

A Role for Insulin-Like Growth Factor-I in the Regulation of Schwann Cell Survival

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During postnatal development in the peripheral nerve, differentiating Schwann cells are susceptible to apoptotic death. Schwann cell apoptosis is regulated by axons and serves as one mechanism through which axon and Schwann cell numbers are correctly matched. This regulation is mediated in part by the provision of limiting axon-derived trophic molecules, although neuregulin-1 (NRG-1) is the only trophic factor shown to date to support Schwann cell survival. In this report, we identify insulin-like growth factor-I (IGF-I) as an additional trophic that can promote Schwann cell survival *in vitro*. We find that IGF-I, like NRG-1, can prevent the apoptotic death of postnatal rat Schwann cells cultured under conditions of serum

withdrawal. Moreover, we show that differentiating Schwann cells in the rat sciatic nerve express both the IGF-I receptor (IGF-I R) and IGF-I throughout postnatal development. These results indicate that IGF-I is likely to control Schwann cell viability in the developing peripheral nerve and, together with other findings, raise the interesting possibility that such survival regulation may switch during postnatal development from an axon-dependent mechanism to an autocrine and/or paracrine one.

Key words: peripheral nervous system; glia; Schwann cell; myelination; apoptosis; insulin-like growth factor-I

Programmed death is a cell fate adopted by multicellular organisms to control cell number in development, homeostasis, and defense. Such physiological cell death, which most often proceeds through a series of well-defined alterations in cellular morphology termed apoptosis, is genetically programmed and results in cell suicide. Importantly, apoptosis must be tightly regulated such that only certain cells are specified to die, and it is now clear that both cell-extrinsic and cell-intrinsic signals can render cells susceptible to apoptosis (Steller, 1995; Fraser et al., 1996; White, 1996). During development, programmed cell death is often controlled by the positive selection of cells via specific ligand–receptor tyrosine kinase interactions.

Approximately 50% of immature postmitotic neurons normally undergo apoptotic death. In this instance, extrinsic signals delimit neuronal number and establish appropriate innervation patterns through competition for limiting target-derived trophins (Barde, 1989; Oppenheim, 1991). Death occurs when neurons fail to secure access to these trophins and consequently are unable to

suppress a constitutive cell death program (Raff, 1992; Raff et al., 1993). Apoptosis also occurs in differentiating oligodendrocytes and Schwann cells, the myelinating glial cells of the CNS and peripheral nervous system, respectively. Approximately 50% of newly generated oligodendrocytes die during development of the rodent optic nerve (Raff et al., 1993; Barres and Raff, 1994), and differentiating Schwann cells undergo apoptotic death during both embryonic (Ciutat et al., 1996) and early postnatal (Grinspan et al., 1996; Syroid et al., 1996; Nakao et al., 1997) development. Cell death in these glial lineages is thought to be in part regulated by the limited availability of axon-derived trophic factors and represents one mechanism whereby the appropriate stoichiometry between glia and the axons they myelinate is achieved (Raff et al., 1993; Barres and Raff, 1994; Zorick and Lemke, 1996).

Understanding the regulation of Schwann cell apoptosis requires an identification of the trophins that control Schwann cell viability. Apoptosis in Schwann cell precursors *in vitro* (Dong et al., 1995) and in postnatal Schwann cells *in vivo* (Grinspan et al., 1996; Trachtenberg and Thompson, 1996; Kopp et al., 1997) and *in vitro* (Syroid et al., 1996) can be prevented by neuregulin-1 (NRG-1), which is normally supplied by axons (Carraway and Burden, 1995; Lemke, 1996). Another set of factors that may mediate Schwann cell survival is the insulin-like growth factors (IGFs)-I and -II. As autocrine and/or paracrine factors, the IGFs are thought to play an important role in the development and regeneration of the nervous system (Hansson, 1993; Ishii et al., 1993; Lewis et al., 1993a; de Pablo and de la Rosa, 1995). IGF-I, for example, regulates oligodendrocyte development and myelination *in vitro* and *in vivo* (McMorris et al., 1986, 1993; McMorris and Dubois-Dalcq, 1988; Saneto et al., 1988; Mozell and McMorris, 1991; Barres and Raff, 1994; Yao et al., 1995). Reduced

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oligodendrocyte numbers and a severe myelination defect have been observed in IGF-I and IGF-I receptor (IGF-I R) mutant mice (Liu et al., 1993; Beck et al., 1995), and a corresponding enhancement of myelin content has been noted in transgenic mice that overexpress IGF-I (Carson et al., 1993; Ye et al., 1995). For the Schwann cell lineage, both IGF-I and IGF-II can potentiate the survival of cultured Schwann cell precursors (Gavrilovic et al., 1995) and can act as mitogens and differentiation factors for postnatal Schwann cells *in vitro* (Schumacher et al., 1993; Stewart et al., 1996). In this report, we provide evidence that IGF-I acts as an autocrine and/or paracrine survival factor for postnatal Schwann cells.

MATERIALS AND METHODS

Cell culture. Schwann cells were prepared from postnatal day 3 (P3) rat sciatic nerve and purified to >99.5% homogeneity essentially as described previously (Brookes et al., 1979). Cells were routinely maintained by plating on poly-L-lysine (100 μ g/ml; Sigma, St. Louis, MO)-coated 10 cm tissue culture Petri dishes (Falcon) in DMEM (Life Technologies, Gaithersburg, MD) containing fetal calf serum (FCS; 10%; HyClone, Logan, UT), forskolin (Fsk; 2 μ M; Calbiochem, La Jolla, CA), and recombinant NRG-1 β [NRG- β ; amino acids 177–228 encompassing the β -type epidermal growth factor-like domain (Lemke, 1996)] (10 ng/ml). These represent the preassay conditions for all experiments. Schwann cells grown on eight-well chamber slides (Lab-Tek) were plated on poly-L-lysine (100 μ g/ml) and laminin (10 μ g/ml; Life Technologies). Schwann cell survival assays were performed as described previously (Syroid et al., 1996).

Growth factors. Human IGF-I and IGF-II were from Boehringer Mannheim (Indianapolis, IN). Human recombinant NRG- β was kindly provided by Dr. Duanzhi Wen (Amgen, Thousand Oaks, CA). Bovine serum albumin (BSA; Sigma; 0.1% final) was added to all growth factors as a carrier protein.

