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## Enhanced expression of transforming growth factor $\beta 1$ in the rat brain after a localized cerebral injury

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### Abstract

It is becoming clear that transforming growth factor  $\beta$  (TGF $\beta$ ) may be a key factor regulating inflammatory and tissue specific wound responses. Because the formation of a glial-collagen scar at CNS lesion sites is thought to contribute to the pathology associated with penetrating CNS injuries, and because in the periphery TGF $\beta 1$  stimulates fibroblast deposition of scar tissue, we used in situ hybridization and immunohistochemistry to investigate the effect of a defined cerebral lesion on the local expression of TGF $\beta 1$ . Induction of TGF $\beta 1$  mRNA and protein is relatively diffuse in the neuropile around the margins of the lesion at 1, 2 and 3 days, but becomes localized to the region of the glial scar at 7 and 14 days. The signal intensity for TGF $\beta 1$  mRNA and protein is maximal between 2 and 3 days and decreases between 7 and 14 days after lesion. The predominant cell types in the neuropile localizing TGF $\beta 1$  mRNA and protein have the morphological characteristics of astrocytes, although macrophages are also detected. An induction of TGF $\beta 1$  mRNA was also observed in endothelial cells of the meninges, hippocampal fissure and choroid plexus, at 2 and 3 days. However, this is dramatically reduced by 7 days and has disappeared by 14 days. These results suggest a role for TGF $\beta 1$ , not only in inflammation, but also in the tissue-specific glial scar formation that occurs in the CNS. Furthermore, they suggest a potential therapeutic use of TGF $\beta 1$  antagonists in the CNS to help limit the pathogenesis associated with matrix deposition in the wound.

### Keywords

Transforming growth factor  $\beta 1$ ; mRNA; Injury; Central nervous system; Glial scar; Astrocyte

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## INTRODUCTION

Injury to the adult mammalian central nervous system (CNS) generates a cascade of cellular events including recruitment of inflammatory cells into the lesion, transient abortive regeneration of damaged neurones, astrocyte proliferation and angiogenesis. The blood-brain barrier is reformed by the tight association of astrocyte end-feet at the wound margins, to make a glia limitans externa which encloses a core of matrix molecules laid down by fibroblasts recruited from the meninges<sup>24</sup>. The glial-mesodermal scar provides a physical barrier that any regenerating neurones have to penetrate in order to re-establish functional connections with their target organs. The limited ability of the adult CNS to recover function after injury relates to unsuccessful nerve regeneration and deposition of a dense, permanent fibrotic scar<sup>5,21</sup>.

Although the mechanisms responsible for post-traumatic CNS degeneration are not defined, interacting growth factors are thought to play crucial roles in determining the outcome of CNS injury. Indeed, there have been a number of recent reports suggesting the involvement of several growth factors, including basic fibroblast growth factor (FGF)<sup>11,22</sup>, acidic FGF<sup>20</sup>, transforming growth factor  $\beta$ 1 (TGF $\beta$ 1)<sup>17,28</sup>, and interleukins 1 and 2<sup>13,29</sup>. Some of these factors may play important neurotrophic roles<sup>1,6,30,43,49</sup> but their role in scar formation has not been studied. While it remains unclear which components of the trophic and cellular responses are helpful and which harmful to functional recovery of the CNS, the agents which promote scar formation presumably impede neuronal regeneration and the establishment of new synaptic connections<sup>5,21,35</sup>.

In addition to the reported effects of TGF $\beta$ 1 as an inflammatory mediator<sup>51</sup>, this factor also influences endothelial cells<sup>4,9,31</sup>, neurones<sup>6</sup> and glia<sup>7,18,39,41</sup>. TGF $\beta$ 1 is a potent desmoplastic agent in peripheral tissues<sup>37</sup>. It is an important regulator of extracellular matrix deposition by fibroblasts, increasing collagen, fibronectin and proteoglycan expression<sup>14,34</sup> and stimulating collagen-matrix contraction<sup>27</sup>. It also increases integrin expression<sup>14</sup>, decreases the synthesis of proteases which degrade extracellular matrix components such as collagenase<sup>8</sup> and transin<sup>23</sup> and increases the expression of protease inhibitors<sup>8</sup>. Thus, many of the activities of TGF $\beta$ 1 are directed towards the production and deposition of fibrous scar tissue within peripheral wounds.

Normally, there are low levels of TGF $\beta$ 1 in the CNS<sup>26,46,50</sup>, but since TGF $\beta$ 1 is released from platelets at the site of injured blood vessels<sup>2</sup> and is also expressed by macrophages<sup>15</sup>, it is likely that breakage of the blood-brain barrier will allow entry of TGF $\beta$ 1 protein into the CNS. We reasoned that after a penetrating CNS lesion, in addition to its action as a mediator of the inflammatory response, TGF $\beta$ 1 may influence gliosis and deposition of the extracellular matrix of the scar and predicted that this factor may be expressed locally in neural tissue following a penetrating wound. Of direct relevance, TGF $\beta$ 1 mRNA has been found to increase in the ipsilateral hippocampus following entorhinal cortex lesions<sup>28</sup> and after ischaemic injury<sup>16</sup> supporting our hypothesis that locally-produced TGF $\beta$ 1 may regulate some of the tissue-specific responses of microvessels, fibroblasts, microglial/macrophages and neurones after injury. However, precisely which cells in the CNS synthesize TGF $\beta$ 1 after injury and how the response differs depending on the type of injury

have not been established. Therefore, we used immunohistochemistry and in situ hybridization in a rat model of penetrating CNS injury to investigate the possibility that TGF $\beta$ 1 mRNA and protein are induced locally at wound sites by injury-responsive cells.

