Transforming Growth Factor α : A Promoter of Motoneuron Survival of Potential Biological Relevance

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Expression of transforming growth factor α (TGF α), a member of the epidermal growth factor (EGF) family, is a general response of adult murine motoneurons to genetic and experimental lesions, $TGF\alpha$ appearing as an inducer of astrogliosis in these situations. Here we address the possibility that $TGF\alpha$ expression is not specific to pathological situations but may participate to the embryonic development of motoneurons. mRNA of TGF α and its receptor, the EGF receptor (EGFR), were detected by ribonuclease protection assay in the ventral part of the cervical spinal cord from embryonic day 12 (E12) until adult ages. Reverse transcription-PCR amplification of their transcripts from immunopurified E15 motoneurons, associated with in situ double-immunohistological assays, identified embryonic motoneurons as cellular sources of the TGF α -EGFR couple. *In* vitro, TGF α promoted the survival of immunopurified E15 motoneurons in a dose-dependent manner, with a magnitude similar to BDNF neuroprotective effects at equivalent concentrations. In a transgenic mouse expressing a human TGF α transgene under the control of the metallothionein 1 promoter, axotomy of the facial nerve provoked significantly less degeneration in the relevant motor pool of 1-week-old mice than in wild-type animals. No protection was observed in neonates, when the transgene exhibits only weak expression levels in the brainstem. In conclusion, our results point to TGF α as a physiologically relevant candidate for a neurotrophic role on developing motoneurons. Its expression by the embryonic motoneurons, which also synthesize its receptor, suggests that this chemokine is endowed with the capability to promote motoneuron survival in an autocrine–paracrine manner.

Key words: EGF; EGFR; erbB1; development; facial nucleus; spinal cord

Gassmann and Lemke, 1997; Burden, 1998). In contrast, links

between members of the first EGF family group and motoneuro-

nal metabolism have been established only under pathological

conditions up to this day. Motoneuronal TGF α upregulation is

thus the only example of enhanced synthesis of an EGF family

 $TGF\alpha$ is one of the members of the epidermal growth factor (EGF) family, which share an EGF-like domain functioning as their receptor binding domain, and act through the erbB family of tyrosine kinase receptors (Hackel et al., 1999). This family can be subdivided into two groups according to the capability of the EGF-like factors to bind or not the EGF receptor (EGFR or erbB1) (Lee et al., 1995). TGF α , the first member of the EGF family to have been discovered after EGF (Todaro et al., 1980), is one of the EGFR ligands that constitute the first group. Products of the neuregulin 1 gene are part of a second group of molecules that do not bind EGFR but do bind erbB3 and erbB4 and profoundly affect motoneuronal metabolism. They have been shown to be synthesized by embryonic motoneurons and to be necessary for proper motoneuronal maturation and, in particular, for the establishment of adequate interactions between the motoneuron and its glial and muscular environment (for review, see

factor by neuronal cells submitted to an aggression. It was found in wobbler mutant mice undergoing degeneration of their cervical spinal motoneurons, in muscle-deficient murine mutants affected by a degenerative process targeting their lumbar spinal motoneurons, and in hypoglossal motoneurons after axonal crush and cut (Junier et al., 1994, 1998; Lisovoski et al., 1997). The tightly correlated time courses of TGFα motoneuronal production with that of astrogliosis development and EGFR expression by reactive astrocytes designated these glial cells as $TGF\alpha$ primary targets. $TGF\alpha$ capability to induce astrogliosis when overexpressed in the CNS was subsequently demonstrated (Rabchevsky et al., 1998). Although TGF α expression was not observed in intact adult motoneurons, the possibility of an involvement of TGF α in motoneuronal metabolism during development remains to be evaluated. The expression of several members of a given growth factor family by developing motoneurons (Elde et al., 1991; Schecterson and Bothwell, 1992; Krieglstein et al., 1998; Kanda et al., 1999; Garces et al., 2000) and of other EGF family members by immature motoneurons support such a possibility. We thus studied whether TGF α is expressed by embryonic motoneurons and sought for its possible function in the developing spinal cord. We report that embryonic motoneurons are cellular

sources for both the factor and its receptor and show that $TGF\alpha$

has the capability to promote motoneurons survival during em-

bryonic and early postnatal development.

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MATERIALS AND METHODS

Mice and rats were housed in an air-conditioned room with access to water and food *ad libitum*. Sprague Dawley rats (Janvier, Le Genest St. Isle, France) were used for experiments on embryos and motoneuron cultures. Mice were used to assay *in vivo* TGF α effects on motoneuronal survival. Metallothionein 1 (MT1)-hTGF α transgenic mice from the MT-42 line were a generous gift from Dr. G. Merlino of the National Cancer Institute (Bethesda, MD) and bred in our animal room. This transgenic line has been described in detail by Jhappan et al. (1990). It carries a 917 bp human TGF α cDNA under the control of the zinc-inducible mouse metallothionein 1 promoter. These transgenes have been obtained through injection of the construct directly into CD1 one-cell mouse embryos (Jhappan et al., 1990). Their age-matched controls corresponded to the parental CD1 strain (Charles River, Saint-Aubin les Elbeuf, France).

RNA extraction from tissue samples. Total RNA was obtained by the method of Chomczynski and Sacchi (1987) from the ventral part of the cervical spinal cord of embryonic day 12 (E12) to postnatal day 60 (P60) rats and the brainstem of postnatal mice. Pregnant females were anesthetized with chloral hydrate (400 mg/kg, i.p.), and the fetuses were retrieved surgically. A longitudinal incision was made dorsally from the neck to the tail of the fetus, and the spinal cord was removed and its ventral part was trimmed out after flattening the spinal cord on its ventral surface. The dorsal columns were discarded, and the ventral halves were collected. The entire length of the spinal cord was dissected out from E12 embryos. At all other embryonic ages and in postnatal animals, only the cervical part of the spinal cord was collected. Postnatal and adult animals received an overdose of pentobarbital (60 mg/kg, i.p.), and their cervical spinal cord was rapidly removed and the ventral part was dissected out as described above. For RNA extraction from the cervical spinal cords, the tissue samples were pooled before the extraction as follows: five animals per tube from E12-E21, four animals per tube from P2-P6, and three animals per tube from P14-P60. For brainstem collection, the mice were decapitated under pentobarbital anesthesia, the brains were rapidly removed, and the brainstem was dissected out under a binocular microscope. An anterior cut was made rostrally to the trapezoid bodies and a posterior cut rostrally to the lateral reticular nucleus. The resulting tissue piece encompassed the entire extent of the facial nuclei, and tissues from two different animals were pooled per tube. All tissues were immediately frozen on dry ice and stored at -80C until further use.

