ASSOCIATION OF CALCIUM BUFFERING CAPACITY AND LOCAL INFLAMMATORY REACTIONS IN INJURY OF MOTOR NEURONS

Ph.D. Thesis

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# TABLE OF CONTENTS

## INTRODUCTION ................................................................................................................... 1
  - General context of the study .............................................................................................. 1
  - Brief overview of amyotrophic lateral sclerosis ............................................................... 2
    - Clinical features ........................................................................................................... 2
    - Pathological findings ................................................................................................. 2
    - Epidemiology .............................................................................................................. 3
  - Identified basic mechanisms of degeneration in ALS ....................................................... 3
    - Glutamate excitotoxicity ............................................................................................. 4
    - Oxidative damage ........................................................................................................ 4
    - Mitochondrial dysfunction .......................................................................................... 5
    - Protein misfolding and aggregation ............................................................................ 5
    - Defects in axonal transport .......................................................................................... 6
    - Immune-inflammatory dysfunction ............................................................................. 6
    - Impairment of the blood-brain/spinal cord barrier ...................................................... 7
    - Calcium-mediated toxicity .......................................................................................... 7
  - A synthesis of the basic mechanisms to a putative pathomechanism underlying ALS .... 8
    - Calcium is at centerstage in degeneration ................................................................... 8
  - Expectations from increased calcium buffering capacity .............................................. 10
  - Proof of concept experiment: protection by calcium binding in ALS ............................ 11
    - Previous experiments relevant to the present study .................................................. 12
    - Problems to be addressed by further experiments ..................................................... 12

## AIMS OF THE STUDY ........................................................................................................ 13

## MATERIALS AND METHODS .......................................................................................... 13
  - Experimental animals ...................................................................................................... 13
  - Maintenance of the transgenic colony ............................................................................ 13
  - Characterization of the motoneuron pools in PV+/+ animals ........................................ 14
  - Axotomy/target deprivation of hypoglossal and oculomotor neurons ............................ 16
  - Axotomy of the sciatic nerve .......................................................................................... 17
  - Electron microscopic detection of calcium ..................................................................... 17
  - Specimen preparation ................................................................................................... 17
Specificity control of the histochemical reaction for calcium ........................................ 18
Semiquantitative evaluation of the intracellular amount of EDDs .......................... 19
Immunohistochemical characterization of inflammatory markers in the spinal cord .... 20
CD11b staining ........................................................................................................... 20
MCP1 staining .......................................................................................................... 21
Evaluation of the functional recovery after sciatic nerve cut .................................... 20
RESULTS ......................................................................................................................... 21
Calcium increase is reduced after injury in hypoglossal motor neurons of PV+/+ mice 22
Inflammatory reaction can be quantified by systematic sampling and background
corrected segmentation using immunohistochemistry ................................. 25
Increased motoneuronal calcium buffer reduces local inflammation after acute lesion . 27
DISCUSSION ................................................................................................................. 31
The role of CaBPs in neuronal degeneration ............................................................ 31
Lesion induced by axotomy ....................................................................................... 32
The role of inflammatory factors in injury of motor neurons ................................. 33
Quantification of the inflammatory markers by light microscopic IHC .................. 34
Relevance of the results in relation to the mechanisms of ALS ......................... 35
SUMMARY .................................................................................................................. 37
REFERENCES .......................................................................................................... 38
ACKNOWLEDGEMENTS .......................................................................................... 50
APPENDIX
INTRODUCTION

General context of the study

Our study is aimed at understanding the mechanisms underlying the selectivity of degeneration of motor neurons during amyotrophic lateral sclerosis (ALS): why do certain groups of motor neurons succumb to the disease while others are spared. Our hypothesis is that the impaired calcium homeostasis (i.e. increased calcium level) plays a pivotal role in the degeneration of motor neurons (Appel et al, 2001), and by its stabilization an increased resistance could be provided (Siklós and Appel, 2005). Accordingly, factors affecting the maintenance of the intracellular calcium equilibrium, such as the calcium buffering capacity, having the potential to reduce overshoots in calcium signaling, might be a significant feature contributing to the selective resistance of these cells.