## MATERIALS AND METHODS

### Materials

All reagents not specified were analytical grade from Sigma Chemical Co., St. Louis, MO. Radioisotopes were obtained from Amersham International, Arlington Heights, IL.

### Animals and surgery

Surgical and animal care procedures were carried out with strict adherence to the guidelines set out in the 'NIH guide for the care and use of laboratory animals', National Institutes of Health Publications no. 80–23.

Groups of adult, male Sprague–Dawley rats (200 g) were anaesthetized (i.m.) with a mixture of acepromazine (1.875 mg/kg), ketamine (3.75 mg/kg) and xylazine (1.9 mg/kg). A stereotactically defined, 4-mm deep rostro-caudal knife-wound incision was made vertically into the occipital cortex, corpus callosum and presubiculum<sup>22</sup>. At 1, 2, 3, 7 and 14 days after surgery, animals ( $n = 4$ /group) were put under deep anaesthesia with the same anaesthetic and perfused transcardially with 300 ml of 0.9% (w/v) saline, 250 ml of 4% (w/v) paraformaldehyde (PFA) in 0.1 M acetate buffer, pH 6.5, followed by 500 ml of 4% (w/v) PFA plus 0.05% (w/v) glutaraldehyde in 0.1 M borate buffer, pH 9.5, using the pH shift method<sup>45</sup>.

### Histology

For in situ and immunocytochemical studies, the perfusion-fixed brains were post-fixed overnight at 4°C in 4% (w/v) PFA in 0.1 M borate buffer containing 10% (w/v) sucrose. They were then rapidly frozen on powdered dry ice in Tissue Tek OCT compound (Miles Laboratories, Napeville, IL) and stored at –80°C. Frozen sections (20  $\mu$ m) were mounted on poly-L-lysine coated slides, air-dried and stored at –80°C until use.

### In situ hybridization

The methods used for the cloning of rat TGF $\beta$ 1 are described elsewhere<sup>33</sup>. In situ hybridization of TGF $\beta$ 1 used the *Hind* III-*Xba*I fragment of 0.985 kbp, derived from the major coding region of the rat TGF $\beta$ 1 precursor, which was subcloned into pBluescript SK<sup>+</sup> (Stratagene, San Diego, CA). The antisense strand of the coding sequence was transcribed using T7 polymerase. A [<sup>35</sup>S]UTP-labelled RNA probe encoding the sense strand of 5' non-coding sequence was prepared with T3 RNA polymerase and used for control tissue sections.

Mounted frozen sections of brain, prepared as described in the histology section of Materials and Methods, were digested with 10  $\mu$ g/ml of proteinase K in 0.1 M Tris (pH 8.0) containing 50 mM EDTA at 37°C for 30 min. Sections were rinsed in deionized water followed by incubation in 0.1 M triethanolamine (TEA), pH 8.0, for 3 min. Sections were then acetylated

for 10 min with 0.25% (w/v) acetic anhydride in 0.1 M TEA, rinsed in 2 ×SSC, dehydrated through a graded series of ethanol washes and air-dried under vacuum for 2 h before hybridization.

Hybridizations with labelled TGF $\beta$ 1 antisense or sense probes (1 × 10<sup>6</sup> cpm/ml) were performed at 55°C overnight in 10 mM Tris (pH 8.0) containing 50% (w/v) formamide, 0.3 M NaCl, 1 mM EDTA, 10 mM dithiothreitol (DTT), 1 × Denhardt's solution and 10% (w/v) dextran sulphate. After hybridization, sections were rinsed for 1 h in 4 ×SSC and treated with 25  $\mu$ g/ml ribonuclease A in 10 mM Tris (pH 8.0) containing 0.5 M NaCl and 1 mM EDTA at 37°C for 30 min. This was followed by increasing high stringency washes of SSC containing 1 mM DTT, finishing with 0.1 ×SSC at 65°C for 30 min.

Slides were then dehydrated through a graded series of ethanol, dried under vacuum and then exposed to  $\beta$ max hyperfilm (Amersham) for 5 days to examine gross changes in mRNA. For microscopic analysis, slides were exposed to Kodak NTB-2 liquid autoradiographic emulsion for 3 weeks at 4°C, processed with Kodak D19 developer, rinsed and fixed with Kodak rapid fixer. The slides were rinsed for 30 min in tap water, counterstained with Harris' haematoxylin and examined by dark field and bright field microscopy.

### Immunoperoxidase staining

The primary antiserum raised against TGF $\beta$ 1 has been previously described and characterized<sup>10</sup>. It is an IgG fraction of a rabbit polyclonal, raised against amino acids 1-30 of rat TGF $\beta$ 1, which was purified by passage over a Protein A-Sepharose column.

Immunoperoxidase staining for TGF $\beta$ 1 in 20- $\mu$ m frozen sections of brain used the ABC Vectastain Elite kit (Vector Laboratories, Burlingame, CA). The tissue sections were washed in PBS and the endogenous peroxidase quenched by incubating with 0.3% (v/v) hydrogen peroxide in PBS for 30 min. The sections were rinsed in PBS and incubated in 1.5% (v/v) goat serum, diluted in PBS containing 0.3% (v/v) Triton X-100, for 30 min to reduce non-specific staining.

