ganglia, and facial target regions of the trigeminal nerve (Bondy et al., 1990). In humans, IGF-I is present in fetal and adult brain and in cerebrospinal fluid (Backstrom et al., 1984; Haselbacher et al., 1985; Han et al., 1988). In vitro, developing oligodendrocyte progenitor cells require IGF-I to progress to adult, myelin-forming oligodendrocytes (McMorris and Dubois-Dalq, 1988; Mozell and McMorris, 1988). IGFs also enhance growth in a

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Address correspondence and reprint requests to Dr. E. L. Feldman at University of Michigan, Department of Neurology, 4414 Kresge III, Ann Arbor, MI 48109, U.S.A.

Abbreviations used: DMEM, Dulbecco's modified Eagle's medium; ECM, extracellular matrix; FBS, fetal bovine serum; IGF, insulin-like growth factor; IGFBP, insulin-like growth factor binding protein; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; NGF, nerve growth factor; SC, Schwann cell; [³H]-TdR, [³H]-thymidine.

variety of neuronal types, including sympathetic and dorsal root ganglia and cloned neuronal cell lines (Martin and Feldman, 1993; Sumantran and Feldman, 1993).

IGFs and IGFFBPs also serve a role in tissue repair and regeneration (Skottner et al., 1990). IGFs are implicated in accelerated wound healing (Lynch et al., 1989; Steenfors and Jansson, 1992; Suh et al., 1992) and bone remodeling; IGFs modulate osteoblast proliferation and matrix formation (Skottner et al., 1990). IGFs enhance tissue regeneration, including pancreas (Smith et al., 1991), muscle (Jennische and Matejka, 1992), and kidney (Matejka and Jennische, 1992; Verstrepen et al., 1993). In the nervous system, IGF infusion enhances motor nerve regeneration (Sjoberg and Kanje, 1989; Near et al., 1992) and, in vitro, stimulates regeneration of adult sensory neurons (Fernihough et al., 1993).

The pattern of IGF expression during nervous system development coupled with its role in repair of nonnervous system tissues lead us to ask whether IGFs and/or IGFFBPs could play a facilitory role in nerve regeneration. As an initial step, we examined IGF-I and IGF-I receptor gene expression after sciatic nerve transection and observed increased transcripts for both factors in lesioned nerves. Schwann cells (SCs) were the main source of IGF-I immunoreactivity until 7 days after nerve transection, when macrophages containing high levels of IGF-I invaded the distal nerve stumps. To address further the role of the IGF axis in SC biology, we determined the in vitro expression pattern of the IGF-I receptor and the IGFFBPs. Transfected MT4H1 rat SCs expressed the IGF-I receptor and two IGFFBPs with molecular mass estimates of 24 and 31 kDa. Addition of IGF-I resulted in SC mitogenesis and enhanced levels of IGFBP-5. We investigated further the mechanism by which IGF-I regulates IGFBP-5 expression and found that IGF-I protected IGFBP-5 from proteolysis. Collectively, these data support the concept that in response to injury, invading macrophages produce IGF-I and SC express the IGF-I receptor, to facilitate axonal regeneration. This regenerative process may be augmented further by the ability of SC to secrete IGFFBPs, which in turn bind SC basement membrane and increase local IGF-I bioavailability.

MATERIALS AND METHODS

Dulbecco's modified Eagle's medium (DMEM) was purchased from GibcoBRL (Gaithersburg, MD, U.S.A.), fetal bovine serum (FBS) from Hyclone (Logan, UT, U.S.A.), and tissue culture supplies from Corning Glass Works (Corning, NY, U.S.A.) and Costar (Cambridge, MA, U.S.A.). Restriction enzymes were purchased from GibcoBRL and New England Biolabs (Beverly, MA, U.S.A.) and [³²P]-dCTP from NEN Du Pont (Boston, MA, U.S.A.). All other chemicals were purchased from Sigma Chemical Co. (St. Louis, MO, U.S.A.).

Sciatic nerve transection

Using aseptic technique, the sciatic nerve of anesthetized (ketamine-rompun, 1 cc/kg) adult Sprague-Dawley male rats (125–150 g; Spartan, Indianapolis, IN, U.S.A.) was exposed at the sciatic notch. After double ligation, the nerve was transected using fine scissors. The nerve stumps were sutured into adjoining muscles. At 3 or 7 days following surgery, the animals were killed by an overdose of ketamine-rompun. Both the distal and proximal nerve stumps were removed, stripped of excess connective tissue, and immediately frozen in liquid nitrogen. The contralateral unlesioned sciatic nerve served as the control. Nerves from 10–12 rats were pooled for each time point. All animal protocols were approved by the institutional animal care and use committee of the University of Michigan.

cDNA probes

A 300-bp rat IGFBP-5 cDNA (Dr. Shunichi Shimasaki, Whittier Institute, La Jolla, CA, U.S.A.), which encodes portions of the mature peptide, in pBluescript SK + plasmid was generated by *Sac*II and *Hind*III digestion. The IGF-I receptor cDNA was a gift from Dr. Haim Werner (Bethesda, MD, U.S.A.). The cDNA was a 265-bp insert excised from pGEM3 by *Eco*RI and *Sma*I digestion. A 1.8-kb *Pst*I–*Pst*I fragment of a chicken β -actin was excised from pAct-1 (Cleveland et al., 1980). All cDNA probes were purified using a Magic PCR Prep DNA Purification Kit (Promega, Madison, WI, U.S.A.).

RNase protection

Riboprobes were prepared from cDNA templates following the manufacturer's protocol (Promega). An exon 3 rat genomic IGF-I fragment was kindly provided by Dr. Peter Rotwein (Washington University, St. Louis, MO, U.S.A.). The rat IGF-I receptor cDNA, inserted in pGEM3, yields a protected band of 265 bp after hybridization (Werner et al., 1989). Probes were prepared and RNase protection performed as described previously, using 10- μ g RNA samples (Martin et al., 1991).