Indeed, the increased level of intracellular calcium in the affected motor neurons was documented in ALS patients (Siklós et al, 1996), which study was paralleled by the demonstration of a correlation between the resistance of such cells against injury and their calcium binding protein (CaBP) content (Alexianu et al, 1994; Elliot and Snider, 1995). This observation was subsequently confirmed by a series of in vitro (Ho et al, 1996), ex vivo (Lips and Keller, 1998; Mosier et al, 2000; Palecek et al, 1999; Vanselow and Keller, 2000; recently reviewed by von Lewinski and Keller, 2005) and in vivo (Obál et al, 2006; Siklós et al, 1998) experiments.

With the advent of the transgenic technology, providing gold standards for the research of motor neuron disease (Nagai et al, 2006; Tovar-y-Romo et al, 2009), a direct method became available to test whether an increased calcium buffering capacity provides protection in ALS-like conditions by breeding transgenic animals simultaneously possessing increased calcium buffering capacity and presenting ALS-specific pathological mechanisms. Regarding the documented universal role of an increased level of calcium in cell destruction (Berridge et al, 1998), and the crucial role of calcium increase in a positive feedback loop of the detrimental processes (Appel et al, 2001), the calcium binding protein parvalbumin (PV)-upregulated mice exhibited only an unexpectedly mild protection in a transgenic model of ALS (Beers et al, 2001). These results reduced our confidence in the neuroprotection strategy based on the increased calcium buffering capacity.
To rule out the possibility that our basic proposal (stabilization of the calcium homeostasis) is not relevant at all in rescuing injured motor neurons, experiments were planned to characterize motor neuron populations in PV-upregulated transgenic mice after a sublethal injury, and tested whether a naturally vulnerable cell type could be indeed transformed to a resistant phenotype in these animals. A positive answer to this question may suggest that while the strategy of neuroprotection built on the improved stability of the calcium homeostasis may be correct, a method based on the increased buffering capacity alone, may not be sufficient. In this case, a more efficient approach should be sought.

**Brief overview of amyotrophic lateral sclerosis**

*Clinical features*

Amyotrophic lateral sclerosis (also known as Lou Gehrig’s disease), the most common adult motor neuron disease (Cleveland and Rothstein, 2001), is a progressive, lethal disorder characterized by degeneration of both upper and lower motor neurons with clinical signs in four areas (brainstem, cervical-, thoracic- and lumbosacral regions) of the central nervous system. Anatomically, the axons of the upper motor neurons (Betz cells, located predominantly in layer V of the motor cortex) connect with the lower motor neurons positioned in the motor nuclei of the brainstem and the anterior horn of the spinal cord. Lower motor neurons project mainly to skeletal muscles. As the disease advances, there are progressive manifestations of combined dysfunctions of lower motor neurons (atrophy, cramps and fasciculations) and upper motor neurons (spasticity and pathological reflexes) in the absence of sensory symptoms (Mulder, 1982). According to the El Escorial- and the revised Airlie House criteria, the diagnosis of clinically definite ALS requires the presence and progression of both upper- and lower motor neuron signs in at least three body regions (Swash, 2000). Clinical, laboratory and neuropathological criteria have also been established for less definitive categories (Cwik, 2006). The relentless progression of muscle atrophy and weakness culminates in respiratory death within 5 years in the majority of the patients (Rowland and Shneider, 2001).

*Pathological findings*

Until the last two decades, the main pathological findings in postmortem tissue of ALS patients were loss of motor neurons, commonly accompanied by astrocytosis, in the precentral...
gyrus, brainstem and spinal cord. Surviving motor neurons showed features of degeneration such as atrophy of the cell body with large number of lipofuscin granules. Secondary changes involved the degeneration of the pyramidal tract, loss of nerve fibers and neurogenic abnormalities in the muscles. However, none of these changes are exclusive for ALS (Hays, 2006). ALS-specific pathological alterations were identified during the past 15-20 years by recognition of intracytoplasmic inclusions in standard pathological (hematoxylin and eosin stained) sections such as Bunina bodies, with an occurrence in the ALS tissue of 65-95% (Hays, 2006). More importantly, after the introduction of immunohistochemical (IHC) analysis into the pathological practice, ubiquitinated inclusions with nearly 100% occurrence and high specificity for ALS were discovered (Kato, 2008). Furthermore, the protein composition of such inclusions, identified also by IHC provided suggestions for the pathobiology of ALS, as well (Strong et al, 2005).