After a 24-h incubation at 4°C with the protein-A purified primary antibody (0.015 mg/ml), diluted in PBS supplemented with 0.3% (v/v) Triton X and 1% (v/v) BSA, the sections were treated with a 1:200 dilution of biotinylated goat anti-rabbit IgG (Vector) for 1 h. This was followed by a 30-min incubation with a biotin-avidin-peroxidase complex (Vector). Finally, the sections were treated for 5 min with 0.5 mg/ml diaminobenzidine in PBS containing 0.01% (v/v) hydrogen peroxide. All steps were separated by buffer washes consisting of PBS with 0.3% (vol/vol) Triton X-100.

The sections were finally washed in PBS, counterstained with Harris' haematoxylin, dehydrated, cleared and mounted. Sections incubated with anti-TGF $\beta$ 1 antiserum preincubated with recombinant TGF $\beta$ 1 or without primary antisera were used as controls. Sections processed with the control procedures failed to stain.

## RESULTS

### General properties of the lesion

Haematoxylin staining of sections through the lesion site at different days after surgery shows the typical transformation of the wound from an open cavity to a closed glial/mesodermal scar (Fig. 1) in both the rostral (Fig. 1A, B) and caudal (Fig. 1C, D) portions of the lesion. The incision is seen to have penetrated the cortex and the corpus callosum and passed through the ventricle, medial to the hippocampal formation. After 7 days, the migration and proliferation of glial cells has produced a well-defined glia limitans which defines the wound margin, and surrounds a core of mesodermal, fibrous elements. Glial proliferation through the neuropile appears especially extensive in the corpus callosum. The contraction of the scar tissue is particularly evident at 7 days (Fig. 1B).

### TGF $\beta$ 1 mRNA and the margin of the lesion

The TGF $\beta$ 1 probe recognizes the expected major 2.5-kb band when the injured brains are processed for Northern blot hybridization (not shown). The time course of the TGF $\beta$ 1 mRNA response in the rostral region of the wound was assessed by in situ hybridization and is shown in Fig. 2. In this study signal for TGF $\beta$ 1 mRNA is very low or negligible in sections through similar regions of the unlesioned adult rat brain (not shown). A clear increase in TGF $\beta$ 1 mRNA signal is observed in sections from lesioned animals at 1, 2, 3, 7 and 14 days post-lesion (Fig. 2A<sub>1</sub>–E<sub>1</sub>). This increase is apparent when the level of signal in sections from lesioned animals is compared to the signal observed in sections taken through the same region of unlesioned brain and processed simultaneously and identically (not shown), or in alternate lesion sections hybridized with the TGF  $\beta$ 1 sense strand probe (Fig. 2A<sub>2</sub>–E<sub>2</sub>).

By in situ hybridization, the TGF $\beta$ 1 mRNA signal is most intense at 2 and 3 days post-lesion (Fig. 2B<sub>1</sub>–C<sub>1</sub>). Interestingly, while the signal is relatively diffuse at 1, 2 and 3 days, it becomes focal when the glia limitans is formed after 7 days. The meninges and presumptive astrocytes (by morphological criteria) adjacent to the site of incision, show intensive TGF $\beta$ 1 mRNA hybridization at 1, 2 and 3 days but the signal has diminished at 7 and 14 days.

The response is very similar in the caudal region of the wound and is presented in Fig. 3. A marked induction of TGF $\beta$ 1 mRNA in the choroid plexus is observed in these sections at 1, 2, and 3 days, and has diminished by 7 days (Fig. 3A<sub>1</sub>–E<sub>1</sub>). When adjacent sections are hybridized with sense probe, the signal is negligible (Fig. 3A<sub>2</sub>–E<sub>2</sub>). As in the rostral part of the wound, the signal for mRNA has localized to the glia limitans at 7 days. Signal in all caudal regions of the wound has diminished by 14 days post-lesion.

The time course of the TGF $\beta$ 1 mRNA response in cells on the margin of the wound is shown with high magnification bright field micrographs in Fig. 4. At 1 and 2 days post-lesion, cells at the wound margin are almost ubiquitously labelled, and these include some neurones (large pale nuclei) (Fig. 4A, B). Cells of glial origin (small dark nuclei) are densely labelled with silver grains 2 days after lesion (Fig. 4C–E). By 14 days few cells are labelled at the margin (Fig. 4E). Adjacent sections hybridized with the sense strand show minimal signal (Fig. 4A<sub>3</sub>–E<sub>3</sub>).

### TGF $\beta$ 1 mRNA and the corpus callosum

At low magnification in bright field view, an extensive distribution of dense signal for TGF $\beta$ 1 mRNA is seen through the corpus callosum after 7 days (Fig. 5A), but no labelling is detected with the sense probe (Fig. 5B, D). Based on morphology, location and staining properties, higher magnification suggests that the labelled cells may be of glial origin (Fig. 5C).

### TGF $\beta$ 1 mRNA and microvasculature

In situ hybridization of TGF $\beta$ 1 mRNA following the cortical wound demonstrates a clear association between TGF $\beta$ 1 mRNA and cells of the microvasculature. Low or negligible message is detected in the normal adult rat brain in this experiment. At 2 and 3 days after injury, TGF $\beta$ 1 mRNA is detected in the meninges and in the endothelium of the hippocampal fissure and choroid plexus (Fig. 6); this signal is significantly diminished by 7 and 14 days post lesion (Fig. 3). *Meninges.* In the meninges, sections hybridized with the antisense probe to TGF $\beta$ 1 mRNA show signal in cells containing darkly stained round nuclei (Fig. 6A<sub>1</sub>). No signal is present when sections are hybridized with sense probe (Fig. 6A<sub>2</sub>).