Ribonuclease protection assay. The antisense TGF α cRNA probe used was obtained by in vitro transcription of a 400 bp EcoRI/HindIII cDNA fragment corresponding to the coding region of the rat $TGF\alpha$ cDNA, subcloned into the pGEM3Z vector (Junier et al., 1991). The transcription was performed as described previously (Junier et al., 1991) in the presence of [32P]CTP. The template was linearized with EcoRI to obtain a 430 nucleotide (nt) [32P]CTP-labeled antisense RNA using SP6 RNA polymerase, 30 of the nucleotides corresponding to the vector multiple cloning site. The antisense EGFR RNA probe was derived from a 578 bp Sau3A1 cDNA fragment of the rat EGFR cDNA (Petch et al., 1990), linearized with AvaII, and transcribed with T7 polymerase. Only 160 bp of this cDNA fragment corresponds to the sequence of the extracellular domain of the full-length EGFR mRNA (Junier et al., 1993). The 169 nt antisense cyclophilin probe was obtained from a PstI/XmnI 111 bp fragment of the rat cyclophilin cDNA (Danielsson et al., 1988), subcloned into the pBluescript vector. The template, linearized with EcoRI and transcribed with T7 RNA polymerase, yielded a 169 nt probe, 58 of which corresponding to the vector multiple cloning site. Levels of the mRNA of interest were assayed as described previously (Junier et al., 1991). Briefly, the tissue total RNA (5 µg) was simultaneously hybridized to the antisense TGF α probe (500,000 dpm) or the antisense EGFR probe and to the cyclophilin antisense RNA probe (5000 dpm) to assess for procedural losses. Cyclophilin mRNA is expressed constitutively in brain tissue (Danielsson et al., 1988). The purified RNA-RNA duplexes were electrophoresed through a 7 M urea-5% acrylamide gel, and the gel was exposed to XAR-5 films for either 18 hr (for cyclophilin signal analysis) or 4-8 d (for TGF α and EGFR signal analysis). Autoradiograms were analyzed by laser densitometry using the cyclophilin mRNA signal to standardize TGF\alpha mRNA values. Results are expressed as the ratio of the TGF α mRNA signal to the cyclophilin mRNA signal.

Immunohistochemistry. E17 and E18 embryos were anesthetized on ice before being perfused transcardially with warm heparinized saline followed by 0.1 m PBS, pH 7.4, containing 4% paraformaldehyde. The spinal cord was immediately removed and post-fixed for 4 hr at 4°C in the same fixative. Tissues were frozen after a 24 hr incubation in a 30%

sucrose cryoprotective medium. Cryostat sections (10 µm thickness) were cut in the frontal plane. Immunohistochemistry was performed on sections mounted on gelatinized slides. The sections were incubated overnight at 4°C in the appropriate dilution of the primary antibody in 0.05 M Tris, 154 mM NaCl, 0.02% BSA, and 0.3% Triton X-100, pH 7.5. Immunohistochemical detection was achieved using either the avidinbiotin complex immunoperoxidase technique and the diaminobenzidine chromogen (Vector Laboratories, Burlingame, CA) or the tyramide signal amplification system coupled to the fluorochrome cyanine 3 (Cy3) or fluorescein (FITC; NEN, Paris, France). $TGF\alpha$ -immunoreactive cells were identified with rabbit polyclonal antibody 1296 (1:5000), which recognizes a peptide sequence (amino acids 137–151) contained within the cytoplasmic domain of the $TGF\alpha$ precursor (pro- $TGF\alpha$) (Gentry et al., 1987). The use of an antiserum that recognizes the intracellular portion of the precursor ensured the identification of the cells that synthesize $TGF\alpha$ as opposed to cells susceptible to bind it. Adjacent sections were stained for the EGF receptor using the rabbit polyclonal antibodies 29.1 (1:250; Sigma, St. Quentin Fallavier, France) or sc-03 [1:100; Santa Cruz Biotechnology (Santa Cruz, CA) and Tebu (Le Perray en Yvelines, France)], which recognize a peptide sequence (amino acids 1005-1016) localized within the intracellular C terminal tail of the EGFR. Alternate sections were stained for choline acetyltransferase (ChAT) using the goat polyclonal antibody AB144P [1:500; Chemicon (Temecula, CA) and Euromedex (Mundolsheim, France)]. Control experiments included omission of either the primary antiserum or the secondary antibody. Specificity of the pro-TGF α and EGFR antibodies in the nervous tissues, including preadsorption of the antibodies with their antigens, has been reported previously (Junier et al., 1991, 1994; Lisovoski et al., 1997; Rabchevsky et al., 1998). Immunofluorescence was observed with a fluorescent microscope (Axioplan 2; Zeiss, Le Pecq, France). Images were acquired on a Cool SNAP camera using the Cool SNAP software (Zeiss). For confocal immunofluorescence, slides were observed using a Zeiss LSM 410 confocal microscope. Two lasers were used depending on the fluorochrome. The excitation wavelengths were 543 nm for Cy3 and 488 nm for FITC, and the emission wavelengths were 570 and 510-525 nm for Cy3 and FITC, respectively. Tissue sections were observed with the 63×. 1.4 numerical aperture lens (Plan-Apochromat; Zeiss, Oberkochen, Germany), the pinhole was set at 20, and the resolution on x-y was of 0.2 and 0.7 μ m for z. Images were prepared for printing using Adobe Photoshop software (Adobe Systems, San Jose, CA).