**Epidemiology**

The incidence of sporadic ALS is reported to be between 1.5 and 2.7 per 100,000 population/year, which corresponds (in agreement with an average of 2-5 years survival) to a point prevalence of 2.7 to 7.4 (Wijesekera and Leigh, 2009). There are conflicting data on gender (Strong, 2003), however, it seems, that the small male preponderance (1.5 male to female ratio) is diminishing with age, suggesting an estrogen-dependence in the pathomechanism (Chio et al, 2009). Although most cases are classed as sporadic ALS (sALS), 5-10% of the patients have a family history (familial ALS; fALS) with often Mendelian inheritance and high penetrance (Wijesekera and Leigh, 2009). Twenty percent of cases with autosomal dominant fALS (and 2% of sALS patients) possess mutations in the Cu/Zn superoxide dismutase (SOD1) gene. Other genes causing familial form of the disease include alsin, senataxin (STX), vesicle associated membrane protein (VAPB), angiogenin, fused in sarcoma/translated in liposarcoma (FUS/TLS) and gene encoding the TAR-DNA binding protein TDP-43 (Valdmanis et al, 2009).

**Identified basic mechanisms of degeneration in ALS**

According to the contemporary view, ALS is considered as a non cell autonomous (Ilieva et al, 2009) and multifactorial disease (Eisen, 2009). Even though its cause is basically unknown (except for the inherited cases constituting the absolute minority of the patients) several
mechanisms have been identified, which either alone, but rather in a concerted action contribute to the degeneration of motor neurons (Boillée et al, 2006). These include: glutamate excitotoxicity, oxidative damage, mitochondrial dysfunction, protein misfolding and aggregation, defects in axonal transport, immune-inflammatory dysfunction, impairment of the blood-brain/spinal cord barrier and calcium-mediated toxicity.

**Glutamate excitotoxicity**

This well recognized mechanism of neuronal death resulting from repetitive cell firing (Choi, 1988) has been implicated in the pathogenesis of ALS for a long time, furthermore, Riluzole, an anti-glutamatergic drug, is currently the only one medicine approved to treat ALS (Rattray and Bendotti, 2006). Several experimental studies and clinical observations document that the toxicity exerted on the motor neurons is mediated by an excess level of extracellular glutamate, due to impairment of the astrocytic glutamate transporter EAAT2 (excitatory amino acid transporter 2), which support a non cell autonomous pathological mechanism in ALS (Rothstein, 2009).

In ALS, the 2-amino-3-(3-hydroxy-5-methylisoxazol-4-yl) propionic acid (AMPA) subtype of glutamate receptors is of particular importance. AMPA receptors, assembled from 4 subunits, are normally impermeable to calcium. However, if the GluR2 subunit is missing, the receptors become highly permeable to this ion (Buckingham et al, 2008). Low expression of the GluR2 protein subunit (Shaw et al, 1999) and decreased levels of the mRNA for GluR2 (Williams et al, 1997) are detected in motor neurons of ALS patients.

**Oxidative damage**

Oxidative stress develops in a system if the balance between the production and neutralization of reactive oxygen species (ROS) is shifted towards production, or the capacity of the repair mechanisms is overwhelmed. The main source of cellular ROS is the leakage of electrons from the mitochondrial respiratory chain which results in production of superoxide anions (O$_2^{-}$) and hydrogen peroxide (H$_2$O$_2$) (Halliwell and Gutteridge, 1999), but other cellular oxidative enzymes in the cytoplasm and the endoplasmic reticulum contribute, as well. The mildly reactive O$_2^{-}$ and H$_2$O$_2$ are further transformed to peroxynitrite (ONOO$^-$) and hydroxyl radicals (•OH), which cause nitration of tyrosine residues of accessible proteins, lipid-
peroxidation and oxidative damage of DNA (Barber and Shaw, 2010), which have been documented in ALS patients (Beal et al, 1997).

The role of oxidative mechanisms in the pathobiology of ALS gained an outstanding attention after identification of missense mutations of the SOD1 gene in a population of the patients (Rosen et al, 1993). A novel (gain of) function of the mutated enzyme has been proposed, which exaggerates the oxidative burden of the affected cells (Julien, 2001). An interesting aspect of the malfunction of the mutant SOD1 has been recently discovered: the mutant enzyme, after association with Rac1, a small GTPase that controls the activation of NADPH oxidase (nicotinamid adenosine dinucleotide phosphate, a multiprotein membrane-associated complex whose catalytic subunit is Nox2), locks Nox2 in its active, superoxide-producing form (Harraz et al, 2008). Thus, instead of removing intracellular superoxide, mutant SOD1 is responsible for overproduction of extracellular $O_2^\bullet-$ of microglial (phagocytic cell) origin, providing additional evidence for the non cell autonomous pathology underlying ALS (Boillée et al, 2008).