**Hippocampus**—While no TGF $\beta$ 1 mRNA is present in the hippocampus of unlesioned animals, TGF $\beta$ 1 mRNA is visible in the endothelial cells of the hippocampal fissure but not in adjacent sections hybridized with the sense strand (Fig. 6B).

**Choroid plexus**—Low to negligible levels of TGF $\beta$ 1 mRNA are present in the choroid plexus of unlesioned animals (Fig. 6C). This is in striking contrast to the lesioned rats in which a dramatic and extensive induction of TGF $\beta$ 1 mRNA is observed in the choroid plexus (Fig. 6D). Labelled arteriole endothelial cells are seen in sections hybridized with antisense probe but not in sections hybridized with the control sense probe (Fig. 6E).

### TGF $\beta$ 1 immunohistochemistry

Immunocytochemical localization of TGF $\beta$ 1 demonstrates a distribution pattern similar to that of its mRNA (Fig. 7). In this study no immunoreactivity is apparent in the normal rat brain (not shown). The predominant cell type that appears to contain TGF $\beta$ 1 after injury has the appearance of astrocytes that are limited to the tissue bordering the glia limitans (Fig. 7). However, immunopositive macrophage-like cells are visible at all time points examined. The time course of the TGF $\beta$ 1 protein response is very similar to that of its mRNA, immunoreactivity reaches a relative peak 3 days after lesion and is considerably diminished by 14 days. At this time point the staining is effectively restricted to the residual cells that have, at higher magnification (not shown), all the morphological characteristics of macrophages.

## DISCUSSION

These results confirm the observations of others<sup>26,46,50</sup> that, in vivo, TGF $\beta$ 1 expression is normally low in the adult CNS. However, there is an immediate and dramatic induction of TGF $\beta$ 1 expression locally within the neuropile after a penetrating cerebral injury, evidenced



by increased TGF $\beta$ 1 mRNA and immunoreactive protein. Furthermore, the results demonstrate that TGF $\beta$ 1 mRNA expression in the choroid plexus and meninges is also markedly increased after CNS lesion.

Enhanced signal for TGF $\beta$ 1 mRNA is detected in the neural tissue at the margins of the lesion throughout the 1–14-day response period examined, which peaks at 2 days but appears much reduced at 14 days. This transient response is paralleled by immunocytochemical localization of TGF $\beta$ 1 protein, which may arise both from the local release of TGF $\beta$ 1 by platelets and macrophages recruited into the wound, and from de novo synthesis of TGF $\beta$ 1 by neural tissues. A recent report by others demonstrated a similar transient elevation of TGF $\beta$ 1 mRNA (measured by northern analysis) in the hippocampus after an entorhinal cortex lesion, with maximal levels occurring 4–6 days after lesion<sup>28</sup>. The differences in timing of the peak levels may be a result of differences in the time courses of the cellular responses to a discrete cortical wound compared to those that occur by deafferentation.

Based on the known functions of TGF $\beta$ 1 and on the cellular responses to this type of cortical lesion, the results suggest that TGF $\beta$ 1, released into the marginal tissues of a CNS wound, may have a number of potential activities. Thus, TGF $\beta$ 1 released by platelets, neutrophils, activated monocytes, macrophages and lymphocytes may initially regulate the inflammatory process by actions on the recruitment, priming and activation of these cells in an autocrine or paracrine manner (reviewed in ref. 52). The observations of immunopositive cells of the monocyte/macrophage lineage within the lesion site at all post-injury time-points examined in this experiment supports this immunoregulatory role for TGF $\beta$ 1. The ability of TGF $\beta$ 1 to up-regulate both its own gene expression and the expression of other growth factor genes suggests a mechanism for the initiation and amplification of its signal within target tissues<sup>25</sup>. This response may underlie the local induction of TGF $\beta$ 1 expression that is observed in damaged CNS tissue.

The many possible roles for TGF $\beta$ 1 after CNS injury include chemoattraction of microglia<sup>53</sup>, which rapidly migrate to the site of CNS injury<sup>13</sup>. Some of these microglia may originate from cells of the monocyte/macrophage lineage<sup>19</sup>. The migration and proliferation of microglia appears critical for subsequent proliferation of astrocytes, and both these cell types appear to secrete multiple growth factors that regulate the many responses to injury, including neuronal sprouting and endothelial cell proliferation<sup>22</sup>. TGF $\beta$ 1 may also stimulate the release or synthesis of neurotoxic cytokines from macrophages or microglia<sup>49</sup>.

The predominant cell-type localizing TGF $\beta$ 1 and its mRNA in the wound margin has the features of the astrocyte, suggesting that this factor could act in a paracrine or autocrine manner to stimulate astroglia after injury. Astrocytes proliferate and migrate in injury-responsive neural tissue and eventually form a glia limitans around the wound margins<sup>24</sup>. However, TGF $\beta$ 1-expressing glial cells are not confined to the wound borders, but also coincide with the medio-lateral proliferation of astrocytes in the corpus callosum. Gliomas are reported to express TGF $\beta$ 1, supporting the notion that glia are capable of TGF $\beta$ 1 synthesis<sup>41</sup>. The action of TGF $\beta$ 1 on astrocytes in vitro is very much dependent on the interaction with other growth factors and on the dosage used<sup>7,17</sup>. For example, in the

absence of other growth factors TGF $\beta$ 1 inhibits growth, but will potentiate the mitogenic effect of basic FGF<sup>17</sup>. TGF $\beta$ 1 may stimulate NGF synthesis in the CNS<sup>18</sup> and NGF is mitogenic for astrocytes<sup>12</sup> as well as neurotrophic<sup>39</sup>. TGF $\beta$ 1 also stimulates glial fibrillary acidic protein<sup>39</sup> and cystatin C mRNA synthesis in astrocyte-like SFME (serum-free mouse embryo) cells<sup>45</sup>, consistent with a role in glial differentiation. The association of TGF $\beta$ 1 with glial cells and its actions as a potent inhibitor of glutamine synthetase implies that it may also impair some glial functions, interfering with their ability to support neurones<sup>43</sup> and thus contributing to the pathology observed after CNS injury.