Facial nerve cut. Cuts of the facial nerve were performed as described previously (Coulpier et al., 1996) on 2- and 8-d-old homozygous MT1hTGF α transgenic mice and their age-matched controls corresponding to the parental CD1 strain (Charles River). Animals were anesthetized on ice and subjected to a unilateral transection of the right facial nerve. The small branch that innervates the caudal auricular muscle and corresponds to motoneurons in the ventromedial part of the facial nucleus was sparred. The success of the axotomy was verified by the sagging of the right facial muscles before the animals were killed. Seven days after the operation, mice were killed and perfused with 4% paraformaldehyde as described above. Sections (30 µm thick) of the brainstem were obtained throughout the entire extent of the facial nucleus and stained with cresyl violet. The nucleoli of the motoneurons were counted every 2 sections over the entire surface of the right and left facial nuclei in each animal. Results are expressed as percentage of total motoneurons counted in the left, unlesioned nucleus (mean ± SEM). Statistical analysis was done using ANOVA, followed by a Fisher's test. Immunofluorescent detection of EGFR in the facial nucleus was performed as described previously. p75 NTR immunodetection was done using the rabbit polyclonal antip75 NTR antibodies (1:500; Promega, Madison, WI) and diaminobenzidine as a chromogen (Vector Laboratories). Counting of the numbers of p75 NTR-immunoreactive motoneurons was performed and analyzed as described above.

Northern blot analysis. Total RNA (10 μ g/lane) was size fractionated in agarose–formaldehyde gels and blotted onto nitrocellulose membranes. The hTGF α transgene and cyclophilin mRNAs were successively identified by hybridization to the following cDNA probes labeled using the random primer method and [32 PJdCTP: 316 bp hTGF α cDNA fragment (Jhappan et al., 1990) and 1 kb rat cyclophilin cDNA (Danielsson et al., 1988). Signals were detected with a PhosphorImager (Molecular Dynamics, Orsay, France) and analyzed with the ImageQuant software (Molecular Dynamics) using the cyclophilin mRNA signal to standardize hTGF α mRNA values (Rabchevsky et al., 1998).

Motoneuron cultures. Cell suspensions of fetal motoneurons were pre-

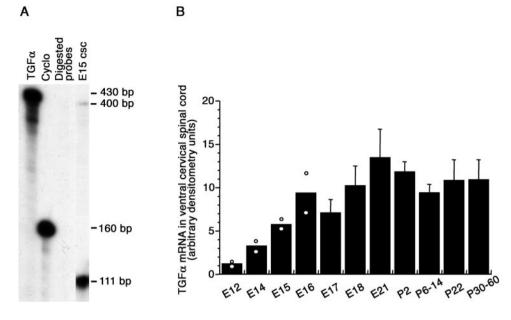


Figure 1. TGFα mRNA levels in the ventral horn of the cervical spinal cord. A, Autoradiogram of TGFα ribonuclease protection assay (4 d exposure, 5 μ g/lane total RNA). $TGF\alpha$, Undigested TGFα cRNA probe; Cyclo, undigested cyclophilin cRNA probe; Digested probes, digested TGFα and cyclophilin cRNA probes; E15 csc, 5 μ g of total RNA from the ventral horn of an E15 rat cervical spinal cord. B, Densitometric analysis of TGFα mRNA signals. Each point represents five (E12–E21), four (P2–P6), and three (P14–P60) animals. Mean \pm SD: n = 3-4.

pared from 15-d-old rat fetuses as described previously (Camu and Henderson, 1992). Motoneurons were isolated and purified using the immunopanning method. The cells were first separated on a metrizamide gradient that segregates large cells. The motoneurons were then isolated using 192 IgG attached previously to a substrate coated with anti-mouse antibodies. Determination of the neuronal enrichment of the cell suspension was performed on aliquots of the cell suspension smeared on a gelatin-coated glass slide. Immunohistochemical detection of the neuronal-specific microtubule-associated protein-2 (MAP-2) and of the motoneuronal-specific transcription factor islet-1 indicated a 92 and 96% enrichment in neurons, respectively, whereas only 2% of the cell suspension was immunolabeled with an antibody recognizing the astrocytespecific marker GFAP. Two thousand five hundred purified motoneurons were seeded in 100 µl of serum-free L15 medium into 96-well plates coated previously with merosin (2 µg/ml; Life Technologies, Cergy Pontoise, France) as described previously (Hanson et al., 1998). One hour after plating, 100 µl of L15 medium alone or containing rat synthetic TGFα (0.1–50 ng/ml; Sigma) or recombinant human brainderived neurotrophic factor (BDNF) (10 ng/ml; Chemicon and Euromedex) was added to each well. Cultures were maintained at 37°C under a 5% CO₂ atmosphere for 3 d. At that time, 88 and 87% of the cells corresponded to motoneurons, as assayed with islet-1/2 and ChAT immunoreactivity, respectively. The vast majority of the remaining cells displayed a neuronal morphology. The cells were then fixed with 4% paraformaldehyde and 0.1% glutaraldehyde in PBS. Immunolabeling with goat anti-ChAT antibody (1:200) was performed as described above using an initial 30 min blocking step in PBS containing 0.2% Triton X-100 and 5% normal rabbit serum and diaminobenzidine as a chromogen. The number of motoneurons was determined over the entire culture dish area. Motoneurons had to exhibit ChAT immunoreactivity and at least three neurites with a length at least twice as long as the soma diameter to be included in the counts. Results are expressed as mean ± SD motoneurons of three independent experiments. Statistical analysis were done using ANOVA, followed by a Fisher's test.