**Mitochondrial dysfunction**

This is a common feature of many neurodegenerative diseases (Chaturvedi and Beal, 2008). Despite of the documented mitochondrial alterations in patients’ biopsy material (Siklós et al, 1996) and in some transgenic models of the disease (Siklós et al, 1998), there is no uniform view how mitochondrial function is actually affected in ALS. Several lines of evidence support that mitochondrial calcium handing mechanism might be altered (see e.g. Damiano et al, 2006), which is fully compatible with the above described excitotoxic mechanisms, thus may be central to the neuronal damage (Boillée et al, 2006). The documented damage evoked by the deposition of mutant SOD1 on the cytoplasmic face of the outer mitochondrial membrane (Vande Velde et al, 2008) together with the notion of an impaired mitochondrial function of astrocytes bearing the mutant enzyme, which promotes the degeneration of motor neurons (Cassina et al, 2008), provide further clue to the cooperative role of different cell types in the disease mechanism.

**Protein misfolding and aggregation**

Another common feature of neurodegenerative disorders that protein aggregates develop, which may be characteristic for the disease and the degenerating neuronal population, such as senile plaques (amyloid beta aggregation) and neurofibrillary tangles (phosphorylated tau
aggregation) in Alzheimer’s disease, Lewy bodies (α-synuclein aggregation) in Parkinson’s disease, prion plaques (aggregation of the scrapie isoform of the prion protein) in Creutzfeld-Jacob disease, inclusions (Huntingtin aggregates) in Huntington’s disease, etc. (Chaudhuri and Paul, 2006). Since the hypothesis of the mode of action of the mutant SOD1, based on the reduced scavenging activity (Beckman et al, 1993), after the first description of mutant SOD1 enzyme in ALS tissue failed, by now it has been replaced with an idea that the mutant protein misfolds, tends to aggregate which confer toxicity to motor neurons. Besides mutant SOD1, other cytoskeletal proteins, various kinases, heat shock proteins and proteins of the endoplasmic reticulum also contribute to the ubiquin-immunopositive, detergent-insoluble material constituting the cytoplasmic inclusions in ALS (Basso et al, 2009). However, recent observations accentuate the role of SOD1 in the pathomechanism documenting that the wild type enzyme may acquire similar toxic properties as the mutant form possesses through oxidative damage (Ezzi et al, 2007), which may offer a convergence point where the mechanisms underlying fALS and sALS merge. Furthermore, misfolded proteins could also be secreted (Urushitani, 2006), which interact with microglial cells, and by triggering the release of noxious substances provides a molecular clue for an intercellular mechanism how does mutant SOD1 exert its toxic effect on motor neurons (Zhao et al, 2009).

**Defects in axonal transport**

Aberrant organization and especially misaccumulation of neurofilaments in spheroids in the cell body and proximal axons of motor neurons is a hallmark of both sALS and fALS. These spheroids may physically compromise the axonal transport mechanism, which has been detected in several transgenic models of ALS (Rao and Nixon, 2003). The mechanisms by which the impaired axonal transport contribute to the degeneration of motor neurons is not clear, but could include inadequate delivery and removal of distal mitochondria, and slow the retrograde transport of trophic factors from the periphery (De Vos et al, 2008). Interestingly, overexpression of the neurofilament heavy (NF-H) gene in a transgenic mouse model of ALS provided neuroprotection by a mechanism assumed to be based on attenuating calcium increases at the perikaryal region of motor neurons through calcium binding by the multiple calcium binding sites of NF-H proteins (Coulliard-Després et al, 1998).

**Immune-inflammatory dysfunction**
The immune privilege of central nervous system (CNS) has been considered as experimentally well defined, however, recent data radically altered this viewpoint: peripheral immune cells can cross the intact blood-brain barrier (BBB), neurons and CNS glial cells actively regulate macrophage and lymphocyte response, and there are immunocompetent cells in the CNS, such as microglial cells (Carson et al, 2006). Although it is still controversial to what extent the role of the adaptive and innate immunity is destructive or protective, their participation in the disease mechanism is undoubted (Appel et al, 2010).