Generation of IGF-I receptor antibody. New Zealand White rabbits were immunized with a peptide corresponding to the C-terminal 14 amino acids (KNERALPLQSSSTC) of the β -subunit of the IGF-I R cross-linked to keyhole limpet hemocyanin with glutaraldehyde. IGF-I R antibody was affinity-purified using an affinity column containing the peptide covalently coupled to cyanogen bromide-activated sepharose beads (Pharmacia, Piscataway, NJ) and then was eluted using 0.1 M glycine, pH 2.5, and 0.1 M ethanolamine, pH 11.5. The two fractions were pooled and dialyzed against PBS buffer (137 mM NaCl, 2.7 mM KCl, 4.3 mM Na₂HPO₄·7H₂O, and 1.4 mM KH₂PO₄, pH 7.3). The IGF-I R antibody was then partially concentrated by centrifugation through a Centrprep 10,000 filter (Amicon, Beverly, MA).

The specificity of the IGF-I R antibody was examined by immunoprecipitation of the IGF-I R from lysates of [³⁵S]methionine-labeled 293 and NIH3T3 cells. A 97 kDa protein corresponding to the IGF-I R β -subunit was selectively immunoprecipitated by the IGF-I R antibody and not by the preimmune serum. Preincubation of the IGF-I R antibody with an excess of the IGF-I R peptide inhibited immunoprecipitation of the 97 kDa protein. Similarly, the mouse monoclonal antibody directed against the α -subunit of the IGF-I R (α IR3; Oncogene Research Products) coimmunoprecipitated the 97 kDa IGF-I R β -subunit. This protein comigrated with the protein immunoprecipitated with the IGF-I R antibody, and both antibodies yielded similar amounts of immunoprecipitated IGF-I R β -subunit. Western blot analyses confirmed that the 97 kDa immunoprecipitated protein corresponded to the β -subunit of the IGF-I R. Furthermore, transfection of NIH3T3 cells with the human IGF-I R cDNA resulted in a dramatic increase in the detection of the 97 kDa IGF-I R protein. Increased tyrosine phosphorylation of the IGF-I R β -subunit was also observed in an anti-phosphotyrosine Western blot after stimulation of cells with IGF-I. Cross-reactivity of the IGF-I R antibody with the insulin receptor was examined using immunoprecipitation and Western blot analyses of cells isolated from IGF-I R $-/-$ mice (Liu et al., 1993). Detection by the Enhanced Chemiluminescence kit (Amersham, Arlington Heights, IL) did not reveal a 97 kDa protein even after prolonged exposures. Data supporting the integrity of the IGF-I R antibody are not shown here (Arbet-Engels et al., 1999). For immunohistochemical analyses, preincubation of IGF-I R antibody with IGF-I R peptide specifically precluded IGF-I R immunoreactivity on P8 rat sciatic nerve sections in a dose-dependent manner (data not shown).

Antibodies. Mouse monoclonal anti-neurofilament (NF) antibody (Sigma) was used at 1:100 dilution. Mouse monoclonal anti-protein zero (P₀) antibody was kindly provided by Dr. J. J. Archelos (Bayerische Julius-Maximilians-Universität, Würzburg, Germany) and used at 1:2000 dilution. Rabbit antiserum to IGF-I (UB2-495) was kindly provided by the National Institute of Diabetes and Digestive and Kidney Diseases (NIDDK), National Hormone and Pituitary Program, by Drs. Louis E. Underwood and Judson J. Van Wyk (University of North Carolina, Chapel Hill, NC) and used at 1:20 dilution. Affinity-purified rabbit polyclonal anti-IGF-I R antibody was used undiluted.

MTT survival assay. MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; Sigma] was added to cells at a concentration of 0.5 mg/ml, and the cells were then further incubated at 37°C for 1 hr. The number of cells in the bottom of each well exhibiting a positive blue granular reaction product was assessed using bright-field microscopy.

[³H]Thymidine incorporation assay. Schwann cells were dissociated by trypsinization, washed once in DMEM containing 10% FCS to inactivate the trypsin, and then washed an additional five times in cold serum-deficient DMEM to remove serum. Cells were plated onto eight-well chamber slides (10,000 cells/well) in either DMEM containing 10% FCS, Fsk (2 μ M), and NRG- β (50 ng/ml) or IGF-I (50 ng/ml), DMEM containing IGF-I only (50 ng/ml), or DMEM containing BSA only. Each culture condition was performed in duplicate. [³H]Thymidine (Amersham) was then added to all wells at a concentration of 0.03 μ Ci/ml, and after a further 24 hr the cells were fixed for 20 min at room temperature in 4% paraformaldehyde/PBS buffer. Processing of slides and assessment of [³H]thymidine incorporation in two independent experiments were as described previously (Syroid et al., 1996). Cells that had four or more silver grains over their nuclei were considered to be [³H]thymidine-positive.

Ribonuclease protection analysis. Total cellular RNA from cultured Schwann cells and Sprague Dawley rat sciatic nerves was prepared and analyzed by ribonuclease protection assay as described previously (Chomczynski and Sacchi, 1987; Krieg and Melton, 1987). Antisense RNA probes were synthesized using the Maxiscript *in vitro* transcription kit (Ambion) from templates containing the following inserts: a 690 base pair *Sau3A*-*NlaIV* fragment containing rat IGF-I genomic sequences encompassing exons 1, 3, and 4 (Adamo et al., 1991); a 265 base pair *EcoRI*-*RsaI* rat IGF-I R cDNA fragment encompassing 5'-untranslated sequences, the coding region for the signal peptide, and the first 53 amino acids of the α -subunit (Werner et al., 1989) (both templates kindly provided by Dr. Derek LeRoith, Diabetes Branch, NIDDK, National Institutes of Health, Bethesda, MD); and a 150 base pair mouse cyclophilin cDNA fragment encompassing 3'-untranslated sequences [nucleotides 586–736 (Hasel and Sutcliffe, 1990)]. This cyclophilin RNA probe gives rise to multiple protected RNA fragments (150 base pairs and lower molecular weight species) when hybridized with the rat RNA used in this study because of several nucleotide mismatches with the corresponding rat cyclophilin cDNA sequence (Danielson et al., 1988). Cyclophilin RNA probes of low specific activity were generated by including 0.05 mM UTP in transcription reactions and using only one-fifth (10 μ Ci) of the usual amount of [α -³²P]UTP (Amersham). RNA (1–18 μ g) or tRNA (10 μ g) was cohybridized with 80,000 cpm of either IGF-I or IGF-I R RNA probe and 30,000 cpm of cyclophilin RNA probe, as indicated. The relative quantity and integrity of RNA used in each experiment were confirmed on agarose gels (data not shown). Protected probe/RNA hybrids were resolved on 6% polyacrylamide and 8 M urea denaturing gels.

Sciatic nerve transection. Adult and P1 Sprague Dawley rats were gas anesthetized using isoflurane, and unilateral sciatic nerve transections were performed just proximal to the sciatic notch as described previously (Zorick et al., 1996). Nerve sections (3–5 mm) were excised to prevent axon regrowth within the time frame of the experiment. Animals were killed either 24 hr or 7 d after transection, at which time both the unlesioned contralateral sciatic nerve and the entire length of the distal stump of the transected nerve were isolated and processed for RNA preparation. Distal stumps of transected nerves were closely examined before isolation to ensure that regeneration was completely precluded.