Immunohistochemistry

For immunohistochemical studies, three animals at each time point were examined following sciatic nerve transection. After they were killed, animals were perfused with 4% paraformaldehyde in 0.1 M phosphate buffer (pH 7.3). Proximal and distal nerve stumps and the contralateral control nerves were dissected and postfixed an additional 3 h at 4°C. Portions of sciatic nerves within 1 cm proximal or distal to the cut site were cryoprotected with sucrose and embedded in OCT (Miles), frozen above liquid nitrogen, and stored at –80°C. Sections (12 μ m) were cut on a cryostat, thaw mounted onto Tespa-coated slides (Pierce Chemicals, Rockford, IL, U.S.A.), and stored at –20°C.

Polyclonal rabbit anti-human IGF-I antibody (UB3-189) was obtained from the NIDDK, National Hormone and Pituitary Program by Drs. Louis E. Underwood and Judson J. Van Wyk (University of North Carolina, Chapel Hill, NC, U.S.A.) and used at 1:1,000 in phosphate buffer. Rabbit antibodies raised against IGFBP-5 were purchased from Upstate Biotechnology (Lake Placid, NY, U.S.A.), and used at 1:500. Frozen sections were processed for immunostaining employing an immunoperoxidase procedure using the chromogen 3,3'-diaminobenzidine as described previously (Sullivan and Feldman, 1994). IGF-I-immunoreactive specificity

was tested by use of either preimmune rabbit serum or primary antibody preabsorbed with excess recombinant IGF-I (1 $\mu\text{g}/\text{ml}$; BioSource, Camarillo, CA, U.S.A.).

Anti-IGFBP-5 is a polyclonal antiserum raised against human IGFBP-5 with low cross-reactivity against other IGFBPs. This antiserum can recognize both human and rat IGFBP-5. IGFBP-5 immunoreactivity was blocked by preabsorption of the antiserum with recombinant human IGFBP-5 (1 $\mu\text{g}/\text{ml}$; Austral Biologicals, San Ramon, CA, U.S.A.).

Cell culture

MT4H1 rat SCs were maintained in DMEM containing 10% FBS, 100 μM ZnSO_4 , and 100 $\mu\text{g}/\text{ml}$ G418 at 37°C in a humidified atmosphere with 10% CO_2 . Upon reaching 80% confluency, cells were subcultured with trypsin-EDTA into T75 flasks. In some experiments, cells were grown in defined glial cell medium, designated G3, containing 50 $\mu\text{g}/\text{ml}$ transferrin, 10 ng/ml biotin, and 30 nM selenium in DMEM (Bottenstein, 1985).

Cell proliferation assays

Cells were washed twice and plated with DMEM containing 0.3% FBS at a density of 3×10^4 cells/cm² in 96-well plates in increasing concentrations of IGF-I (0.1, 0.3, 1, and 3 nM). Cell proliferation was measured at 6, 24, and 48 h after plating with the MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] colorimetric assay (Hansen et al., 1989).

[methyl-³H]Thymidine ([³H]TdR) incorporation studies

MT4H1 cells were plated in 6-well plates (2.5×10^4 cells/cm²) in DMEM with 10% FBS and allowed to grow for 2 days. Medium was removed, and cells were rinsed twice with DMEM and incubated with increasing concentrations of IGF-I (0, 1, and 3 nM) for 24 h at 37°C. After 24 h, [³H]TdR (6 μCi , 6,700 Ci/mol; NEN Du Pont) was added to each well for a 2-h incubation at 37°C, and cells were processed as described previously (Feldman and Randolph, 1991).

Northern analysis

SCs were plated (3×10^6 cells per T75 flask) in DMEM and 10% FBS. After 2 days, the medium was removed, and cells were rinsed twice in DMEM and placed in G3 media only (G3 control) or G3 media containing increasing concentrations (1, 3, and 10 nM) of IGF-I or des(1-3)IGF-I. Des(1-3)IGF-I is an IGF-I analogue with a sixfold higher affinity for the IGF-I receptor but a lower affinity for the IGFBPs compared with IGF-I (Cascieri et al., 1991). After 24 h, the culture media were collected, RNA was isolated, and northern analysis and densitometry were performed as described previously (Martin and Feldman, 1993).

Western ligand blotting

Western ligand blots on conditioned media were prepared as described previously (Feldman and Randolph, 1994). Rinsed membranes were air dried and either placed directly on photostimulable storage phosphorimaging plates and scanned using a phosphorimager (Molecular Dynamics PhosphorImager, Sunnyvale, CA, U.S.A.) or exposed to X-Omat film (Eastman Kodak, Rochester, NY, U.S.A.) at -80°C for 1-7 days in the presence of an intensifying screen.

Immunoblotting

Samples of concentrated conditioned media (50 μg of protein) were mixed at a 5:1 ratio with 10 \times sodium dodecyl sulfate-sample buffer [100 mM Tris (pH 8), 10 mM EDTA, 10% sodium dodecyl sulfate, 100 mM dithiothreitol, 0.1% bromophenol blue, and 20% glycerol], and immunoblots were prepared, incubated for 2 h with IGFBP-5 antiserum (1:1,000), and visualized with enhanced chemiluminescence using the method of Siciliano and colleagues (1992).

Protease activity experiments

SC conditioned media were collected as described above. Briefly, confluent SC monolayer was washed twice, then incubated in G3 media without growth factors, and collected after a 24-h incubation. The protease activity experiments were performed in cell-free conditions: 100 ng of recombinant human IGFBP-5 was added to 50 μl of DMEM or conditioned media in the presence or absence of 10 nM IGF-I. Samples were incubated at 37°C for 2, 6, or 24 h, then separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis under reducing conditions. Western immunoblot analysis using IGFBP-5 antiserum was performed as described above.

RESULTS

IGF-I and IGF-I receptor expression following nerve transection

To examine IGF-I and IGF-I receptor gene expression during wallerian degeneration, we isolated RNA from proximal and distal nerve stumps 3 and 7 days after nerve transection. Nerves from 10-12 animals were pooled for each time point in each experiment, with the uncut contralateral sciatic nerve serving as the control. We detected low levels of IGF-I and IGF-I receptor mRNA in total, uncut sciatic nerves at each time point (Fig. 1A and B). Following nerve transection, there was an increase in IGF-I gene expression in both the proximal and distal stumps, with maximal IGF-I expression observed in the distal stump at 7 days (Fig. 1A). IGF-I receptor signal also increased in both stumps after nerve transection; however, there was no obvious difference between day 3 and day 7 (Fig. 1B).