**Impairment of the blood-brain/spinal cord barrier**

The involvement of the blood vessels in the pathomechanism of ALS has already been suggested by the observation that the deletion of the hypoxia-response element in the (angiogenic) vascular endothelial growth factor (VEGF) results in an ALS-like phenotype in mice (Oosthuyse et al, 2001). Furthermore, the participation of the BBB or blood-spinal cord barrier (BSCB) in the pathogenesis of ALS is supported by the recent results from biochemical (Zhong et al, 2008) and morphological studies (Garbuzova-Davis et al, 2007a, b), obtained with the transgenic mouse model of ALS, showing that disruption of the BBB and BSCB might precede motor neuron degeneration in the spinal cord. Although impairment of the BBB is not a prerequisite to initiate autoimmune responses in the CNS, the accompanying increased exposure of the CNS parenchyma to blood-borne products, particularly to humoral components of the immune response, and the decreased efficacy of the efflux transporters to remove potentially toxic molecules (e.g. excitatory transmitters) may augment the degenerative processes (Carson et al, 2006). In this aspect, the documented presence of immunoglobulin G (IgG) in spinal cord and motor cortex of ALS patients (Engelhardt and Appel, 1990), and the ability of this purified IgG to interact with voltage-gated calcium channels and increase calcium influx might be of primary importance (Engelhardt et al, 1995; Pullen and Humphreys, 2000).

**Calcium-mediated toxicity**

In excitable cells, like neurons, ionic calcium plays an universal second messenger role, converting electrochemical signal to physiological responses, thus its concentration and intracellular distribution require tight regulation (Brini and Carafoli, 2000). A concentration gradient of Ca\(^{2+}\), spanning four orders of magnitude across the plasma membrane of neurons presents a huge diving force, thus guarantees fast signal transduction in physiological
condition, but could easily lead to large increases in \([\text{Ca}^{2+}]\), and cellular malfunction, if the control mechanisms are compromised (Nicotera and Orrenius, 1998; Sattler and Tymianski, 2000). Impaired \(\text{Ca}^{2+}\) homeostasis evoked by injury or alterations of \(\text{Ca}^{2+}\)-regulating proteins may launch several mechanisms some of them may lead directly to neuronal death (Mattson, 2007). These processes include activation of cystein proteases, calpains resulting in downregulation of \(\text{Na}^{+}/\text{Ca}^{2+}\) exchanger and impairing \(\text{Ca}^{2+}\) extrusion (Celsi et al, 2009), activation of caspases that degrade a variety of cytoskeletal proteins, membrane receptors and metabolic enzymes (Chan and Mattson, 1999) and activation/induction of pro-apoptotic proteins such as Bax, Par-4, p53 (Culmsee and Mattson, 2005). Besides these processes, numerous indirect mechanisms are launched, as well, leading to apoptotic or necrotic cell death (Marambaud et al, 2009).

In ALS, the relevance of the altered calcium level and the impaired calcium homeostasis have been documented in motor axon terminals from patients’ muscle biopsies (Siklós et al, 1996), as well as in \textit{in vitro} (Colom et al, 1997) and \textit{in vivo} models of ALS-like injury of motor neurons (Siklós et al, 1998).

**A synthesis of the basic mechanisms to a putative pathomechanism underlying ALS**

Although its etiology is unknown, several mechanisms have been proposed, briefly described above, which most likely through their synergistic action, impair the function and destroy the structure of motor neurons. For the sake of argumentation disregarding the (unknown) initial cause, we demonstrate below, that the individual destructive processes are interconnected in such a way that they mutually augment each other with a positive feedback, thus constitute a complex pathomechanism in which calcium ions play a pivotal role.