The temporal and spatial change in TGF $\beta$ 1 localization in the neuropile may also initiate the migration of fibroblasts into the wound, a phenomenon which occurs 4 days after injury<sup>24</sup>. TGF $\beta$ 1 is a potent chemoattractant for fibroblasts to the site of injury<sup>32</sup> as well as controlling fibroblast deposition of extracellular matrix proteins and the synthesis of degradative enzymes (reviewed in ref. 36). A key role for TGF $\beta$ 1 in the regulation of fibrosis at CNS wound sites is strongly suggested. The contraction of the scar tissue associated with the CNS injury observed here and in other studies after lesioning<sup>5,24</sup> is consistent with the role TGF $\beta$ 1 is purported to play in the periphery<sup>27</sup>.

In peripheral tissues, TGF $\beta$ 1 also plays an important role in regulating angiogenesis<sup>37</sup> and in the CNS TGF $\beta$ 1 may also regulate the neovascularization induced by injury in the damaged neuropile and in the deposited scar tissue<sup>3</sup>. This activity is supported by our observation of increased levels of TGF $\beta$ 1 mRNA in vascular endothelial cells of the microvasculature of the tissue around the lesion site.

The transient increase in TGF $\beta$ 1 mRNA observed in the meninges is of particular interest and may also relate to changes in the local microvasculature. This region is outside the blood-brain barrier and consists of blood vessels, fibroblasts and macrophages. The local induction of TGF $\beta$ 1 in the meninges following injury may additionally play a role in promoting the early repair of the blood-brain barrier, by stimulating fibroblast migration and deposition of extracellular matrix proteins. The effect of TGF $\beta$ 1 may depend on its interaction with other growth factors, especially basic FGF<sup>4,9,31</sup>. Therefore the absence of basic FGF in the meninges but not in the margin of the wound after injury<sup>11,22</sup> may signify different functions of TGF $\beta$ 1 in these two regions.

As in the meninges, endothelial cells in the choroid plexus and hippocampal fissure also appear to transiently and strongly express TGF $\beta$ 1 mRNA. The role of TGF $\beta$ 1 in the choroid plexus is unknown, although it may involve attracting microglia which have phagocytized debris and are clearing it into the ventricle or vessels. Clearly, the CSF may serve as a route for the wider distribution of newly synthesized TGF $\beta$ 1. TGF $\beta$ 1 is present in the choroid plexus on both the ipsilateral and contralateral sides of the lesion. Whether the extensive bilateral distribution of TGF $\beta$ 1 mRNA throughout the lateral ventricles is due to an injury-associated soluble substance which circulates in the CSF and induces TGF $\beta$ 1 mRNA in the ventricle or due to other factors (for example, interaction with microglia or nervous stimulation) is unclear.



Because the increase in TGF $\beta$ 1 synthesis following a cortical knife wound is an early response to CNS injury, the physiological implications of these findings are particularly interesting. A role for TGF $\beta$ 1 in the acute inflammatory response to injury is certainly indicated. Although TGF $\beta$ 1 mRNA is induced in endothelial cells of the local neuropile, meninges, choroid plexus and hippocampal fissure, its function on the microvasculature in these situations remains to be defined, although an autocrine angiogenic activity could be implied. Of major importance, the distribution of TGF $\beta$ 1 and its mRNA in the wound and the known roles of TGF $\beta$ 1 in peripheral tissues are indicative of a role in formation of the glia limitans externa and the associated fibrous matrix of the CNS scar. Several investigators have suggested that the presence of a glial scar after CNS injury can have profound deleterious effects on neuronal survival and neuritic extensions. If the formation of the glial/mesodermal scar is dependent on the well-described ability of TGF $\beta$ 1 to stimulate astroglia and matrix deposition, then the findings described here point to the therapeutic value of developing TGF $\beta$ 1 antagonists to block matrix deposition and enhance the functional recovery from CNS injury and trauma.