Immunohistochemical staining. Immunohistochemical staining of rat sciatic nerve sections was performed as described previously (Zorick et al., 1996). For fluorescence microscopy, immunolabeled sections were incubated for 1 hr at 25°C with anti-rabbit/FITC and/or anti-mouse/Texas Red secondary antibodies (1:100; Jackson ImmunoResearch, West Grove, PA). Frozen cross sections of rat sciatic nerves were prepared as described previously (Syroid et al., 1996; Zorick et al., 1996).

RESULTS

IGF-I inhibits Schwann cell death *in vitro*

We have shown previously that Schwann cells isolated from P3 rat sciatic nerve undergo apoptosis *in vitro* after serum withdrawal and that β -isoforms of NRG-1 (NRG- β) can prevent this apoptotic death (Syroid et al., 1996). To assess further the survival requirements of postnatal Schwann cells, we examined whether IGF-I or IGF-II can promote Schwann cell viability *in vitro* under serum-free conditions using an MTT incorporation assay. MTT is a chromogenic reagent that is converted to a granular blue product when metabolized by active mitochondria. As shown in Figure 1A, only 15–20% of cells remain viable 3 d after serum withdrawal when plated in DMEM in the absence of exogenous trophic factors (e.g., in BSA alone). However IGF-I (50 ng/ml), like NRG- β , can effectively maintain Schwann cell number at ~80% of the initial plating number. In contrast, IGF-II at the same concentration only partially inhibits Schwann cell death, a result that may reflect the reduced affinity of IGF-II for the IGF-I R (Jones and Clemmons, 1995). The ability of IGF-I to prevent Schwann cell death was dose-dependent, exhibiting a dose response similar to that of NRG- β (Fig. 1B). Schwann cells cultured in serum-deficient DMEM containing IGF-I converted MTT into high levels of the chromogenic blue reaction product (Fig. 1C), an activity that was also dose-dependent (data not shown). These observations are indicative of strong metabolic activity in these cells, which were elongated and bipolar in morphology and displayed extended processes, features characteristic of healthy cultured Schwann cells. The few weakly MTT-positive Schwann cells observed in serum-deficient DMEM containing only BSA were rounded and lacked processes (Fig. 1C, arrows). Both these cells and MTT-negative cells exhibited extensive nuclear fragmentation and chromatin condensation (Syroid et al., 1996), which are characteristic features of apoptotic cells (compare also with Fig. 2C,D). Viable cells cultured in DMEM containing only IGF-II gave rise to a less intense MTT reaction product than did those cultured in IGF-I (data not shown), results that are in keeping with the more modest ability of IGF-II to promote Schwann cell survival (Fig. 1A).

IGF-I is a survival factor for postnatal Schwann cells

To address the possibility that IGF-I-mediated maintenance of Schwann cell number under serum-free conditions (Fig. 1A) might be caused by the induced proliferation of a subset of cells refractory to cell death, we performed [³H]thymidine incorporation assays. As shown for a positive control, the majority of cells cultured under proliferative conditions in the combined presence of Fsk and NRG- β incorporated [³H]thymidine (Fig. 2A; 86% of nuclei labeled; 1566 nuclei counted) (Syroid et al., 1996). Similarly and as reported previously (Schumacher et al., 1993; Stewart et al., 1996), IGF-I is also mitogenic for Schwann cells when combined with Fsk (Fig. 2B; 87% of nuclei labeled; 1811 nuclei counted). However, cells subjected to serum withdrawal were not labeled when cultured in DMEM containing IGF-I alone (Fig. 2C; 0.01% of nuclei labeled; 1721 nuclei counted), indicating that IGF-I is not mitogenic for Schwann cells under these conditions. As expected, cells grown in DMEM containing only BSA (most of which were dead or dying) also failed to incorporate [³H]thymidine (Fig. 2D; 0.002% of nuclei labeled; 1614 nuclei counted). These results demonstrate that IGF-I can function as a survival factor for cultured postnatal Schwann cells.

Schwann cells express the IGF-I receptor *in vitro*

The IGF-I R is a receptor tyrosine kinase composed of two ligand-binding extracellular α -subunits associated with two transmembrane kinase domain-containing β -subunits that transduces both IGF-I and IGF-II signals (Jones and Clemmons, 1995; LeRoith et al., 1995). It has been demonstrated previously that proliferating Schwann cells express the IGF-I R *in vitro* (Schumacher et al., 1993; Stewart et al., 1996). To examine whether Schwann cells cultured under apoptotic, serum-free conditions (those used for the assays of Fig. 1) also express the IGF-I R, a ribonuclease protection analysis was performed using total RNA derived from rat Schwann cells at 4 and 24 hr after plating in unsupplemented DMEM and from Schwann cells maintained for 48 hr in DMEM containing serum and stimulated with 20 μ M Fsk, conditions that lead to the induction of myelin-specific genes and thus represent an *in vitro* paradigm for differentiating Schwann cells (Lemke and Chao, 1988). As shown in Figure 3A, the IGF-I R RNA probe was specifically protected under all conditions and gave rise to the expected 265 base pair-protected fragment (Werner et al., 1989). These results demonstrate that the IGF-I R precursor mRNA is expressed both in Fsk-stimulated Schwann cells and in Schwann cells cultured in the absence of serum.

Schwann cells are induced to express IGF-I *in vitro* when switched to unsupplemented medium

Because Schwann cells require the exogenous provision of IGF-I to prevent their apoptotic death *in vitro* (Fig. 1A), we reasoned that cells cultured under apoptotic, serum-free conditions do not express IGF-I. To examine this possibility directly, a ribonuclease protection assay was performed using total RNA isolated from both Fsk-stimulated Schwann cells and Schwann cells subjected to serum withdrawal, as described above. As shown in Figure 3B, IGF-I mRNA was not detected in Fsk-stimulated Schwann cells even after extended autoradiographic exposure. As expected, IGF-I expression was also absent in preassay Schwann cells—Schwann cells cultured in DMEM containing serum and 2 μ M Fsk (data not shown). Interestingly, a very low level of IGF-I mRNA was detected after Schwann cells were cultured for 4 hr in unsupplemented DMEM, and this expression was upregulated after 24 hr (Fig. 3B), a time at which ~50% of cells are already dead (Fig. 1A) (Syroid et al., 1996). The multiple protected IGF-I RNA probe fragments evident in Figure 3B derive from a heterogeneous population of IGF-I transcripts resulting from multiple major transcription initiation sites within exon 1 of the IGF-I gene and correspond to the previously reported protected fragments of 573, 530, and 428 base pairs. [Additional lower molecular weight protected fragments were also detected and probably reflect additional downstream transcription initiation sites (data not shown) (Adamo et al., 1991).] Because the majority of Schwann cells become committed to a pathway of apoptotic death at 2–4 hr after serum withdrawal (Syroid et al., 1996), an IGF-I-mediated autocrine mechanism in early IGF-I-producing cells may account for the 15–20% of cells that survive prolonged serum-deficient culture (Fig. 1A). These surviving cells are unlikely to represent a distinct Schwann cell population present in the nerve at the time of dissociation, because much previous work (e.g., Lemke and Chao, 1988) has suggested that essentially all neonatal Schwann cells, independent of their differentiation state at the time of dissociation, are phenotypically plastic and revert to a common embryonic phenotype after being placed in DMEM supplemented with serum alone.