To determine which cell type expressed IGF-I in the transected nerve model, we employed immunohistochemistry using anti-IGF-I antiserum. Currently, an antibody to rat IGF-I receptor is not available; therefore, we were unable to perform similar cellular immunolocalization of the receptor. In normal nerves, IGF-I staining was primarily present on SCs, fibroblasts, and capillary endothelial cells. In transverse sections, immunoreactive IGF-I was present in SC cytoplasm, forming a rim around the unstained myelin sheath (Fig. 2A). The proximal stumps of both day 3 and day 7 displayed a pattern of IGF-I immunoreactivity similar to intact nerves.

In the distal stumps 3 days after transection, morphological profiles characteristic of nerve injury were observed, including swollen axons and an increased number of endoneurial cells. IGF-I immunoreactivity

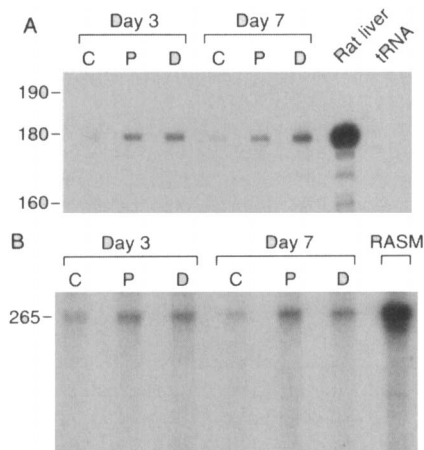


FIG. 1. IGF-I and IGF-I receptor expression following sciatic nerve transection. RNA was isolated from proximal (P) and distal (D) nerve stumps 3 and 7 days after sciatic nerve transection, as described in Materials and Methods. Nerves from 10–12 animals were pooled for each time point in each experiment. **A:** IGF-I RNase protection assay. An anti-sense RNA probe was transcribed from a 1-kb segment of the IGF-I gene that contained 182 bp of exon-3. Liver RNA protected the appropriate size fragment, whereas RNA from the uncut control nerve (C) produced a barely detectable fragment. Autoradiograph was exposed for 3 days. Data are from one of three representative experiments. **B:** IGF-I receptor RNase protection. An anti-sense RNA probe was transcribed from a 293-bp IGF-I receptor cDNA. Rat aortic smooth muscle (RASM) RNA protected the expected 265-bp fragment, whereas RNA from the uncut control nerve (C) produced a barely detectable fragment. RNA from nerve proximal (P) and distal (D) to the cut protected the expected 265-bp fragment at 3 and 7 days, with enhanced protection distal to the cut (D) at 7 days. Autoradiograph was exposed for 2 days. Data are from one of two representative experiments.

was detected in SCs, fibroblasts, and capillary endothelial cells (Fig. 2B). IGF-I immunoreactivity was not observed when the antiserum was preabsorbed with excess recombinant IGF-I, indicating antiserum specificity (data not shown).

Seven days after nerve transection, the pattern of IGF-I immunoreactivity continued. In addition, macrophages, which were strongly IGF-I positive, appeared within the endoneurial space and along the perineurial border, especially in the distal nerve stumps (Fig. 2D). The IGF-I immunoreactivity within SCs as well as other cellular components remained. In the distal stumps at 7 days, most axons had already degenerated; however, SCs ensheathing axonal debris could still be recognized (Fig. 2C). The quantity of S-100-positive cells rose significantly, indicating SC proliferation. In comparison with the proximal stumps and the control, IGF-I immunoreactivity decreased within the distal stump by day 7. However, more IGF-I-positive macrophages were detected (Fig. 2D), suggesting macrophages are the primary source of IGF-I at this later time point.

IGF-I promotes SC growth and DNA synthesis

In vitro models provide a valuable means of assessing growth factor action on specific cell types. To understand better the role of the IGF axis in the PNS, we employed cultured MT4H1 SCs, a rat cell line in which synthetic promoter elements from the mouse metallothionein-I promoter control the expression of SV40 T antigen. This cell line resembles untransfected secondary SC in both morphology and expression of P₀ and glial fibrillary acidic protein (Peden et al., 1989; Knight et al., 1993).