**Calcium is at centerstage in degeneration**

A strong relationship exists between excessive \(\text{Ca}^{2+}\) influx and glutamate-triggered neuronal injury (Arundine and Tymiansky, 2003), first proposed in ALS twenty years ago (Rothstein et al, 1990). As a consequence of overstimulation, increased amount of calcium enters motor neurons through AMPA receptors, permeable to \(\text{Ca}^{2+}\) probably due to lack of astrocytic control to upregulate the GluR2 subunit (Van Damme et al, 2007). Elevated calcium, through activation of enzymes like phospholipase A2, calpain, xantin-oxidase, increase the production of free radicals (Mitumoto et al, 1997) directly or indirectly after the uptake of \(\text{Ca}^{2+}\) by
mitochondria (Carriedo et al, 2000). An important consequence of the oxidative stress on membrane proteins (ion pumps and exchangers) is an increased calcium influx, thus the process amplifies itself (Ermak and Davies, 2001). Nevertheless, other proteins such as SOD1, providing the main intracellular antioxidant defense may suffer oxidative damage, as well. Thus, since oxidative stress may equally develop either due to the presence of a mutant SOD1, or to wild type enzyme underwent oxidative damage gaining similar toxic properties as the mutant form (Kabashi et al, 2007), the calcium-mediated amplification cascade persists regardless of the initial trigger.

Another feedback loop of destructive processes with converging effect on motor neurons is associated with glial cells, particularly with astrocytes (Julien, 2007). Although the exact mechanisms are still debated, how do these cells contribute to the degeneration of motor neurons (Foran and Trotti, 2009), certain components of this cascade have already been identified. First, it has been demonstrated that glial glutamate transporter EAAT2 is prone to oxidative injury provided by ROS produced in motor neurons (Rao et al, 2003; Rao and Weiss, 2004), with a consequence of reduced capacity of astrocytes to maintain glutamate level at the synaptic cleft, which as discussed above leads to calcium-mediated toxicity. Another aspect of the interplay between astrocytes and microglial cells is associated with their known activation and proliferation in response to injury. Microglia even in minor pathological events start to proliferate (Kreutzberg, 1996), and reactive microglia are equipped to produce both components (oxygen radicals and NO) of highly toxic ONOO−, which may induce membrane perturbations, and capable to inhibit the function of membrane proteins, like glutamate transporters (Trotti et al, 1996), contributing to elevated glutamate levels and excitotoxicity. Microglial NO production – in physiological conditions – is under the negative feedback control of astrocytes by the release of transforming growth factor β (TGF-β) (Vincent et al, 1997). However, astrocytes maintain this inhibitory control only as long as they remain in their differentiated state. If they are forced to undergo pathological activation, astrocytes may aggravate oxidative neuronal damage by the activation of their own inducible NO-synthase (Schubert et al, 2001). Furthermore, non-differentiated proliferative astrocytes lack of inward rectifying chloride channels, responsible for maintaining their intact membrane potential (Ferroni et al, 1995). Thus, pathologically activated astroglial cells, reverting to their primitive state (Aschner, 1998), may loose their ability to regulate potassium homeostasis.
and also to buffer glutamate), thus, through increased depolarization of excitable cells directly contribute to excitotoxicity. Thus, considerable degree of degeneration may result from the local glial reaction next to the injured motor neurons which converts oxidative stress to calcium-mediated excitotoxic lesion (Barger et al, 2007).

Finally, an important aspect of the impairment of the BBB and BSBB is emphasized, which may lead to an increased concentration of immunoglobulins, at least locally at the lesion sites. The phenomenon possess significance in the pathomechanism of ALS, since the presence of ALS IgG indeed could be demonstrated in the brain and spinal cord of ALS patients (Engelhardt and Appel, 1990) and also in a transgenic rat model of ALS (Nicaise et al, 2009). Purified IgG from the sera of ALS patients is directed against L-type voltage gated Ca$^{2+}$-channels (Smith et al, 1992). Further experiments have also documented that ALS IgG can interact and alter the function of neuronal N-/ P-/ Q-channels, as well (Llinas et al, 1993). The interaction of ALS IgG with these calcium channels enhances inward calcium current (Mosier et al, 1995) and increases intracellular calcium (Colom et al, 1997) as was demonstrated in a motor neuron cell line.

### Expectations from increased calcium buffering capacity

Although several approaches may be feasible how to stabilize intracellular calcium homeostasis, clinical observations and harmonizing experimental data pointed to the possibility that by means of increasing the intracellular buffering capacity through overexpressing calcium binding proteins this goal could be achieved.