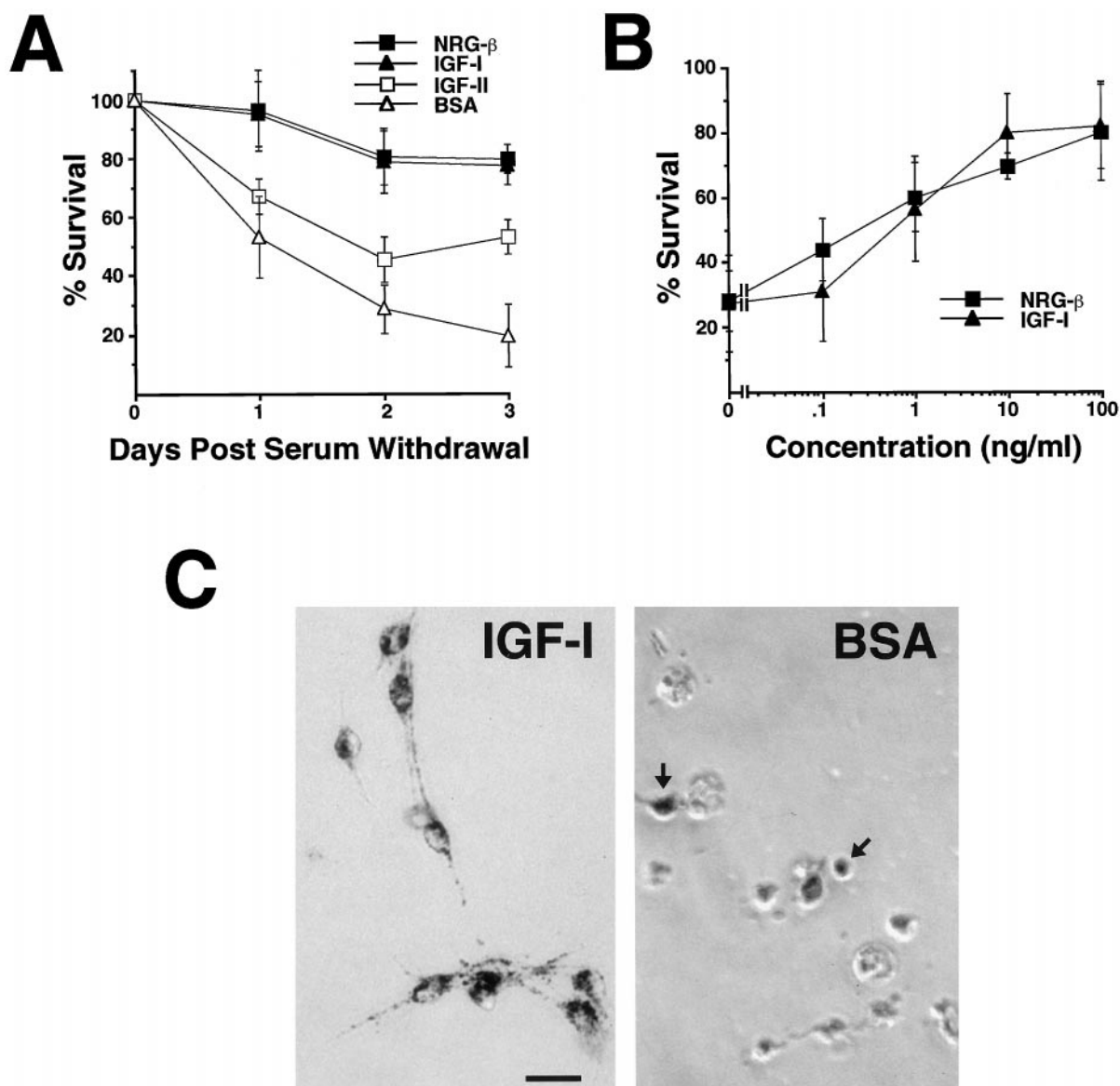


Figure 1. IGF-I promotes the survival of postnatal Schwann cells *in vitro*. Schwann cells were washed five times in serum-deficient DMEM to remove serum and then plated on multiple microwell plates in DMEM either in the presence of the factor indicated or with BSA. Viable cells were identified and counted using an MTT incorporation assay at daily intervals over a 3 d period. *A*, IGF-I, like NRG- β , can promote Schwann cell survival, whereas BSA cannot. IGF-II can only partially inhibit Schwann cell death. *B*, Dose–response curves for IGF-I- and NRG- β -mediated Schwann cell survival 3 d after serum withdrawal shows the cellular morphology and the strong MTT positivity of Schwann cells cultured in DMEM in the presence of IGF-I, as analyzed by phase contrast microscopy. Arrows denote the few remaining viable, weakly MTT-positive Schwann cells in the BSA-negative control. Factor concentration was 50 ng/ml unless otherwise indicated. Error bars represent the SD from the mean in survival activity derived from at least three separate experiments. Scale bar, 5 μ m.

Schwann cells express the IGF-I receptor in the developing sciatic nerve

To assess the *in vivo* relevance of our *in vitro* results, the expression of both the IGF-I R and IGF-I was examined in rat sciatic nerves during the first 2 postnatal weeks, the time at which Schwann cells are maximally susceptible to apoptotic death (Grinspan et al., 1996; Syroid et al., 1996; Nakao et al., 1997). Sciatic nerves were isolated at various postnatal days, and either total RNA was prepared and used for ribonuclease protection analyses to examine mRNA expression or nerves were processed for sectioning and subjected to immunohistochemical analyses to examine protein expression.