IGF-I is mitogenic in both neuronal and glial cells

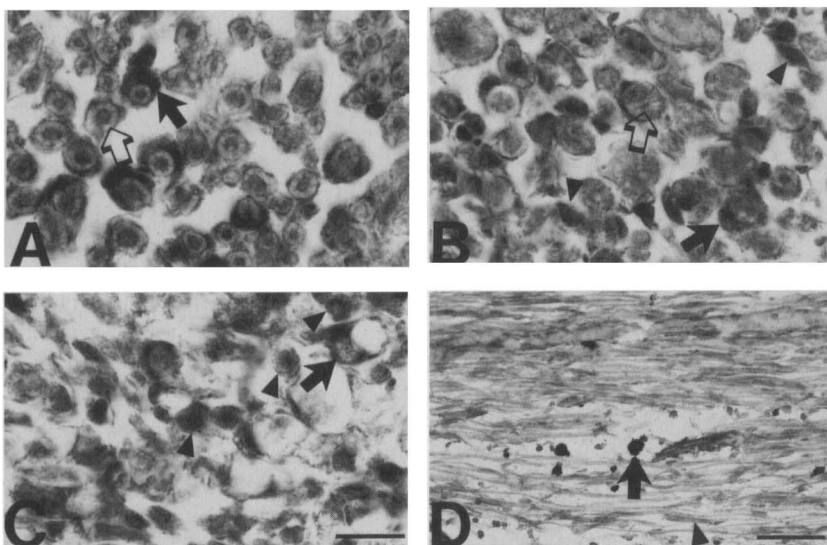


FIG. 2. IGF-I immunoreactivity after nerve transection. Rat sciatic nerves were transected and IGF-I immunohistochemistry was performed 3 and 7 days after surgery using polyclonal antiserum for IGF-I. **A:** IGF-I immunoreactivity in intact nerves. SCs (arrows) were immunostained, but not the myelin sheath (white arrow). **B:** IGF-I immunoreactivity at the distal stump at 3 days. Both SCs (arrow) and endoneurial cells (arrowheads) were IGF-I positive. Axonal degeneration was observed (white arrow). **C:** IGF-I immunoreactivity in distal nerve stumps at 7 days. Axonal degeneration was prominent. IGF-I-immunopositive SC either wrapped around axonal debris (arrow) or were proliferating (arrowheads). **D:** Longitudinal section of 7-day distal stump. Invading macrophages (arrow, D) appeared to be a major IGF-I source at this point. Less IGF-I immunoreactivity remained within SCs (arrowhead). Sections were representative of nerves of three animals with similar results. Bar = 20 μ m (A, B, C), or 100 μ m (D).

(DiCicco-Bloom and Black, 1988; McMorris and Dubois-Dalcq, 1988; Mozell and McMorris, 1988). To determine the effects of IGF-I on SC mitogenesis, SCs were grown in DMEM with 0.3% FBS with increasing concentrations of IGF-I for 3 days and cell number was determined by MTT assay (Hansen et al., 1989). IGF-I at 0.1 and 0.3 nM had no effect on growth at 6 and 24 h, but a modest effect was noted at 48 h compared with cells grown in 0.3% FBS alone. However, at 1 and 3 nM IGF-I, doses known to stimulate glial and neuroblast proliferation (DiCicco-Bloom and Black, 1988; McMorris and Dubois-Dalcq, 1988; Mozell and McMorris, 1988), IGF-I increased cell number within 6 h with a subsequent dose-dependent increase at 24 and 48 h (Fig. 3A). We next examined the effect of IGF-I on [³H]thymidine incorporation as a reflection of DNA synthesis. As observed with the MTT assay, IGF-I enhanced [³H]thymidine incorporation threefold after 24 h (Fig. 3B). These results demonstrate that IGF-I is mitogenic in MT4H1 SCs; however, although these data support the concept that IGF-I is mitogenic in primary SCs, they must be interpreted with caution because the current experiments were performed on transfected SCs only.

IGF-I and IGF-I receptor expression in cultured SCs

We speculated that cultured SCs would express IGF-I and IGF-I receptor mRNA similar to the results we observed in the transected sciatic nerve of the rat. To explore this possibility, RNA was isolated from cells grown in serum, defined G3 media, or G3 media with increasing concentrations of IGF-I (Peden et al., 1989; Knight et al., 1993). IGF-I mRNA was not detected with northern analysis or RNase protection (data not shown). As with the transected sciatic nerve preparation, MT4H1 SCs expressed mRNA for the IGF-I receptor (Fig. 4).

Effects of IGF-I on IGFBP gene and protein expression

The IGFbps play an integral role in the expression and action of the IGFs (Elgin et al., 1987; Clemmons, 1991; Shimasaki and Ling, 1991; Rosenfeld et al., 1993). Multiple neuronal and glial cell lines, including astrocytes, express one or more IGFbps at the gene and protein level (McCusker et al., 1990; Cheung et al., 1991; Clemmons, 1991). We initially examined IGFBP expression by western ligand blotting in MT4H1 SCs grown in defined G3 media for 24 h. When ¹²⁵I-IGF-II was used as the ligand, two IGFbps with molecular mass estimates of 31 and 24 kDa were detected (see Fig. 5A). In contrast, the 24-kDa IGFBP could not be detected by ¹²⁵I-IGF-I (Fig. 5B). The estimated molecular masses suggested that the 31-kDa band corresponded to IGFBP-5 and the 24-kDa band to IGFBP-4 or IGFBP-6 (Adamo et al., 1992; James et al., 1993; Jones et al., 1993). IGF-I treatment increased the intensity of the 31-kDa band in a dose-