Reverse transcription-PCR amplification. Rat embryonic (E15) motoneurons were purified as described above, pelleted, and immediately frozen on dry ice. RNA was extracted from 10^6 cells using Trizol following the instructions of the manufacturer (Life Technologies). After DNase digestion, 200 ng of total RNA was reverse transcribed using oligo-dT as a primer and the Omniscript reverse transcriptase following the instructions of the manufacturer (Qiagen, Courtaboeuf, France). Substitution of RNA with H_2O served as control of the reverse transcription (RT). PCR amplification was performed using the Taq polymerase from Life Technologies and the following primers: $TGF\alpha$ upper primer, TGG AGA ACA GCA CGT CC; $TGF\alpha$ lower primer, TGG AGA CAC CAG AGT GTT CTG TG; EGFR upper primer, TGG AGA CAC CAG AGT GAT GTG TGG AG; and EGFR lower primer, GAG CTG CAG CGC TGG GGG TCT CGG GCC AT. The amplified cDNAs corresponded to nt 212–575 of the rat $TGF\alpha$ cDNA and to nt 2863–3116 of the rat EGFR

cDNA. PCR was done using a hot-start procedure and the following amplification program: 15 sec at 94°C, 1 min at 55°C, and 2 min at 72°C, 35 cycles. PCR products were purified on a 2% agarose gel containing ethidium bromide before being transferred onto a Nytran membrane (Amersham Pharmacia Biotech, Orsay, France). The membranes were prehybridized for 4 hr and hybridized overnight at 37°C with either a [32 P]dCTP-labeled TGF α or [32 P]dCTP-labeled EGFR cDNA probe (10 cpm/ml) corresponding to nt 1–400 and nt 2863–3116 of the TGF α and EGFR rat cDNAs, respectively. The prehybridization and hybridization solutions were composed of 50% formamide, 5× SSC, 25 mM Na₂HPO4, pH 6.8, 1× Denhardt's solution, 5% dextran sulfate, 1% SDS, and 20 μ g/ml denaturated ssDNA. The membranes were washed twice for 15 min in 1× SSC and 0.1% SDS at room temperature and twice in 0.25× SSC and 0.1% SDS at 37C for 15 min before be exposed to BIOMAX-MS Kodak films (Eastman Kodak, Rochester, NY).

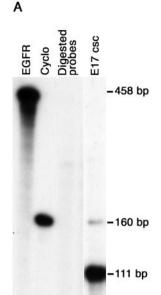
RESULTS

Developmental expression of TGF α and EGFR mRNA in the ventral horn of the rat cervical spinal cord

TGF α mRNA levels were evaluated by ribonuclease protection assay at different developmental stages in the ventral part of the rat cervical spinal cord, in which motoneurons are located. Ribonuclease protection assay of TGF α and cyclophilin mRNAs yielded a 400 and 111 bp protected band, respectively (Fig. 1A). TGF α mRNA signal was detected at 12 d of gestation (E12), the earliest embryonic age examined. TGF α mRNA levels increased progressively throughout the gestation to reach adult levels at birth (Fig. 1B). Ribonuclease protection of the EGFR and cyclophilin mRNA yielded a 160 and 111 bp protected band, respectively (Fig. 2A). Like TGF α mRNA, EGFR mRNA was detected in the spinal cord of E12 embryos. Its levels increased at subsequent embryonic ages and exhibited little variations up to the second week of postnatal life, before decreasing at 3 postnatal weeks (Fig. 2B).

RT-PCR amplification of TGF α and EGFR transcripts from E15 immunopurified motoneurons

To determine whether motoneurons could be one of the cellular sources of $TGF\alpha$ and/or EGFR mRNA detected by ribonuclease protection assay in the ventral spinal cord, RT-PCR amplification of $TGF\alpha$ and EGFR transcripts was performed on total RNA extracted from immunopurified E15 motoneurons. Immunohistochemical detection of the neuronal-specific protein MAP-2 and



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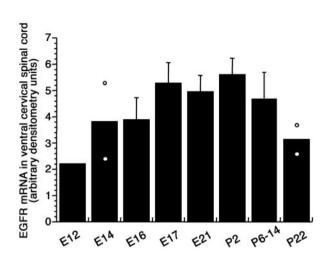


Figure 2. EGFR mRNA levels in the ventral horn of the cervical spinal cord. A, Autoradiogram of EGFR ribonuclease protection assay (8 d exposure, 5 μ g/lane total RNA). EGFR, Undigested EGFR cRNA probe; Cyclo, undigested cyclophilin cRNA probe; Digested probes, digested EGFR and cyclophilin cRNA probes; E17 csc, 5 μ g of total RNA from the ventral horn of a E17 rat cervical spinal cord. B, Densitometric analysis of EGFR mRNA signals. Each point represents five (E12–E21), four (P2–P6), and three (P14–P22) animals. Mean \pm SD; n=3-5.

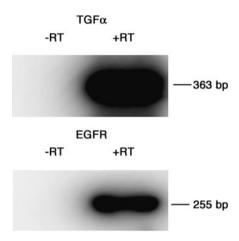


Figure 3. Southern blot hybridization of the RT-PCR amplification of the TGF α and EGFR transcripts from E15 purified motoneuron total RNA. The hybridization probes corresponded to the rat TGF α or the EGFR cDNA probes. -RT, The reverse transcriptase solution was omitted from the PCR reaction. +RT, PCR reaction in the presence of the reverse transcriptase solution.

of the motoneuronal-specific transcription factor islet-1 indicated a 92 and 96% enrichment in neurons, respectively, whereas only 2% of the cell suspension was immunolabeled with an antibody recognizing the astrocyte-specific marker GFAP. RT-PCR amplification yielded a 363 and 255 bp band corresponding to the size expected for TGF α and EGFR cDNAs, respectively. The identity of the RT-PCR-derived products was verified through Southern blot hybridization with the rat TGF α and EGFR cDNA probes (Fig. 3).