As shown in Figure 4*A*, ribonuclease protection detected the IGF-I R proreceptor transcript as early as the day of birth (P0), and mRNA expression was maintained thereafter, into the mature nerve. Because Schwann cells make up the vast majority of mRNA-containing cells in the peripheral nerve, it is likely that they are responsible for the observed IGF-I R mRNA expression. To examine directly which cells express the IGF-I R, immunohistochemical analyses were performed on P10 sciatic nerve cross sections using a polyclonal antibody to the IGF-I R (Arbet-Engels et al., 1999). This antibody was generated using a synthetic peptide comprising the C-terminal 14 amino acids of the β -subunit of the IGF-I R (see Materials and Methods) and was

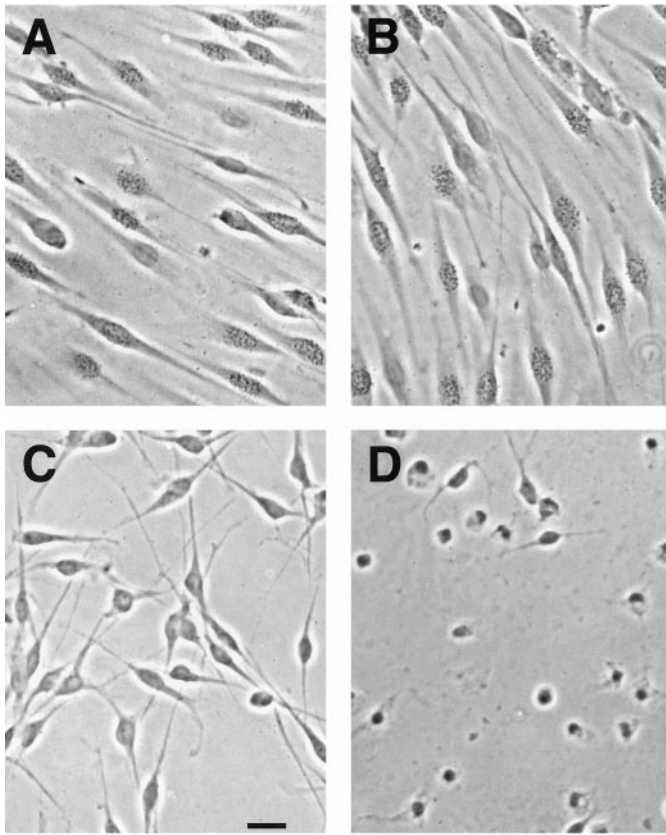


Figure 2. IGF-I is a Schwann cell survival factor *in vitro*. Schwann cells were subjected to serum withdrawal and then cultured for 24 hr on eight-well chamber slides in DMEM containing either 10% FCS, NRG- β (50 ng/ml), and Fsk (2 μ M) (A); 10% FCS, IGF-I (50 ng/ml), and Fsk (2 μ M) (B); IGF-I (50 ng/ml) (C); or BSA (D). [3 H]Thymidine (0.03 μ Ci/ml) was then added, and the cells were further incubated for 24 hr, after which the cells were fixed and [3 H]thymidine incorporation was assessed by phase contrast microscopy. Schwann cells cultured in the presence of IGF-I under conditions of serum withdrawal (C) do not incorporate [3 H]thymidine, indicating that IGF-I is mediating survival activity. Scale bar, 5 μ m.

chosen for our study because it generates much less background than do commercially available antibodies against the IGF-I R α -subunit (data not shown). As shown in Figure 5A, Schwann cells of P10 sciatic nerve exhibit strong IGF-I R immunoreactivity. IGF-I R expression was detected from the day of birth onward (data not shown), results that are in agreement with the IGF-I R transcript expression profile (Fig. 4A). As reported previously (Caroni and Grandes, 1990; Lewis et al., 1993b; Reinhardt et al., 1993), we find that axons also display low IGF-I R immunoreactivity (Fig. 5A).

Schwann cells express IGF-I in the developing sciatic nerve

To determine whether IGF-I is expressed in the rat sciatic nerve during postnatal development, a ribonuclease protection analysis was performed using total RNA derived from sciatic nerves isolated at various stages of postnatal development. As shown in Figure 4B, multiple IGF-I transcripts were detected at the day of birth, and mRNA expression was maintained thereafter, both throughout postnatal development and within the adult nerve. To examine whether Schwann cells account for this expression, double-labeling immunohistochemical analyses were performed

on cross sections of P10 rat sciatic nerve using a polyclonal antiserum to IGF-I, an antiserum that has been used previously to examine IGF-I protein expression *in vivo* in rat sciatic nerve (Cheng et al., 1996), and using antibodies to axon-specific NF and peripheral myelin P₀. As shown in Figure 5B, IGF-I immunoreactivity strongly localizes to the cytoplasm of Schwann cells. Consistent with the IGF-I mRNA profile (Fig. 4B), Schwann cell expression of IGF-I is maintained throughout postnatal development and in the mature nerve (data not shown). Although IGF-I appears to be excluded from the myelin sheath (marked by P₀), a very low level of IGF-I immunoreactivity also localizes to axons [Fig. 5B, bottom (IGF-I alone)]. These results, taken together, demonstrate that Schwann cells are the major source of IGF-I in the postnatal rat sciatic nerve and are in agreement with previous work implicating axons as an additional source of IGF-I in developing and mature peripheral nerves (D. E. Syroid and G. Lemke, unpublished observations) (Hansson et al., 1987; Garcia-Segura et al., 1991; Lievre et al., 1991).

Developing Schwann cells maintain expression of the IGF-I receptor and IGF-I after loss of axonal contact

There is considerable evidence implicating IGF-I as an important trophic molecule in the regeneration of the nervous system (Hansson, 1993; Ishii et al., 1993; Lewis et al., 1993a; de Pablo and de la Rosa, 1995), and consistent with this view, several studies report the induction of IGF-I in both injured and regenerating peripheral nerves. For example, increased IGF-I immunoreactivity has been noted in both Schwann cells and axons proximal to the site of lesion in adult sciatic nerve and, to a lesser degree, in Schwann cells of the distal nerve stump undergoing Wallerian degeneration (Hansson et al., 1986, 1987, 1988; Hansson, 1993). Upregulation of IGF-I and IGF-I R mRNA has also been reported in both proximal and distal nerve stumps after mature sciatic nerve transection (Glazner et al., 1994; Pu et al., 1995; Cheng et al., 1996). Interestingly, mature Schwann cells remain viable, at least over several weeks, after loss of axonal contact in the adult nerve (D. E. Syroid, T. J. Kilpatrick, and G. Lemke, unpublished observations) (Weinberg and Spencer, 1978; Grinspan et al., 1996), whereas developing Schwann cells in the neonatal nerve undergo apoptosis within 24 hr of transection (Grinspan et al., 1996; Trachtenberg and Thompson, 1996; Kopp et al., 1997). Because IGF-I is a potent survival factor for postnatal Schwann cells and because Schwann cells synthesize IGF-I during postnatal development (this study), it is conceivable that the observed induction of Schwann cell death in neonatal nerves undergoing Wallerian degeneration may be caused by a corresponding impairment in IGF-I signaling. Expression of the IGF-I R and IGF-I was therefore examined in both neonatal (P1) and adult rat sciatic nerves during Wallerian degeneration. Unilateral sciatic nerve transections were performed, and both the distal stump of lesioned nerves and the intact contralateral nerves were isolated 24 hr after transection; total RNA was prepared and used for ribonuclease protection analyses. As an additional control, IGF-I R and IGF-I expression was examined 7 d after transection in adult distal nerve stumps, a sufficient period in the mature nerve for axonal degeneration. As shown in Figure 6, A and B, expression of the IGF-I R proreceptor transcript and the multiple IGF-I transcripts, respectively, in the distal stump of transected P1 nerves is maintained at approximately the same level relative to that in transected adult nerves, at both 24 hr and 7 d after transection. Moreover, IGF-I R and IGF-I expression in transected nerves is no different relative to that in the corre-