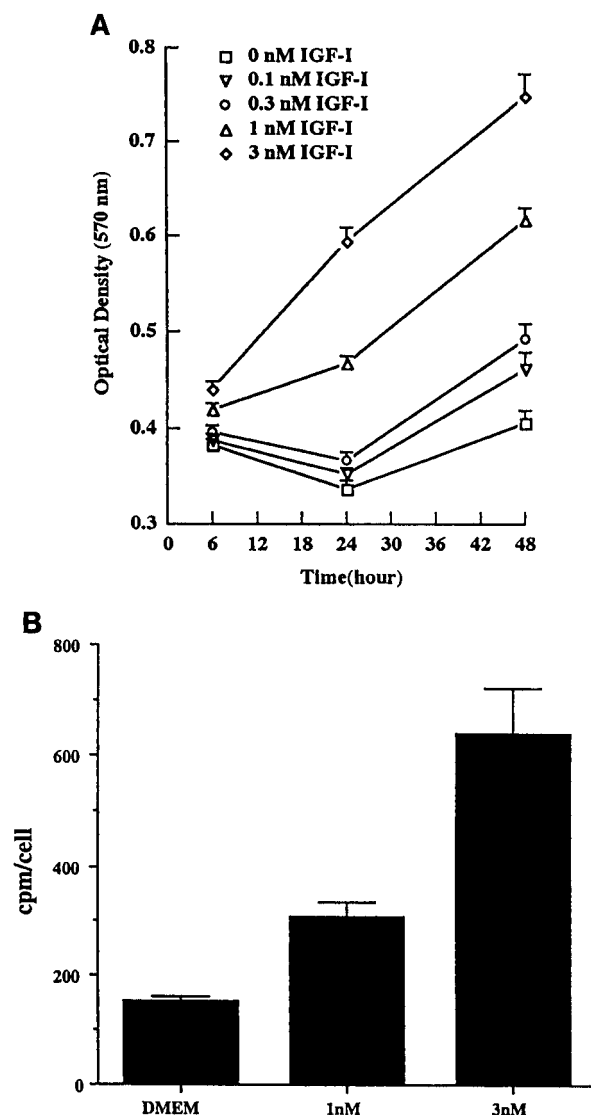


FIG. 3. IGF-I enhances SC proliferation. **A:** Cells were rinsed and plated in 96-well plates (3×10^4 cells/cm²) in DMEM containing 0.3% FBS \pm IGF-I. Cell number was measured by MTT assay after 6, 24, or 48 h. Relative optical density (570 nm) values are means of eight wells \pm SEM. Cell number was significantly higher at each time with 1 and 3 nM IGF-I treatment and at 48-h treatments ($p < 0.05$ compared with 0 nM IGF-I values, by unpaired, two-tailed *t* test). All data are from representative experiments performed three times. **B:** Effects of IGF-I on [³H]-TdR incorporation. Cells were plated at 2.5×10^4 cells/cm² in DMEM + 10% FBS for 2 days, then rinsed in DMEM and treated with DMEM \pm IGF-I. [³H]TdR incorporation and viable cell number (determined by trypan blue dye exclusion) were measured in parallel wells after 24 h. Values in B, expressed as counts per minute per cell, are means of triplicate wells \pm SEM. All data are from representative experiments performed three times.

dependent fashion compared with untreated cells with no change in the 24-kDa form (Fig. 5A). To verify the identity of the 31-kDa form of IGFBP, an immunoblot was probed with an IGFBP-5 antibody. Under reducing conditions, the immunoblot for IGFBP-5 re-

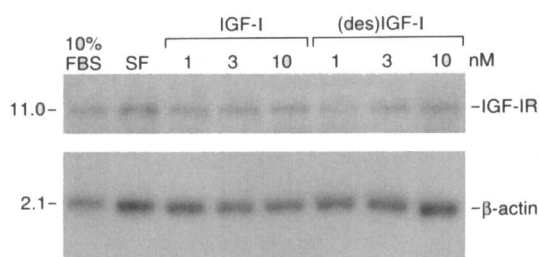


FIG. 4. Northern analysis of IGF-I receptor (IGF-IR) in MT4H1 SC. MT4H1 SC were plated at 4×10^4 cells/cm² in DMEM + 10% FBS. After 2 days, the medium was removed and the cells were placed in G3 media + 1, 3, or 10 nM IGF-I. Northern analysis on 20 μ g of RNA was performed with ³²P-labeled cDNA probes for IGF-IR, IGFBP-5, and β -actin, as indicated in Materials and Methods. RNA transcript sizes in kilobase units are listed at the left of the autoradiogram. Autoradiograph was exposed for 7 days. Data are from one of three representative experiments.

vealed a 23-kDa band that exhibited a dose-dependent expression in response to IGF-I treatment (Fig. 5C).

IGF-I enhancement of IGFBP-5 expression in SC conditioned media could represent an IGF-I receptor-mediated event or, alternatively, IGF-I protection of IGFBP-5 from proteolysis (Conover and Kiefer, 1993; Conover et al., 1993). To distinguish between these two possibilities, cells were treated with an IGF-I analogue, des(1-3)IGF-I, that has a sixfold higher affinity for the IGF-I receptor but reduced affinity for IGF-BPs (Cascieri et al., 1991). Des(1-3)IGF-I treatment of SCs decreased the level of measurable IGFBP-5 compared with IGF-I treatment (Fig. 5D). This implies that IGF-I enhances the presence of IGFBP-5 via a receptor-independent mechanism (Conover et al., 1993). Finally, equal amounts of protein were electrophoresed for each treatment, making it unlikely that changes in cell number were responsible for the differences observed in the amount of detectable IGFBP-5.

Because IGF-I can alter IGFBP gene expression in multiple cell lines (Krywicky and Yee, 1992), we next examined the effect of IGF-I and IGF analogue treatment on MT4H1 SC IGFBP-5 gene expression. Northern blots were prepared from cells treated for 24 h in G3 media alone or with the addition of 1, 3, or 10 nM IGF-I or des(1-3)IGF-I. Changes were not observed in the 6.0-kb IGFBP-5 transcript with these treatments (Fig. 6).

IGFBP-5 proteolysis in SC conditioned media

To demonstrate further that IGF-I enhances the presence of IGFBP-5 via a receptor-independent mechanism, we examined the effect of IGF-I on IGFBP-5 stability in cell-free MT4H1 SC conditioned media. rhIGFBP-5 (100 ng) was incubated in 50 ml of either DMEM alone (control) or MT4H1 SC-conditioned media in the presence or absence of 10 nM IGF-I (Fig. 7). After 2 h of incubation, a single 34-kDa band was detected in the control, corresponding to the published

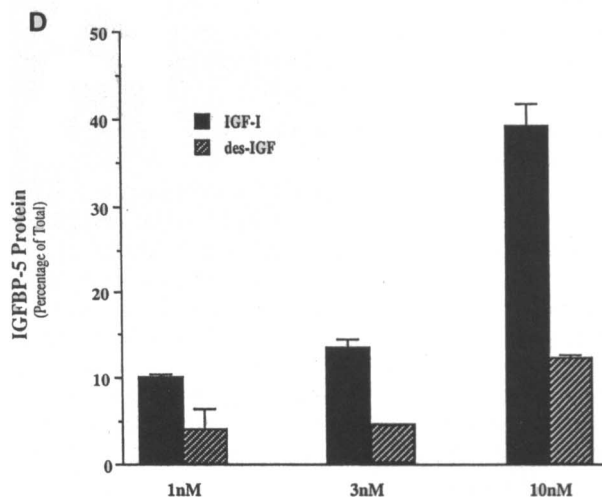
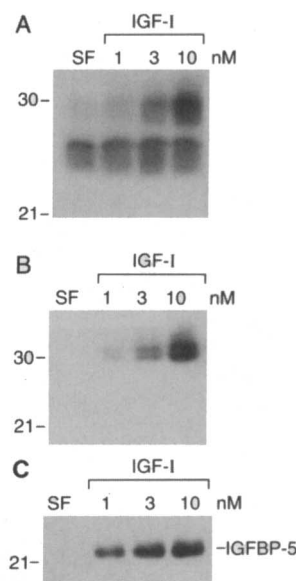