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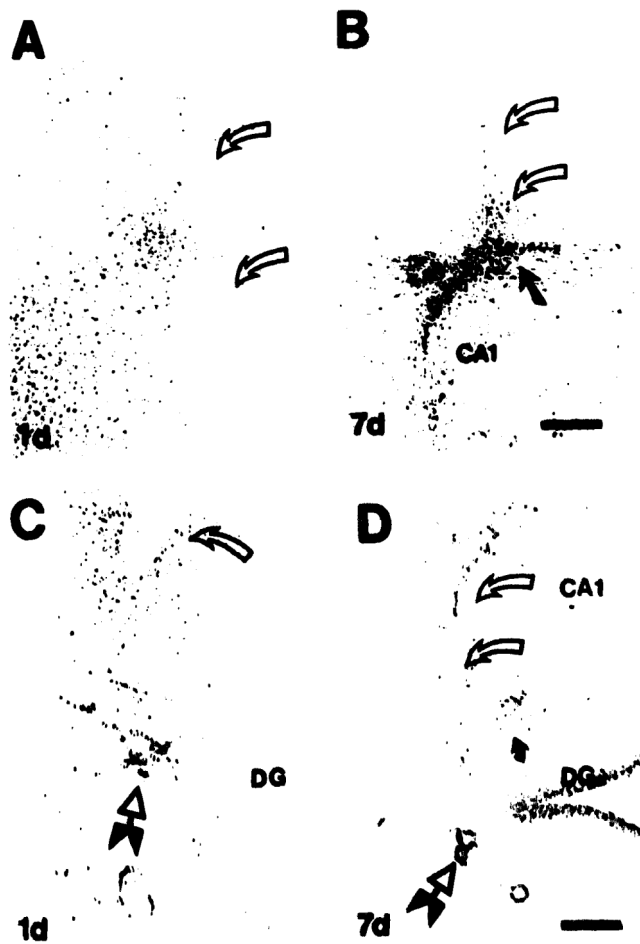
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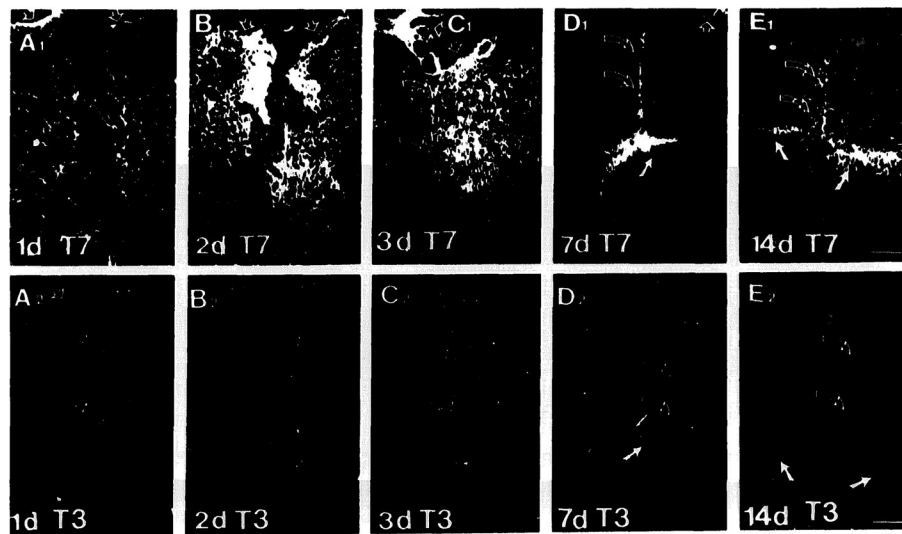
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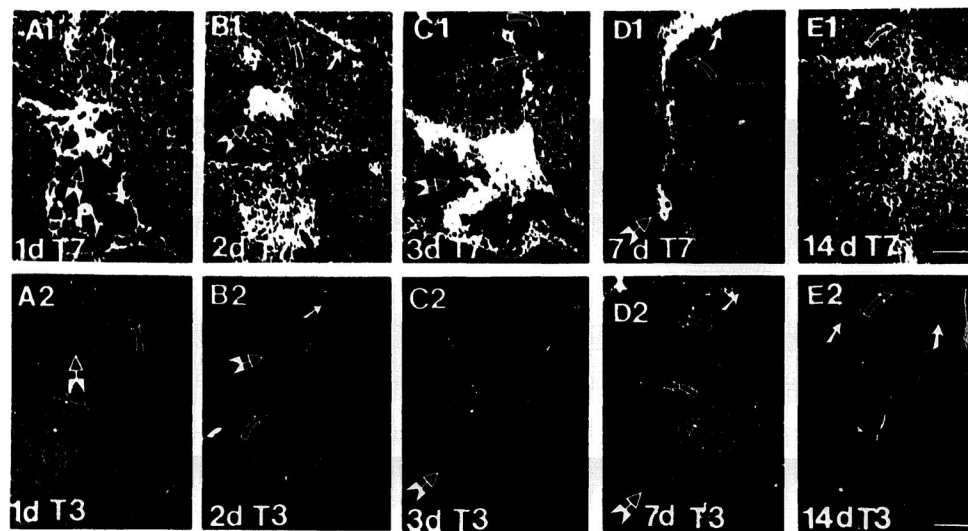
**Fig. 1.** Bright field micrograph of the brain 1 day and 7 days after injury. The caudal (panels A and B) and rostral (panels C, D) portions of the lesion are shown. The curved open arrows show the region of the lesion in the cortex at 1 day (panels A, C) and 7 days (panels B and D) after the lesion. Bar = 250  $\mu$ m. Note that the glial scar is well established by 7 days. The corpus callosum (large closed arrow), hippocampal fissure (closed curved arrow), the choroid plexus (straight open arrow with tail), dentate gyrus (DG) and CA1 region of the hippocampus are shown for orientation.