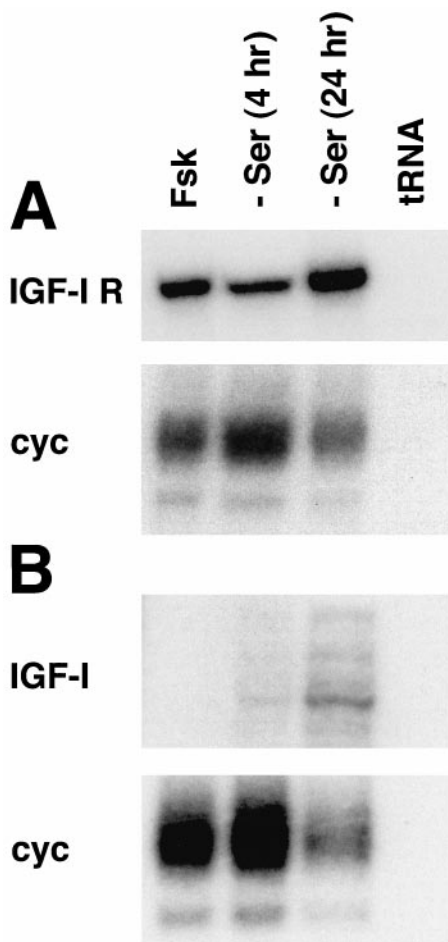


Figure 3. Schwann cells express both *IGF-I R* and *IGF-I* mRNAs *in vitro*. Ribonuclease protection analyses were performed using total RNA (6 or 18 μ g per hybridization reaction with *IGF-I R* or *IGF-I* RNA probes, respectively) isolated from subconfluent Schwann cells cultured either in DMEM containing 10% FCS and 20 μ M Fsk (*Fsk*) or in DMEM alone (*-Ser*) for 4 or 24 hr or using 10 μ g of *tRNA* as a negative control, as indicated. *A*, The *IGF-I R* is expressed by Schwann cells grown in serum- and *Fsk*-supplemented DMEM and by cells cultured in DMEM alone. Autoradiographic exposure was 10 hr. *B*, *IGF-I* is not expressed in Schwann cells grown in DMEM containing serum and forskolin; however a very low level of *IGF-I* mRNA can be detected in Schwann cells cultured for 4 hr in unsupplemented DMEM, and this expression is upregulated after 24 hr. Autoradiographic exposure was 63 hr. *Bottom (A, B)*, The relative cyclophilin RNA level (*cyc*) from corresponding cohybridization reactions using a cyclophilin-loading control probe is shown. Autoradiographic exposure in these panels was 18 hr (*A*) and 5 hr (*B*).

sponding contralateral control nerves. Previous reports have suggested that these mRNAs are even upregulated in adult sciatic nerves undergoing Wallerian degeneration (Glazner et al., 1994; Pu et al., 1995; Cheng et al., 1996), although we have not observed this. In any event, our results indicate that developing Schwann cells in the neonatal sciatic nerve maintain expression of both *IGF-I* and the *IGF-I R* after loss of axonal contact and that the level of expression appears to be comparable with that of Schwann cells found in mature nerves undergoing Wallerian degeneration.

DISCUSSION

In this study, we demonstrate that *IGF-I* is a survival factor for postnatal rat Schwann cells *in vitro* and that *IGF-I*-mediated

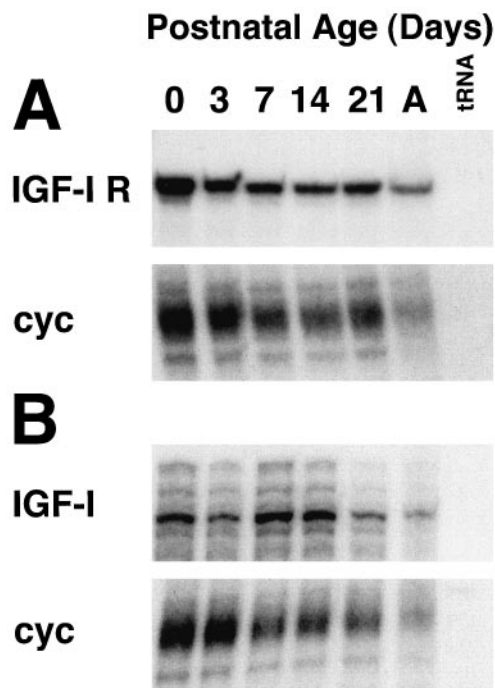


Figure 4. The *IGF-I R* and *IGF-I* mRNAs are expressed in the sciatic nerve throughout postnatal development. Ribonuclease protection analyses were performed using total RNA (6 μ g per hybridization reaction) isolated from adult rat sciatic nerves (*A*) and nerves at various stages of postnatal development or using 10 μ g of *tRNA* as a negative control, as indicated. *A, B*, The *IGF-I R* (*A*) and *IGF-I* (*B*) are expressed throughout postnatal development and in the adult nerve. Autoradiographic exposure was 18 hr (*A*) and 24 hr (*B*). *Bottom (A, B)*, The relative cyclophilin RNA level (*cyc*) from corresponding cohybridization reactions using a cyclophilin-loading control probe is shown. Autoradiographic exposure in these panels was 75 hr (*A*) and 24 hr (*B*).

survival activity is almost certainly transduced via the *IGF-I R*. We also demonstrate that differentiating Schwann cells in the rat sciatic nerve express both *IGF-I* and the *IGF-I R* throughout postnatal development and that *IGF-I* and *IGF-I R* expression by both developing and mature Schwann cells is maintained during Wallerian degeneration. These results implicate an *IGF-I*-mediated autocrine loop in Schwann cell survival regulation.