Cellular sources of pro-TGF α and its receptor in the embryonic rat cervical spinal cord

Immunohistochemical procedures were used to verify the presence of $TGF\alpha$ and EGFR proteins in the cervical spinal cord and to identify their cellular sources in 17- and 18-d-old embryos. In the 17-d-old embryo, pro- $TGF\alpha$ -immunoreactive cells were observed in the mediolateral and ventral parts of the spinal cord (Fig. 4A) in which motoneurons, identified by their immunore-

activity for the motoneuron marker ChAT, were observed (Fig. 4G). In addition, some pro-TGF α -immunoreactive cells were gathered along the medial axis of the cervical spinal cord (data not shown). Fluorescent pro-TGF α immunohistochemistry yielded similar results in E18 spinal cord embryos. In the ventral part of the spinal cord, $TGF\alpha$ -immunoreactive cells appeared as large-sized neurons (Fig. 4B, C). In the 17-d-old embryo, EGFRimmunoreactive cells displayed a localization similar to pro-TGF α -immunoreactive cells (Fig. 4D). Fluorescent EGFR immunohistochemistry yielded similar results in E18 spinal cord embryos, EGFR-immunoreactive cells appearing as large-sized neurons in the ventral part of the spinal cord (Fig. 4E,F). Motoneuronal identity of the pro-TGF α and EGFRimmunoreactive cells observed in the ventral horn of the embryonic spinal cord was further confirmed with doubleimmunofluorescent staining using either TGF α and ChAT antibodies or EGFR and ChAT antibodies (Fig. 5). Confocal microscopy showed that neurons immunoreactive for $TGF\alpha$ were also immunoreactive for ChAT (Fig. 5A-C). ChAT immunolabeling appeared diffused throughout the cytoplasm (Fig. 5A), whereas $TGF\alpha$ immunostaining exhibited a punctated aspect (Fig. 5B). Likewise, EGFR-immunoreactive neurons were also ChAT immunoreactive (Fig. 5D-F), with the EGFR signal being preferentially localized at the plasma membrane (Fig. 5E,F).

$\mathsf{TGF}\alpha$ effects on embryonic rat motoneuron survival cultured in serum-free medium

The observation of TGF α and EGFR expressing motoneurons during the period of developmental cell death led us to evaluate the effects of this growth factor on the survival of motoneurons immunopurified from E15 embryonic spinal cord. The percentage of motoneurons in the cellular preparation was assessed with the 4D5 monoclonal antibody, which recognizes the motoneuron-specific transcription factors islet-1/2 (Ericson et al., 1992), and with an anti-ChAT polyclonal antibody. The purification procedure yielded a 88 and 87% enrichment in motoneurons, as assayed with islet-1/2 and ChAT immunoreactivity, respectively. Motoneurons were seeded at low density in serum-free medium to avoid accumulation of motoneuron-derived growth factors and addition of EGF-like factors known to be abundant in serum.

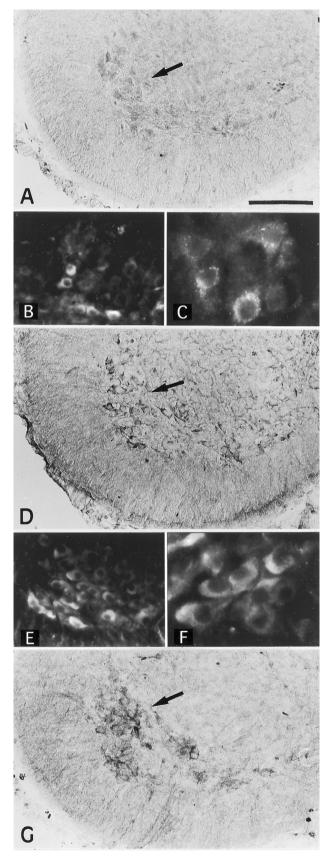


Figure 4. Immunohistochemical localization in the cervical spinal cord of 17- and 18-d-old rat embryos of the TGF α precursor, its receptor EGFR, and ChAT, a motoneuron marker. A, TGF α precursor immunolabeling in 17-d-old cervical spinal cord. The arrow points to some of the labeled cells in the ventral region of the spinal cord. B, C, Immunofluo-

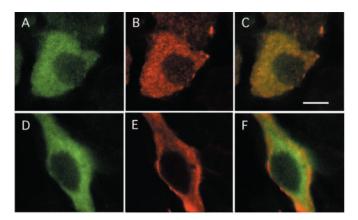


Figure 5. Confocal visualization of ChAT and TGF α precursor (A–C) and of ChAT and EGFR (D–F) immunofluorescent labelings in the ventral part of the cervical rat spinal cord of an 18-d-old embryo. TGF α precursor-immunoreactive neurons (B) are also ChAT immunolabeled (A). C, Overlay of ChAT (green) and TGF α (red) immunofluorescent signals depicted in A and B. Likewise, EGFR-immunoreactive neurons (E) are also ChAT immunoreactive (D). F, Overlay of ChAT (green) and EGFR (red) immunofluorescent signals depicted in D and E. Scale bar, $10~\mu m$.

Motoneuronal survival was assessed after 3 d in culture. At that time, glia-like cells were rare (zero to five cells per well), including in the presence of $TGF\alpha$. Well developed ChAT-immunoreactive cells displaying a large soma and at least three neurites with a length at least twice as long as the largest soma diameter were taken into account (Fig. 6). BDN F, a neurotrophin well known to be trophic for cultured embryonic motoneurons (Henderson et al., 1993), was used as a positive control. A statistically significant positive effect of $TGF\alpha$ was observed on motoneurons survival. $TGF\alpha$ promoted motoneurons survival within a concentration range of 0.1–50 ng/ml, the maximal effect being observed at 50 ng/ml, the highest concentration tested. At 10 ng/ml, $TGF\alpha$ effect on motoneuron survival was similar to the survival-promoting effect of 10 ng/ml BDNF (Fig. 7).