FIG. 5. IGFBP profile after IGF-I or des(1-3)IGF-I treatment. MT4H1 SC were plated at 4×10^4 cells/cm² in DMEM + 10% FBS. After 2 days, the medium was removed and the cells were placed in G3 media \pm 1, 3, or 10 nM IGF-I, or des(1-3)IGF-I. After 24 h, the media were collected and western ligand blot analysis was performed as described in Materials and Methods. Data are from one of three representative experiments. **A:** Western ligand blot analysis after IGF-I treatment. Using ¹²⁵I-IGF-II as the ligand, a dose-dependent increase was observed in a 31-kDa form with no change observed in a 24-kDa form under non-reducing condition. Autoradiograph was exposed for 5 days. **B:** The 31-kDa form was resolved into 29/31-kDa doublet when samples were electrophoresed on 12.5% sodium dodecyl sulfate-polyacrylamide gel. The 24-kDa form IGFBP was not detected when ¹²⁵I-IGF-I was the ligand. Autoradiograph was exposed for a week. **C:** Immunoblot of IGFBP-5. A similar dose-dependent pattern was observed on IGFBP-5 immunoblot. IGFBP-5 migrated to 23 kDa under reducing conditions. **D:** Phosphorimaging analysis of western ligand blots. Western ligand blots were placed directly on a photostimulable storage phosphorimaging plate and scanned using a phosphorimager. The percentage of counts in the total scan was determined from three separate experiments and the results are expressed as a percentage of total counts \pm SEM.

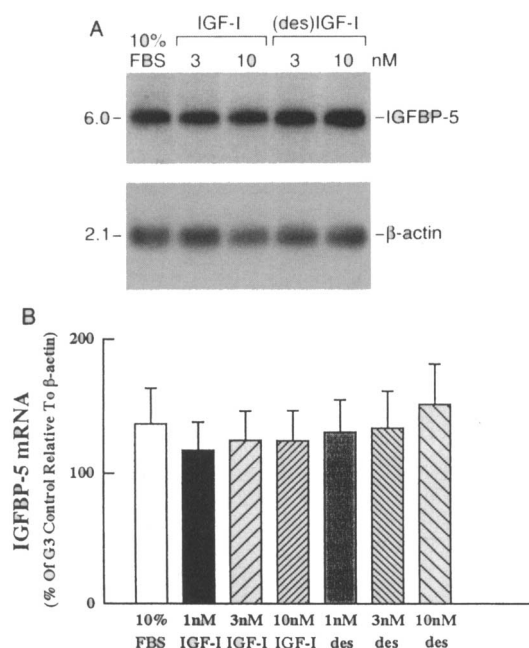


FIG. 6. Northern analysis of IGFBP mRNA after IGF-I and des(1-3)IGF-I treatment. MT4H1 SC were plated at 4×10^4 cells/cm² for 2 days in DMEM + 10% FBS, rinsed in DMEM as outlined in Materials and Methods, and then treated with 3 or 10 nM IGF-I or des(1-3)IGF-I. **A:** Autoradiograph from northern blot of IGFBP-5 and β -actin mRNA. RNA was isolated 24 h after treatment. Northern analysis on 20- μ g RNA samples was performed with ³²P-labeled cDNA probes for IGFBP-5, and β -actin, as indicated in Materials and Methods. RNA transcript sizes in kilobase units are listed at the left of each autoradiograph. Autoradiographs were exposed for 3 days (IGFBP-5) and 1 day (β -actin). **B:** Densitometric analyses of IGFBP-5 mRNA. Autoradiographs from multiple exposures of the experiments shown in A and at least one additional experiment were quantitated using relative optical densities, as described in Materials and Methods. Values are means \pm SEM of the ratio of IGFBP-5 mRNA relative optical density to β -actin relative optical density, expressed as percentage of serum-free untreated control.

molecular mass for rhIGFBP-5 under reducing conditions (Fig. 7, lane d) (Kiefer et al., 1992; Conover and Kiefer, 1993). The same band was also present in the conditioned media after 2 h with or without 10 nM IGF-I treatment with similar densities to the control (Fig. 7, lanes b and c). After 6 h of incubation, the same 34-kDa band was present in the control (Fig. 7, lane h) and in the conditioned media treated with IGF-I (Fig. 7, lane g). In contrast, in untreated conditioned media, a 23-kDa band appeared after 6 h of incubation and coincided with decreased density of the 34-kDa band (Fig. 7, lane f). These data imply that IGF-I protects IGFBP-5 from proteolysis by an IGFBP-5 protease, in agreement with previous reports (Conover and Keifer, 1993; Conover et al., 1993; Clemmons et al., 1994).

IGFBP expression in transected nerves

To demonstrate that SCs also express IGFBP-5 *in vivo*, we examined IGFBP-5 immunoreactivity in the

transected nerve model. Similar to IGF-I immunostaining, IGFBP-5 immunoreactivity was detected in SCs (Fig. 8A, arrows) as well as axons (Fig. 8A, white arrows) in intact nerves. Reaction product was homogeneously distributed throughout the SC cytoplasm with an intense rim of immunostaining in intact sciatic nerve. Unlike IGF-I, IGFBP-5 immunoreactivity of SCs was not influenced by nerve transection. IGFBP-5 immunoreactivity remained localized within SCs in the proximal and distal stumps 3 days after transection (Fig. 8B and C). The IGFBP-5 antiserum also stained proliferating SCs in the distal stump on day 7 (Fig. 8D). In contrast, the axonal IGFBP-5 immunoreactivity faded after nerve transection (compare Fig. 8A and B).