Schwann cell number is tightly controlled during early postnatal development, and this regulation is mediated by the action of multiple axon-associated signals that modulate both Schwann cell proliferation and apoptotic death. Schwann cell and axon numbers are eventually matched such that within the first postnatal week in rodents, a one-to-one relationship between myelinating Schwann cells and large diameter axons has largely been attained, whereas nonmyelinating Schwann cells remain associated with multiple axons (Webster, 1993). The critical role of axons in the regulation of Schwann cell development during this period can primarily be ascribed to the provision of trophic factors (Reynolds and Woolf, 1993; Mirsky and Jessen, 1996; Zorick and Lemke, 1996). The postulated role of *NRG-1* as an axon-derived Schwann cell mitogen (Fig. 2*A*) (Marchionni et al., 1993; Levi et al., 1995; Morrissey et al., 1995) and survival factor (Grinspan et al., 1996; Syroid et al., 1996; Trachtenberg and Thompson, 1996; Kopp et al., 1997) fits well within this mechanistic context, because *NRG-1* is expressed in both developing and mature sensory and motoneurons (Carraway and Burden, 1995; Lemke, 1996). The results presented in this study identify *IGF-I* as an additional

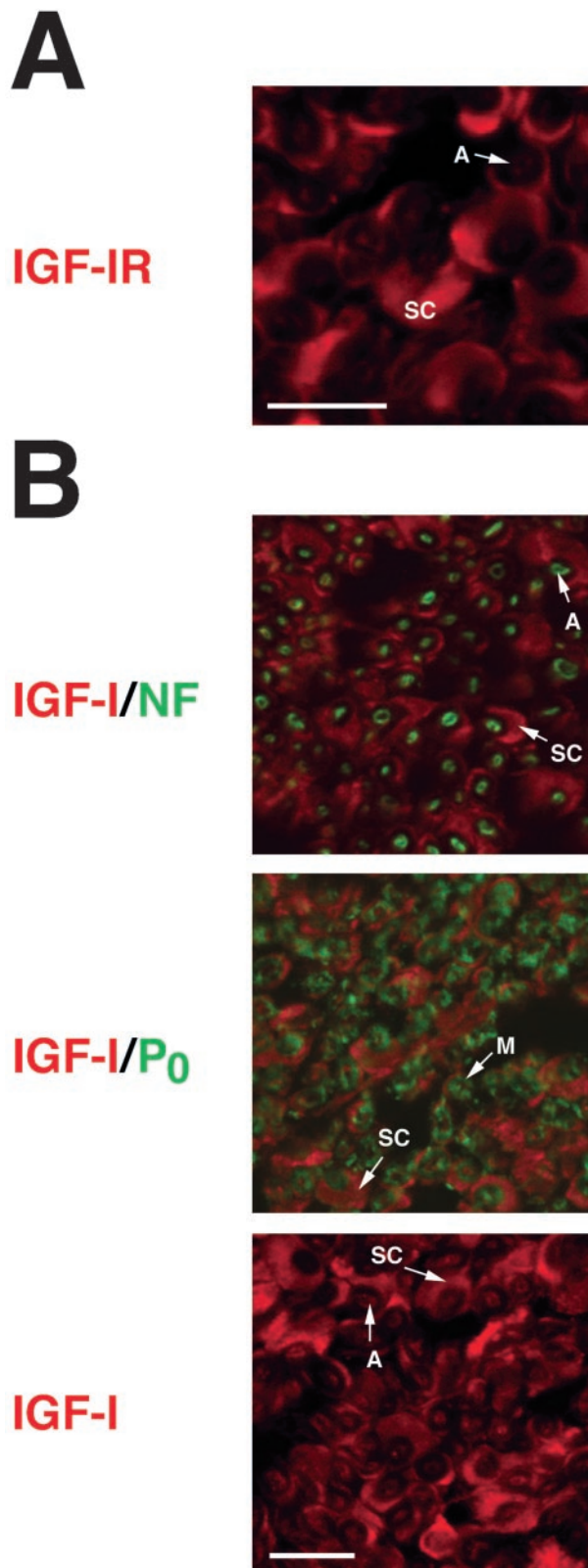


Figure 5. Schwann cells express *IGF-IR* and *IGF-I* throughout postnatal development. Immunohistochemical analyses were performed on cross sections of P10 rat sciatic nerve to examine *IGF-IR* and *IGF-I* protein expression. *A*, *IGF-IR* protein (red) is localized primarily to Schwann cells (SC), whereas axons (A) also express a very low level of the *IGF-IR*. *B*, Double-labeling analyses were performed to examine *IGF-I* protein expression (red) relative to axon-specific NF protein expression (green) and myelin-specific P₀ protein expression (green), as indicated. *IGF-I* is

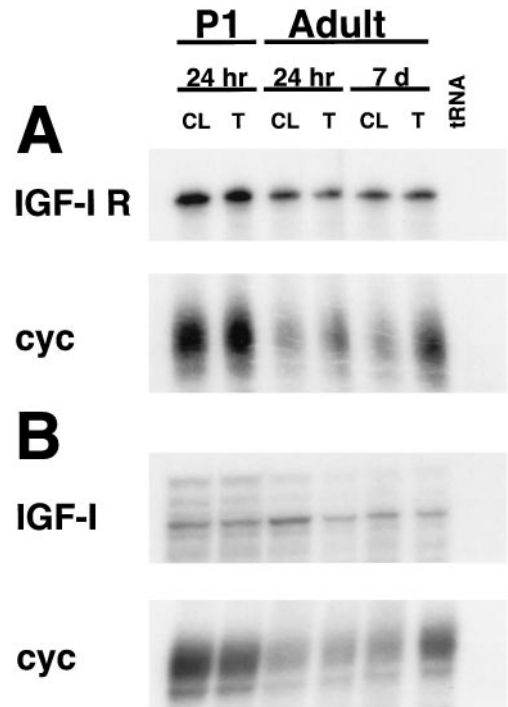


Figure 6. Expression of *IGF-IR* and *IGF-I* mRNA is maintained in the neonatal rat sciatic nerve during Wallerian degeneration. Unilateral transections of P1 and adult rat sciatic nerve were performed, and ribonucleic acid protection analyses were performed using total RNA (1 μ g per hybridization reaction) isolated from both the intact contralateral nerve (CL) and the transected distal nerve stump (T) at 24 hr and 7 d after transection or using 10 μ g of *tRNA* as a negative control, as indicated. *A*, *B*, *IGF-IR* (*A*) and *IGF-I* (*B*) expression in the degenerating neonatal nerve is maintained at approximately the same level relative to that in the contralateral control nerve and is equivalent to that in degenerating adult nerves, at both 24 hr and 7 d after transection. *Bottom* (*A*, *B*), The relative cyclophilin gene expression (*cyc*) from corresponding cohybridization reactions using the cyclophilin RNA probe is shown. We find that cyclophilin mRNA levels (which, like β -actin or glyceraldehyde phosphate dehydrogenase, are commonly used for loading controls in protection assays) are consistently lower in adult nerves relative to that in neonatal nerves (see Fig. 4), whereas cyclophilin mRNA levels increase in adult nerves undergoing Wallerian degeneration for an extended period (7 d). Autoradiographic exposure was 96 hr (*A*) and 40 hr (*B*).