Axotomy-induced motoneuronal death in 1-week-old MT1-hTGFlpha mice

The ability of $TGF\alpha$ to promote motoneuron survival *in vivo* was explored using the paradigm of facial nerve cut, which triggers motoneuronal death in young postnatal mice. Mice overexpressing an hTGF α transgene in the CNS were used. In this mouse strain, the transgene expression is under the control of the metallothionein 1 promoter. Because metallothionein 1 is expressed at barely detectable levels in embryonic and early postnatal ro-

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rescent staining of the TGF α precursor in the ventralmost part of the cervical spinal cord gray matter of 18-d-old rat embryos at low (B) and high (C) magnifications. D, EGFR immunolabeling in 17-d-old cervical spinal cord. The arrow points to some of the EGFR-immunoreactive cells localized in the ventral portion of the spinal cord. E, F, Immunofluorescent staining in the ventralmost part of the cervical spinal cord gray matter of 18-d-old rat embryos of the EGFR at low (E) and high (F) magnifications. Note that EGFR-immunoreactive neurons appear more numerous in E than TGF α precursor-immunoreactive neurons in F. G, ChAT immunohistochemical labeling. Immunolabeled neurons are localized in the mediolateral and ventral regions of the spinal cord in 17-d-old cervical spinal cord. The arrow points to some of the labeled cells in the ventral region of the spinal cord. In B, C, E, and F, the original immunofluorescent signal appeared red (cyanine 3 fluorochrome). Scale bar: A, D, G, 150 μ m; B, E, 140 μ m; C, F, 56 μ m.

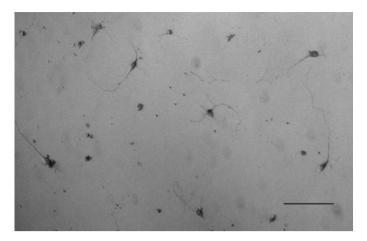


Figure 6. ChAT-immunoreactive neurons after 3 d in culture in the presence of 1 ng/ml TGFα. Scale bar, 100 μ m.

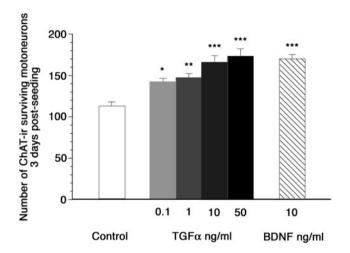
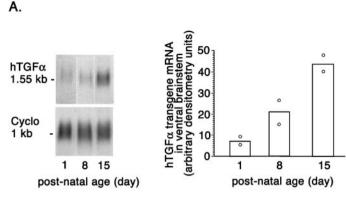


Figure 7. Effect of TGFα on spinal motoneuron survival. Number of motoneurons was determined after 3 d in culture in serum-free medium (white bar), in the presence of 0.1–50 ng/ml TGFα (gray and black bars), or in the presence of 10 ng/ml BDNF (hatched bar). Cells were taken into account when exhibiting ChAT immunoreactivity and displaying a large soma and at least three neurites with a length at least twice as long as the largest soma diameter. The number of motoneurons was determined over the entire culture dish area. Means \pm SD; n=7–10; three independent experiments. *p<0.0005; ***p<0.0005; ***p<0.0001; ANOVA followed by a Fisher's test.

dents brains (Waalkes and Klaassen, 1984; Choudhuri et al., 1996; Penkowa et al., 1999), hTGF α mRNA levels were first evaluated by Northern blot assays at 1, 8, and 15 postnatal days in the mid-brainstem, in which facial nuclei are localized (Fig. 8A, left). hTGFα mRNA levels were low at birth and increased through postnatal development to reach six times higher levels at postnatal day 15, the oldest age examined (Fig. 8A, right). Vulnerability of facial motoneurons to axotomy-induced death has been reported to decrease with aging. It has been shown that axotomy induces death of 90% of the motoneurons in 2-d-old mice, of 50% in 1-week-old mice, and of 10% in 2-week-old mice (Kuzis et al., 1999). Facial nerve cuts were therefore performed on 8-d-old animals, a time at which half of the facial motoneurons is sensitive to axotomy-induced death and significant amounts of hTGF α mRNA are detected in the MT1-hTGF α transgenic mice brainstem. In addition, facial motoneuron response to axotomy was monitored in 2-d-old animals, a time at which the $TGF\alpha$



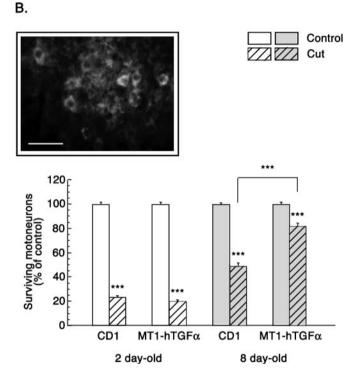


Figure 8. Effect of TGF α on axotomy-induced motoneuronal death in the facial nucleus of controls and MT1-hTGF α transgenic mice. A, Human TGF α transgene expression in the ventral brainstem of 1-, 8-, and 15-d-old MT1-hTGF α mice. Left, Northern blot analysis of hTGF α mRNA levels. Cyclophilin mRNA served as an internal standard. Right, Densitometric analysis of hTGF α mRNA signals. Note the increase in hTGFα mRNA levels as the animals age. Each point represents two animals. B, Quantification of facial motoneurons on the unlesioned and lesioned facial nuclei 7 d after lesion in control (CD1) and MT1-hTGF α transgenic mice operated at either 2 or 8 d of age. Results are represented as percentages of surviving motoneurons in the lesioned facial nucleus 7 d after axotomy relative to motoneuron numbers in the contralateral, unlesioned facial nucleus. Unhatched bars, Unlesioned nucleus. Hatched bars, Lesioned nucleus. Mean \pm SEM; n = 4; ANOVA followed by the Fisher's test; ***p < 0.001. *Inset*, Immunohistochemical localization of EGFR in the facial nucleus of 1-week-old MT1-hTGF α transgenic mice. Scale bar, $75 \mu m$.

transgene is expressed at low levels. The examination of facial nuclei 7 d after the transection of the facial nerve at 2 d of age revealed a dramatic reduction in the number of neurons in control mice, and, as expected, similar observations were made in transgenic mice. Eighty percent of the facial motoneurons were lost