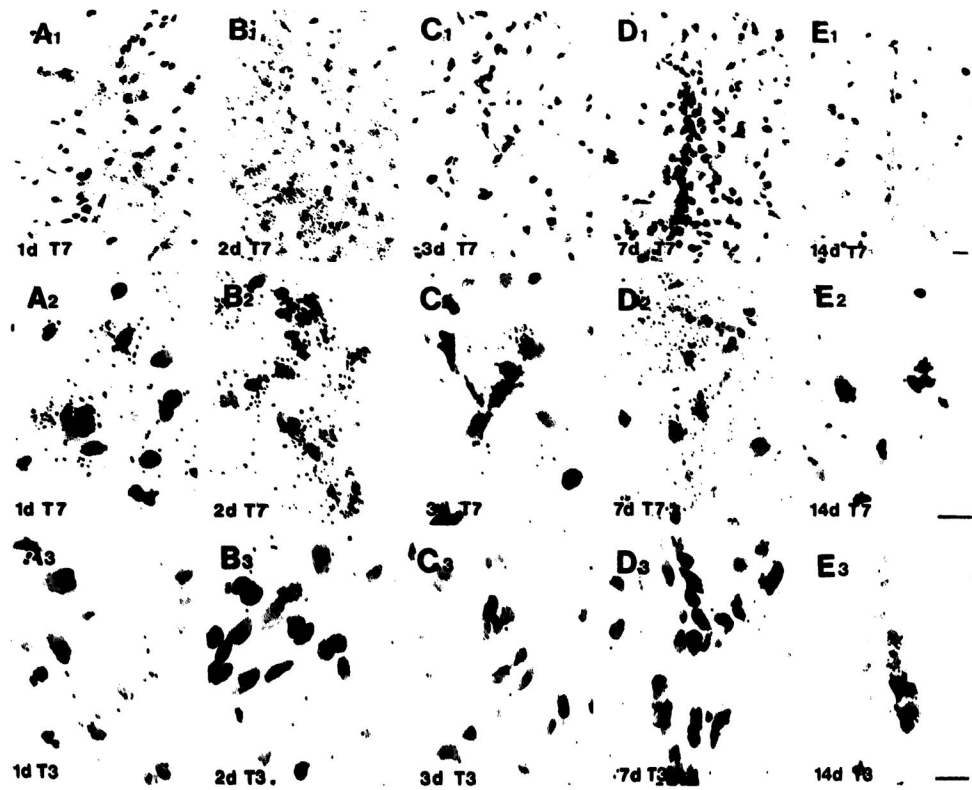
**Fig. 2.**

Dark field micrographs showing  $TGF\beta 1$  mRNA in the caudal region of the wound.  $TGF\beta 1$  mRNA (T7) is observed at 1 day (panel A<sub>1</sub>), 2 days (panel B<sub>1</sub>), 3 days (panel C<sub>1</sub>), 7 days (panel D<sub>1</sub>) and 14 days (panel E<sub>1</sub>) following the lesion. When sections are hybridized with the sense strand (T3) the signal is minimal (panels A<sub>2</sub>–E<sub>2</sub>). The curved open arrows show the margin of the lesion.  $TGF\beta 1$  mRNA, but not the control signal, is also observed in the corpus callosum (panels D<sub>1</sub> and D<sub>2</sub>) as shown by the closed straight arrows and in the meninges (panels A<sub>1</sub>–D<sub>1</sub> and A<sub>2</sub>–E<sub>2</sub>) as shown by straight open arrows. The signal is most intense at 2 and 3 days after lesion. At 7 days and 14 days the signal is less diffuse and only present within the boundary of the well defined glial scar. Bar = 250  $\mu$ m.