DISCUSSION

The IGF family of ligands, receptors, and binding proteins is instrumental in normal nervous system development (Daughaday and Rotwein, 1989; Liu et al., 1991) and in the repair of nervous tissues (Sjoberg and Kanje, 1989; Jennische and Matejka, 1992; Near et al., 1992; Fernyhough et al., 1993). In this study, we examined the pattern of IGF-I and IGF-I receptor gene expression after sciatic nerve transection. In both proximal and distal nerve segments, we observed increased IGF-I and IGF-I receptor transcripts, with a selective enhancement of IGF-I mRNA in the nerve segment distal to the lesion. Immunohistochemical examination of transected nerves confirmed the presence of IGF-I production by SCs. Our findings parallel those

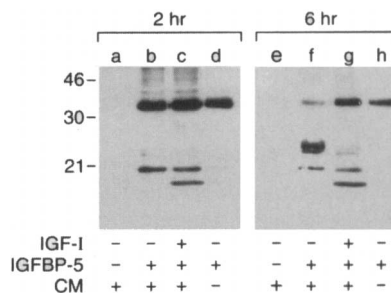


FIG. 7. IGFBP-5 immunoblots revealed IGF-I protection of IGFBP-5 from proteolysis in SC conditioned media (CM). SC conditioned media (50 μ l) were incubated with or without 100 ng of rhIGFBP-5 in the presence or absence of 10 nM IGF-I at 37°C for 2 or 6 h. The samples were separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis under reducing conditions, followed by immunoblotting for IGFBP-5. IGFBP-5 was undetectable in SC conditioned media (a, e). There was no difference between the 34-kDa IGFBP-5 intensities of the conditions with (c) or without (b) IGF-I in SC conditioned media after 2 h of incubation. In contrast, obvious degradation was observed after 6 h of incubation. The intensity of the 34-kDa band without IGF-I (f) decreased significantly in comparison with the conditions plus IGF-I or positive control (g, h) accompanying the appearance of a 23-kDa proteolytic fragment (f). Data are from one of the three experiments with similar results.

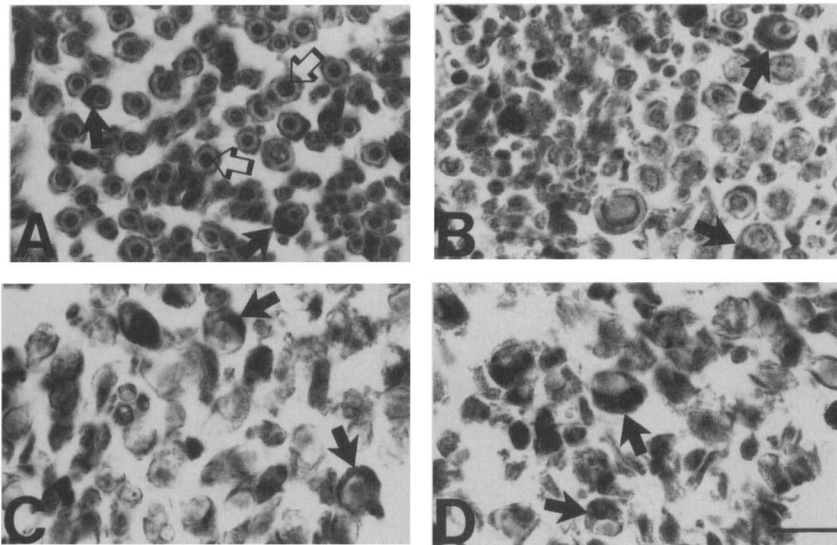


FIG. 8. IGFBP-5 immunoreactivity in nerve transection model. IGFBP-5 immunohistochemistry studies were performed using a polyclonal antiserum against IGFBP-5. **A:** Intact nerve section. **B:** Proximal nerve stump of 3 days. **C:** Distal nerve stump of 3 days. **D:** Distal nerve stump of 7 days. The IGFBP-5 immunoreactivity was consistently detected in SC cytoplasm (black arrows, A, B, C, D) as well as axons in intact nerves (A, white arrows). Representative sections are from three animals. Bar = 20 μ m.

of Hansson and colleagues who observed patterns of enhanced SC production of IGF-I with peripheral nerve injury (Hansson et al., 1986).

Similar to our results with IGF-I and IGF-I receptors, SCs, including those that ensheath motor axons, increase production of nerve growth factor (NGF) and NGF receptors following nerve injury (Taniuchi et al., 1986, 1988; Heumann et al., 1987). Transection of selective ventral roots entering the sciatic nerve results in increased NGF receptor expression by SCs associated with degenerating motor neurons, but not intact sensory neurons (Taniuchi et al., 1988). Like IGF-I and NGF, transforming growth factor- β expression increases with nerve transection, particularly in the distal nerve stump (Scherer et al., 1993). In contrast, the expression of ciliary neurotrophic factor, which is axon dependent, decreases in the distal stumps of transected nerves (Friedman et al., 1992).

We also found that IGF-I promoted *in vitro* SC growth in a dose-dependent manner. Several polypeptide growth factors are mitogenic to SC *in vitro*, including transforming growth factor- β , acidic and basic fibroblast growth factors, and platelet-derived growth factor (Eccleston, 1992). These factors elevate intracellular cyclic AMP which, in the presence of serum, results in SC proliferation (Eccleston, 1992). High-dose insulin can substitute for serum in the presence of both fibroblast growth factor and platelet-derived growth factor and the cyclic AMP agonist, forskolin (Eccleston, 1992). Because the mitogenic effects of high-dose insulin are mediated via the IGF-I receptor (Flier et al., 1986), the growth-promoting effects of insulin on SC suggest not only the presence of IGF-I receptors on SC but also a physiological role for IGF-I in SC growth.