(mean \pm SEM of motoneurons corresponding to 1377 \pm 28 in intact nucleus vs 325 \pm 19 in axotomized nucleus in control mice and 1367 ± 25 vs 273 ± 19 in transgenic mice; n = 4) (Fig. 8B). Most surviving motoneurons were observed in the ventromedial part of the nucleus, which corresponds to the nonsectioned caudal auricular branch of the facial nerve. In 8-d-old control mice, axotomy also triggered facial motoneuron death 7 d after lesion but to a lesser extent than in 2-d-old animals. Fifty percent of the motoneurons were lost (mean ± SEM of motoneurons corresponding to 1359 \pm 22 in intact nucleus vs 666 \pm 39 in axotomized nucleus in control mice; n = 4) (Fig. 8B). In 8-d-old transgenic mice, surviving facial motoneurons observed 7 d after axotomy appeared more numerous than in control mice. Only 18% of the motoneurons were lost (mean \pm SEM of motoneurons corresponding to 1193 \pm 21 in intact nucleus vs 978 \pm 33 in axotomized nucleus; n = 4) (Fig. 8B). Observation of numerous EGFR-immunoreactive motoneurons in the facial nucleus of these mice indicated that these cells are endowed with the capability to be directly affected by $TGF\alpha$ (Fig. 8B, inset). Such a low percentage of motoneurons death observed after the axotomy of the facial nerve of 8-d-old transgenic mice is, in control mice, attained only when the axotomy is performed after 2 weeks of postnatal life (Kuzis et al., 1999). The possibility, nevertheless, remained that motoneuron protection observed in the 8-d-old transgenic mice resulted from an accelerated maturation of the facial motoneurons. To address this possibility, motoneuronal expression of the low-affinity neurotrophin receptor $p75^{\,\mathrm{NTR}}$ was used as an index of motoneuron maturation in the intact facial nucleus. p75 NTR motoneuronal expression is indeed known to decrease progressively during the first 2 weeks of postnatal life before disappearing in the adults (Eckenstein, 1988; Yan and Johnson, 1988; Ernfors et al., 1989). The numbers of p75 NTRimmunoreactive motoneurons in 2-week-old transgenes and controls were similar and represented 2.4-3.4% of the population of facial motoneurons (control mice 32 ± 9.5 vs transgenic mice $40.7 \pm 6.6 \,\mathrm{p}75^{\mathrm{NTR}}$ -immunoreactive motoneurons; mean \pm SEM; n = 3). In 1-week-old transgenes, p75 NTR-immunoreactive facial motoneurons were 4.4 times more numerous than in 2-week-old control animals but represented only 10.4% of the population of facial motoneurons instead of 16.2% in 1-week-old controls (transgenic mice 142.3 ± 5.7 vs control mice 223 ± 14.5 ; mean \pm SEM; n = 3; p = 0.0001).

DISCUSSION

Studies over the past 8 years have revealed that developing motoneurons synthesize at least one class of the EGF family of growth factors, the neuregulin1 gene products. Our results show that, in rat embryonic motoneurons, this family extends to $TGF\alpha$. They also indicate that this expression might have functional consequences by providing evidences *in vitro* and *in vivo* of a role for $TGF\alpha$ as a promoter of motoneuron survival.

A functional autocrine–paracrine loop indicated by TGFlpha and EGFR coexpression

Localization of $TGF\alpha$ in embryonic nervous tissues is only known in the forebrain (Kornblum et al., 1997). Our results show that $TGF\alpha$ and EGFR transcripts can be detected in the rat spinal cord from 12 d postcoitum until adult ages. Most cells immunoreactive for the $TGF\alpha$ precursor and its receptor were localized in the ventral and mediolateral part of the cervical spinal cord. This distribution matches that observed for motoneurons identified by their ChAT immunoreactivity and their large soma size (Altman

and Bayer, 1985; Phelps et al., 1990). Double-immunohistological assays showing that pro-TGF α - and EGFR-immunoreactive cells are indeed ChAT immunoreactive, associated with RT-PCR amplification of the TGF α and EGFR transcripts from suspensions of purified motoneurons containing at most 2% of GFAP-immunoreactive cells, confirmed motoneurons as cellular sources of the TGF α -EGFR couple.

TGF α promotes motoneuron survival

TGF α and EGFR coexpression in embryonic motoneurons during the period of developmental cell death raised the possibility that TGFα affects regulation of neuronal survival through an autocrine-paracrine mode of action. The increase in motoneuron survival observed in culture in the presence of $TGF\alpha$ indicates that this factor has the capability to promote survival of embryonic motoneurons, an observation similar to that reported by Hanson et al. (1998) with a single TGF α concentration of 50 ng/ml. TGF α effect is dose dependent and, at the highest concentrations, similar in magnitude to the survival stimulatory effect of equivalent concentrations of BDNF in our culture conditions. EGFR expression by motoneurons and the lack of significant glial contamination in the cultures (zero to five cells per culture well, including in the presence of $TGF\alpha$) indicate that astrocytes do not mediate $TGF\alpha$ effects on motoneurons. This singles out motoneurons among the neuronal populations known to be responsive to $TGF\alpha$ neurotrophic effects, which, up to this day, have been shown to occur in most cases through the astrocytes intermediacy (for review, see Junier, 2000).