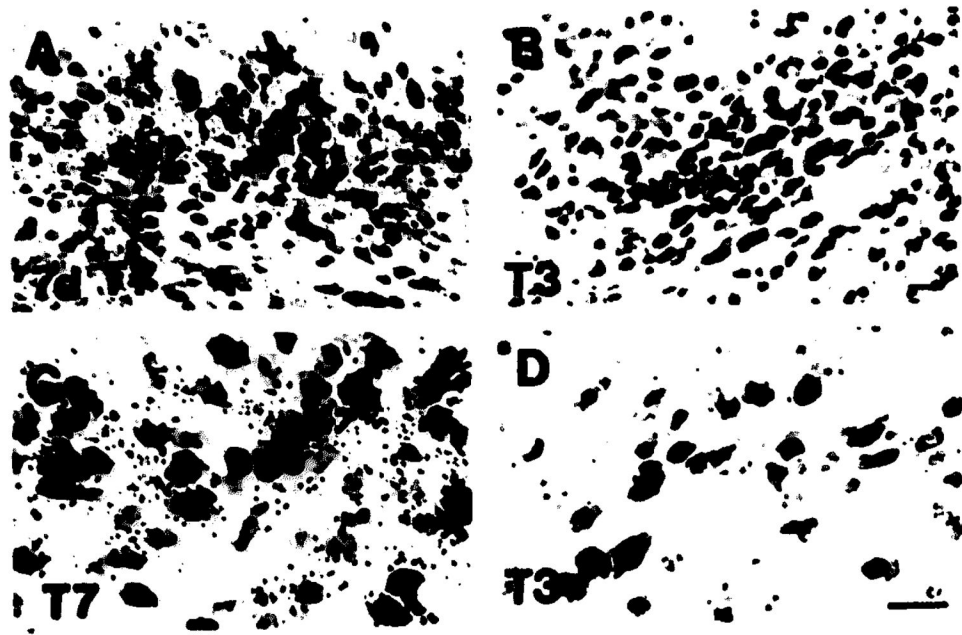




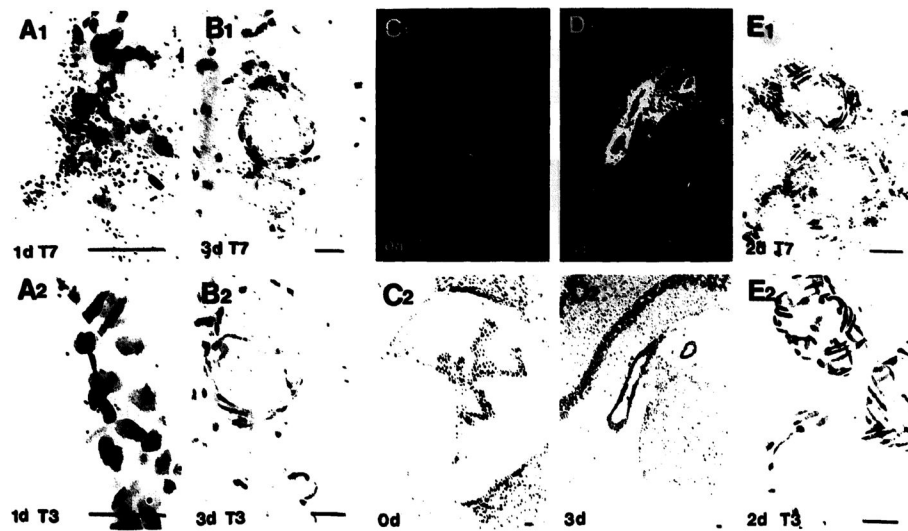
**Fig. 3.** Dark field micrographs showing  $TGF\beta 1$  mRNA in the rostral region of the wound.  $TGF\beta 1$  mRNA (T7) is observed at 1 day (panel A<sub>1</sub>), 2 days (panel B<sub>2</sub>), 3 days (panel C<sub>1</sub>), 7 days (panel D<sub>1</sub>) and 14 days (panel E<sub>1</sub>) following the lesion. When sections are hybridized with the sense strand (T3) the signal is minimal (panels A<sub>2</sub>–E<sub>2</sub>). The curved open arrows show the margin of the lesion.  $TGF\beta 1$  mRNA, but not the control signal, is also observed in the corpus callosum (panels B<sub>1</sub>, D<sub>1</sub>, E<sub>1</sub> and B<sub>2</sub>, D<sub>2</sub>, E<sub>2</sub>) as shown by the closed straight arrows and in the choroid plexus (panels A<sub>1</sub>–D<sub>1</sub> and A<sub>2</sub>–E<sub>2</sub>) as shown by straight open arrows with tails. The signal appears considerably diminished by 14 days. Bar = 250  $\mu$ m.



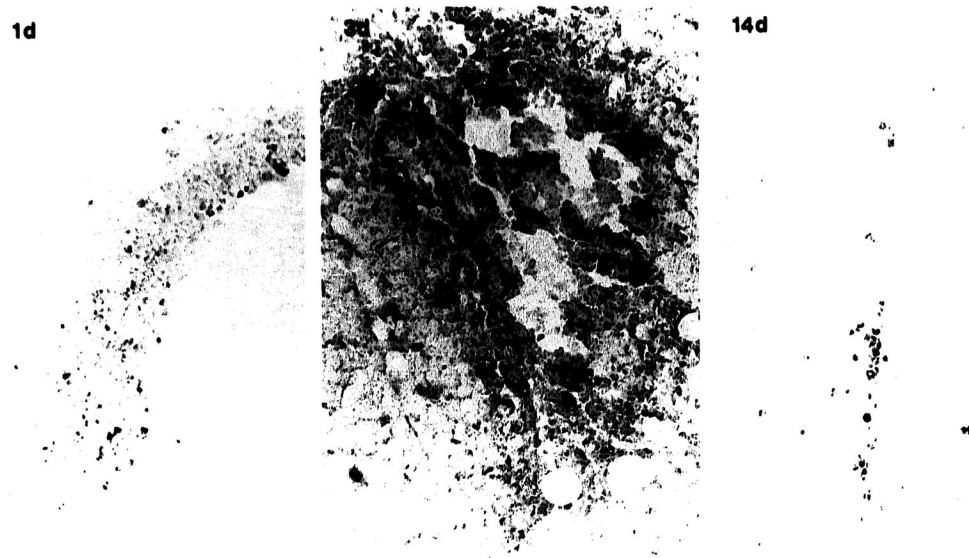
**Fig. 4.** High magnification bright field micrographs of cells in the margin of the wound. TGF $\beta$ 1 mRNA (T7) is observed associated with selective cells in the margin of the lesion at 1 day, (panel A) 2 days (panel B), 3 days (panel C), 7 days (panel D) and 14 days (panel E) after the lesion but not in cells hybridized with the sense strand (T3, row 3). Initial signs of the glial scar are seen at 3 days and are well defined by 7 days, at which time the mRNA appears less diffuse. High magnification reveals that these cells are typically glial cells (row 2). Bar = 10  $\mu$ m.



**Fig. 5.** The presence of TGF $\beta$ 1 mRNA in the corpus callosum 7 days following a defined lesion. There is an intense signal for TGF $\beta$ 1 mRNA (Panels A and C) in the corpus callosum at the point of lesion penetration. Control sections show no signal (T3, panels B and D). Bar = 10  $\mu$ m.



**Fig. 6.** TGF $\beta$ 1 mRNA in the meninges, hippocampal fissure and choroid plexus after injury. Intense TGF $\beta$ 1 mRNA (T7) is observed in the meninges (panel A) and hippocampal fissure (panel B) but not in the control sections hybridized with the sense strand (T3). In the choroid plexus, TGF $\beta$ 1 mRNA is almost non-detectable in uninjured animals (0 day, C1) but very intense following injury (panels D<sub>1</sub>, E<sub>1</sub>). Control sections show no signal (T3). Bright field of low-power micrographs are shown in panels C<sub>2</sub> and D<sub>2</sub>. Bar = 25  $\mu$ m.



**Fig. 7.** Immunolocalization of TGF $\beta$ 1 after injury. Immunoreactive TGF $\beta$ 1 is seen diffusely in the neuropile along the borders of the lesion after 1 day and this increases at 3 days. Staining is residual by 14 days and mostly confined to the macrophages remaining in the center of the wound.

Bar = 10  $\mu$ m.