SC may not be the sole source of polypeptide growth factors in the PNS. Trophic factors may also be pro-

duced by different cell types present in transected nerves, including macrophages, endothelial cells, and fibroblasts (Blexrud et al., 1990). In the current study, we found that macrophages, which infiltrate injured nerves and remove the tissue debris, serve as another source of IGF-I. It is interesting that activated macrophages not only remove tissue debris but also promote SC growth. *In vitro*, conditioned media of cultured macrophages activated by phagocytosis of myelin debris promote cultured SC growth via mitogenic peptide factors (Baichwal et al., 1988). Our results suggest one of these mitogenic peptide factors could be IGF-I, which was highly expressed by macrophages in the distal stump 7 days after nerve transection. In contrast to the expression of both IGF-I and the IGF-I receptor by SC *in vivo*, we did not observe IGF-I gene expression *in vitro* in MT4H1 SC in the current study. These observations imply that (a) the ability of SCs to express IGF-I *in vivo* is lost *in vitro*; (b) the observed *in vivo* expression of IGF-I within SCs of the cut nerve is secondary to other IGF-I-producing cells, including fibroblasts (Clemmons and Van Wyk, 1985), endothelial cells (Delafontaine and Lou, 1993; Fath et al., 1993), and/or macrophages (Nagaoka, 1990); (c) the IGF-I immunoreactivity in nerve sections is either receptor or binding protein associated; (d) IGF-I immunoreactivity decreases in the distal stumps at 7 days indicate that the ability of SC to express IGF-I is axon dependent. An IGF-I *in situ* hybridization study of the proximal and distal nerve stumps as well as the neurons of the spinal cord and dorsal root ganglia would be one way to distinguish between these possibilities.

We did observe IGF-I receptor expression in MT4H1 SC. These data parallel our findings on IGF-I receptor expression in proximal and distal transected sciatic nerve segments. Collectively, our results support the concept that one role of growth factor receptors

on SC surfaces is to present growth factors to regenerating axons, e.g., IGF-I binding to the IGF-I receptor is one means by which SC could increase the local bioavailability of IGF-I to regenerating axons.

An alternative means of presenting IGF-I to regenerating axons is via the IGFBPs. In the current study, we speculated that SCs secrete IGFBPs which, in turn, bind to SC plasmalemmal matrix and serve as reservoirs for IGF-I. Immunohistochemical studies demonstrated the presence of IGFBPs in the SCs of transected nerves. We then determined whether IGFBPs were produced by MT4H1 SC. Utilizing western ligand blots, immunoblots, and northern analysis, we report that MT4H1 SCs secrete IGFBP-5 and a 24-kDa IGFBP which, according to the molecular mass, could be IGFBP-4 or IGFBP-6 (Kiefer et al., 1992). The evidence that this 24-kDa IGFBP can be detected only by the ligand ^{125}I -IGF-II suggests it is IGFBP-6 because IGFBP-6 binds preferentially to IGF-II and cannot always be detected by ligand blots using ^{125}I -IGF-I (Kiefer et al., 1992).

Although the precise mechanisms remain unknown, IGFBP-5 is associated with cellular growth and differentiation (Mohan et al., 1989; Andress and Birnbaum, 1992; James et al., 1993; Jones et al., 1993). In vitro, IGFBP-5 regulates osteoblast proliferation (Mohan et al., 1989; Andress and Birnbaum, 1992), and expression of IGFBP-5 markedly increases during muscle cell differentiation (James et al., 1993). It is likely that IGFBP modulation of IGF bioavailability underlies the profound effects IGFBPs have on growth. Recently, Clemmons and colleagues have demonstrated that IGFBP-5 can bind to fibroblast ECM and sequester IGF-I for local fibroblast growth (Jones et al., 1993). The ability of ECM components to dissociate IGFBP-5 from an IGF-I/IGFBP-5 complex further supports the concept that IGFBP-5 targets IGF-I near cell surface IGF-I receptors. Thus, the interactions between SC ECM, IGFBP-5, and IGF-I receptors may regulate local IGF-I bioavailability (Arai et al., 1994).

In cultured fibroblasts and osteoblasts, IGFBP-5 appears to be a stimulatory factor for IGF's mitogenic effects. Clemmons and colleagues have demonstrated that ECM-associated IGFBP-5 (but not free IGFBP-5 in the media) can potentiate IGF-I's mitogenic effects (Jones et al., 1993). Similarly, a truncated 24-kDa IGFBP-5, isolated from osteoblast-conditioned media, can enhance IGF-I-dependent osteoblast proliferation (Andress and Birnbaum, 1992). The present study reveals the colocalization of IGF-I and IGFBP-5 in a model of nerve transection, implying that IGFBP-5 may play a positive role in modulating IGF-I trophic effects on SC growth, and on axonal regeneration in PNS.

Our experiments also suggest that IGF-I-induced changes in IGFBP-5 are not mediated via the IGF-I receptor. Although IGF-I enhanced the presence of IGFBP-5, des(1-3)IGF-I, an analogue that retains the

ability to bind the IGF-I receptor but has a marked decreased affinity for the IGFBPs (Forbes et al., 1988), did not increase the concentration of IGFBP-5. Thus, an IGF analogue that cannot bind IGFBP-5, but can bind the IGF-I receptor, cannot mimic IGF-I's effects. We also found that IGF-I protected IGFBP-5 from proteolysis in a cell-free system. These data imply that the IGF-I-dependent increased concentration of IGFBP-5 is a receptor-independent phenomenon and represents the protection of an extracellular IGF-I/IGFBP-5 complex from proteolysis by an IGFBP-5 protease. In agreement with the current results, Conover and colleagues reported that addition of IGF-I enhanced IGFBP-5 levels in vitro in osteosarcoma-conditioned media via a receptor-independent mechanism (Conover et al., 1993) and that IGF-I protected IGFBP-5 from proteolysis (Conover et al., 1993; Durham et al., 1994). Similar results were also observed in a fibroblast system by Clemmons and colleagues, and a calcium-dependent serine protease responsible for IGFBP-5 proteolysis has been identified and isolated (Clemmons et al., 1994). Our results suggest the presence of a similar protease that can cleave IGFBP-5 into proteolytic fragments in SC conditioned media.

In summary, although recent work (Eccleston, 1992), including the current study, emphasizes the importance of neurotrophic factors in nerve growth, the mechanisms underlying successful axonal regeneration and reformation of axon-SC contacts remain unknown. One theory is that neurotrophins bind to SC surfaces producing a trophic factor-enriched SC plasmalemmal matrix that enhances axonal regrowth (Eccleston, 1992). The present study supports this concept; we have shown that, to facilitate regeneration in response to nerve injury, SCs produce IGF-I and IGF-I receptor. This regenerative process is augmented further by the ability of SCs to secrete IGFBPs, which, in turn, may bind SC basement membrane and increase local IGF-I bioavailability.

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