trophic molecule that may regulate Schwann cell viability during postnatal development. However, because Schwann cells themselves are the major IGF-I⁺ cell type in the neonatal nerve, our data suggest that axon-independent mechanisms also play an important role in controlling Schwann cell development. Because IGF-I, like NRG-1, is also capable of acting as a strong mitogen for cultured Schwann cells when assayed in combination with Fsk (Fig. 2*A,B*) (Schumacher et al., 1993; Stewart et al., 1996), it is likely that these molecules function together as important signals in the regulation of multiple stages (e.g., proliferation and survival) of Schwann cell development.

What are the molecular mechanisms that regulate Schwann cell viability during peripheral nerve development, and how does this

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highly expressed by Schwann cells (SC). Although *IGF-I* immunoreactivity is excluded from the myelin sheath (M), a very low level of *IGF-I* protein also localizes to axons (A). This is most evident in the *bottom section* immunostained for *IGF-I* alone. Images were collected using a scanning confocal microscope. Scale bars, 10 μ m.

relate to the matching of Schwann cell and axonal numbers? The finding that transection of neonatal rat optic nerve gives rise to widespread oligodendrocyte cell death (Fulcrand and Privat, 1977; David et al., 1984; Barres et al., 1993) led Raff and colleagues to conclude that oligodendrocyte survival is critically dependent on axon-derived signals during postnatal development. These investigators have proposed a model in which the competition for limiting axon-derived trophic molecules serves to match correctly the number of oligodendrocytes to the number and length of axons they myelinate (Raff et al., 1993; Barres and Raff, 1994). Axonal interactions also control the survival of early postnatal Schwann cells. Transection of neonatal rat sciatic nerve results in an upregulation of Schwann cell apoptosis, and this axotomy-induced cell death can be prevented by the exogenous provision of NRG-1, indicating that NRG-1 is one such axon-derived trophic factor for Schwann cells (Grinspan et al., 1996; Trachtenberg and Thompson, 1996; Kopp et al., 1997). Exogenous NRG-1 can also block naturally occurring Schwann cell death (Grinspan et al., 1996), indicating that the amount or availability of NRG-1 in the neonatal nerve may be limiting. These observations suggest that during early postnatal development, excess Schwann cells that either have lost axonal contact or are otherwise unable to secure limiting trophic support would become susceptible to apoptotic death and thus would be eliminated from the developing nerve.

However, the survival of only a fraction of postnatal Schwann cells is axon-dependent, because previous studies have shown that only a subset of Schwann cells are induced to undergo apoptotic death after transection of neonatal sciatic nerves (Grinspan et al., 1996; Trachtenberg and Thompson, 1996). These studies also demonstrate a progressive decrease in the proportion of Schwann cells subject to axotomy-induced apoptosis as transections are performed at progressively later stages of postnatal development. Indeed, little Schwann cell death can be detected after transection of mature sciatic nerves (Syroid, Kilpatrick, and Lemke, unpublished observations) (Weinberg and Spencer, 1978; Grinspan et al., 1996). These observations suggest that the survival requirements for Schwann cells switch to an axon-independent mechanism at later stages of postnatal development. Our results raise the interesting possibility that this later control of Schwann cell viability may, at least in part, require an IGF-I-mediated autocrine and/or paracrine loop. Such a loop would inhibit Schwann cell apoptosis after nerve injury and thereby promote axonal regeneration, which critically depends on the presence of Schwann cells (Fawcett and Keynes, 1990; Bunge, 1993; Son et al., 1996). In agreement with this hypothesis, mature peripheral nerves maintain regenerative capacity after injury (Fawcett and Keynes, 1990), while functional reinnervation by neonatal axons is impaired (McArdle and Sansone, 1977; Thompson and Jansen, 1977; Betz et al., 1980; Dennis and Harris, 1980), a difference that may be partially ascribed to the loss of Schwann cells through apoptotic death in neonates (Grinspan et al., 1996; Trachtenberg and Thompson, 1996; Kopp et al., 1997). Consistent with this model, Schwann cell expression of both IGF-I and the IGF-I R is maintained within injured and regenerating adult peripheral nerves (Fig. 6) (Hansson et al., 1986, 1987, 1988; Hansson, 1993; Glazner et al., 1994; Pu et al., 1995; Cheng et al., 1996).

Although our work strongly supports the hypothesis that IGF-I is a key component of Schwann cell survival regulation, the data presented above also indicate that IGF-I signaling by itself is unlikely to explain the observed switch to axon-independent survival that is seen in mature nerves, because we find that both

the ligand and its receptor are expressed by Schwann cells from the day of birth. Why do some Schwann cells in the neonatal nerve undergo apoptosis during early postnatal development even though they express IGF-I (Figs. 4B, 5B)? Why do some developing Schwann cells remain susceptible to apoptotic death after loss of axonal contact, unlike mature Schwann cells in adult nerves undergoing Wallerian degeneration, even though both Schwann cell populations continue to express IGF-I R and IGF-I (Fig. 6)? There are several possibilities. (1) The subset of Schwann cells in the developing nerve that normally undergoes cell death and the enhanced proportion that die after transection may remain dependent on axonally derived trophic support and thus may represent a subset of Schwann cells that does not yet express IGF-I, (2) cell survival signaling pathways downstream of the IGF-I R may be impaired in the subset of immature Schwann cells that are susceptible to apoptosis, and (3) axon-derived or other Schwann cell-derived factors, in addition to IGF-I, may act in concert with IGF-I to support the survival of these early postnatal Schwann cells *in vivo*. With respect to this latter possibility, recent work has implicated NRG-1 itself in an autocrine and/or paracrine regulation of Schwann cell viability in developing and regenerating nerves (Raabe et al., 1996; Carroll et al., 1997; Rosenbaum et al., 1997).

In summary, the results reported above indicate that IGF-I, like NRG-1, is likely to play an important role in the control of Schwann cell viability during postnatal development and raise the interesting possibility that this regulation may switch in the early neonate from an axon-dependent mechanism to autocrine and/or paracrine signaling. In addition to the established roles for IGF-I in the development and maintenance of the CNS, our work provides further evidence that IGF-I is an important trophic molecule for peripheral nerve modeling and regeneration.

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