TGF α is not the only growth factor endowed with the capability to act in an autocrine-paracrine manner on motoneurons. The growth factors $TGF\beta_2$ and $TGF\beta_3$, as well as neurotrophin-3, are synthesized by embryonic motoneurons together with their receptor and promote embryonic motoneuronal survival (Flanders et al., 1991; Henderson et al., 1993; Gouin et al., 1996; Krieglstein et al., 1998). To examine whether $TGF\alpha$ promoting-survival action on motoneurons extends to in vivo situations, we used the paradigm of facial nerve cut in early postnatal mice. This model is widely used to test the ability of growth factors to promote motoneuron survival in situ (Houenou et al., 1994). In addition, axotomized motoneurons die in an apoptotic manner, as during physiological developmental death (de Bilbao et al., 2000; Vanderluit et al., 2000), and are protected by the same molecules (Sendtner et al., 1990, 1992; Yan et al., 1992, 1995; Henderson et al., 1993, 1994; Koliatsos et al., 1993; Dubois-Dauphin et al., 1994; Kato and Lindsay, 1994; Martinou et al., 1994; Farlie et al., 1995; Vejsada et al., 1995; Coulpier et al., 1996). In the MT1hTGF α mice, the transgene expression is driven by the metallothionein 1 promoter, and metallothionein 1 synthesis appears only during late postnatal development (Waalkes and Klaassen, 1984; Choudhuri et al., 1996; Penkowa et al., 1999). Accordingly, low amounts of the transgene were expressed in the midbrainstem of 2-d-old transgenic mice. In contrast, facial motoneurons are protected from axotomy-induced cell death in 1-weekold mice when expression of human TGF α transgene increases in the brainstem. Approximately 60–65% of the facial motoneurons disappearing in control mice are rescued from death in MT1hTGF α mice. The possibility exists that TGF α overexpression leads to an accelerated maturation of the motoneurons and, hence, to an enhanced resistance to the deleterious effects of axotomy. Although a reduced amount of p75 NTR-immunoreactive motoneurons was observed in the transgenic mice facial nucleus (10% of the facial motoneuron population in 1-week-old transgenic mice vs 16% in age-matched controls), it is not sufficient to account for the increase in motoneuron survival observed in the transgenic mice. Hence, if accelerated motoneuronal maturation indeed occurs in these transgenic mice, it accounts only partly for the TGF α neuroprotective effects obtained in this in vivo model. In addition, detection of numerous EGFRimmunoreactive facial motoneurons in 1-week-old intact facial nucleus suggests that facial motoneurons can be directly affected by TGFα. The possibility remains, however, that other EGFRexpressing cells, not clearly detectable in our assay, also mediate part of the $TGF\alpha$ neuroprotective effects. Altogether, the present results indicate that TGFα, like other factors trophic for motoneurons (Oppenheim, 1996; Sendtner et al., 1996), is available in situ in the developing spinal cord, its receptor is expressed by embryonic motoneurons, and its survival-promoting effects can be disclosed in both in vitro and in vivo assays. These data give credence to the existence of a biological relevant role of $TGF\alpha$ in motoneuronal development.

However, the extent of this biological relevance remains to be evaluated. Resolution of this issue depends in part on examination of mice lacking TGF α or its receptor. The EGFR null mice exhibit embryonic or early postnatal lethality depending on the genetic background, essentially as a result of placental or lung defects (Miettinen et al., 1995; Sibilia and Wagner, 1995; Threadgill et al., 1995). They display severe impairment of brain development (Threadgill et al., 1995; Kornblum et al., 1998; Sibilia et al., 1998), but alterations in motor nuclei have not been explored. In contrast, $TGF\alpha$ null mice develop normally (Luetteke et al., 1993; Mann et al., 1993), and the only CNS defect reported to date corresponds to the lack of a nigral dopaminergic subpopulation (Blum, 1998). To our knowledge, no exploration of the $TGF\alpha$ -/- motoneuronal systems has been undertaken. The absence of major CNS defect, at least at the clinical level, may stem from the existence of functional redundancies between the different EGFR ligands. Indeed, at least two other EGFR ligands, heparin-binding-EGF and amphiregulin, have been detected in the spinal cord (Hayase et al., 1998) (MPJ, unpublished results). It may also result from the existence of subtle interventions for $TGF\alpha$, the factor mediating in part the action of other promoters of motoneuron survival. Its absence would thus lead only to discreet functional modifications. Development of $TGF\alpha$ blocking antibodies should allow to definitely evaluate whether $TGF\alpha$ is physiologically involved in the control of motoneuron development.

$\mathsf{TGF}\alpha$: another member of the EGF family to be expressed by embryonic motoneurons

Our results show that embryonic motoneurons express another EGF family member in addition to neuregulin1. However, marked differences exist between these factors. Neuregulin1 ensures the adequate interactions between motoneurons and their environment. It affects Schwann cell metabolism at all stages of their development and the establishment of the neuromuscular junction, both Schwann and muscle cells expressing neuregulin1 receptors (Burden, 1998; Mirsky and Jessen, 1999). In contrast, our results indicate that the motoneurons themselves constitute $TGF\alpha$ targets in the embryo and that the factor fulfils a trophic role toward the motoneurons. An effect of $TGF\alpha$ on Schwann and muscle cells cannot, however, be excluded. Its receptor has been described in these cells that, at least *in vitro*, may be sensitive to its mitogenic and/or differentiative actions (Lim and Hauschka, 1984; Olwin and Hauschka, 1988; Toma et al., 1992; Sastry et al.,

1996; Halse et al., 1999). This, associated with the capability of the different EGF family members to interact through heterodimerizations of their respective receptors (Beerli and Hynes, 1996; Graus-Porta et al., 1997; Lenferink et al., 1998), allows interactions between neuregulin1 and $TGF\alpha$ to be envisioned. Another difference stems from the switch in $TGF\alpha$ cellular sources during development. Neuregulin1 products are synthesized by motoneurons throughout life (Chen et al., 1994) and, in the adult, participate to the maintenance of the neuromuscular junction (Sandrock et al., 1997). In contrast, TGFα source changes from motoneurons to white matter astrocytes and few small-sized neurons in mature animals. At this stage, $TGF\alpha$ motoneuronal expression may be observed only under pathological situations, and reactive astrocytes and Schwann cells constitute then its potential targets (Toma et al., 1992; Junier et al., 1994; Lisovoski et al., 1997). Inversely to $TGF\alpha$, motoneuronal neuregulin1 expression declines in adults after axotomy (Bermingham-McDonogh et al., 1997), suggesting that TGFα could compensate in part neuregulin1 loss.

Conclusion

Our results allow $TGF\alpha$ to be added to the list of growth factors capable of acting on developing motoneurons. It is now acknowledged that factors trophic for motoneurons may derive not only from their muscle targets but also from local sources (Oppenheim, 1996; deLapeyrière and Henderson, 1997). In this context, $TGF\alpha$ appears as a putative autocrine–paracrine growth factor for motoneurons.

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