Historically, first, a correlation could be established, based on clinical and pathological data, that certain population of motor neurons, possessing higher level of calcium binding proteins calbindin-D$_{28k}$ (CB) and parvalbumin (PV), such as those innervating voluntary sphincter- and external oculomotor muscles do not degenerate during the disease (Ince et al, 1993; Alexianu et al, 1994). These observations culminated in a statement that “Parvalbumin is a marker of ALS-resistant motor neurons” (Elliot and Snider, 1995). On this basis, motor neuron pools were selected with relatively easy experimental manipulation possibilities, considered as “prototypes” of vulnerable- and resistant cell, like hypoglossal- and oculomotor neurons, respectively, which were further characterized with electrophysiological methods. In these studies, the correlation between the CaBP content and the ability of the motor neurons
to resist calcium-mediated injury was confirmed (Lips and Keller, 1998; Vanselow and Keller, 2000).

Next, exploiting the discovered potency of the ALS IgG to induce calcium influx, its ability to kill motor neurons was directly tested and proved \textit{in vitro}, using a motor neuron cell line (Smith et al, 1994). This (VSC 4.1) cell line provided the possibility of direct testing their resistance against ALS IgG-mediated toxicity in differentiation and transfection experiments. Undifferentiated VSC 4.1 motor neurons, while they contained calbindin-D$_{28K}$ and parvalbumin, remained resistant to the effects of IgG. Yet, after their differentiation, once they had lost these proteins, the same IgG resulted in calcium influx and caused cell death (Ho et al, 1996). If these cells were transfected with calbindin-D$_{28K}$ cDNA, the level of this calcium-binding protein remained high, even in differentiated cells and the cell remained viable when treated with ALS IgG (Ho et al, 1996).

Finally, the ability of CaBPs to lend resistance in ALS-like conditions was directly tested \textit{in vivo}, as well, in the G93A mutant SOD1 model. Indeed, qualitatively, with a combined light microscopic immunocytochemical and electron microscopic characterizations of oculomotor- and spinal motor neurons, a correlation could be established between the resistance and PV content of motor neurons, which was paralleled with an inverse correlation between the resistance and the intracellular calcium level (Siklós et al, 1998).

Thus, it seemed a reasonable goal to provide resistance of “oculomotor type” to vulnerable neurons by means of an artificial upregulation of parvalbumin expression, at least in suitably designed transgenic animal strains.

\textbf{Proof of concept experiment: protection by calcium binding in ALS}

The acid test of the protective role of calcium buffering capacity concept in ALS is the production and characterization of a model, in which ALS-like stress and increased CaBP level simultaneously exist. In an effort along this line, double transgenic mice possessing mSOD1 transgene, and concurrently exhibiting increased PV levels have been developed. Although these animals exhibited improved parameters with regard to morphological- and physiological characteristics, ultimately could not be cured, furthermore, the gain in the survival data was just above the level, to be accepted as significant (Beers et al, 2001). This
unexpected low efficacy of protection by means of upregulated calcium buffering capacity, in view of the pivotal role of calcium in cell destruction was surprising. Obviously, next, the reason(s) of the unexpectedly low benefit of this intervention was decided to be investigated, with the hope of finding ways to improve neuroprotection on the same principles, i.e. to stabilize the intracellular calcium homeostasis, if feasible. Along this line, our primary aim was to rule out possible methodological pitfalls, since some of the previous results were based on either morphological or physiological examinations of different groups of cells, and sometimes have been only partially quantified. Thus, our aims in this direction were: setup of a lesion → detection assay based on a uniform intervention (e.g. surgery, like axotomy), and a standardized characterization (e.g. on a morphological basis) of the intracellular calcium changes of the same group of cells, as well as, knowing their involvement in the injury, setup of an assay to quantify the degree of the participation of the non-motoneuronal cells in the injury.

Previous experiments relevant to the present study
Most importantly, an acute injury, i.e. axotomy/target deprivation experimental paradigm has been established, to quantify motoneuronal calcium increase in pools of motor neurons (oculomotor-, hypoglossal motor neurons) with documented different vulnerability to injury in ALS. These experiments, using electron microscopic histochemical methods and light microscopic immunohistochemistry, quantitatively documented a correlation between the resistance of motor neurons against injury and their CaBP level contrasted by an enduring increase in intracellular calcium content in CaBP-low cell types (Obál et al, 2006). Next, more importantly, these experiments provided a reference for comparisons whether an artificial increase is CaBPs could alter the profile of calcium response in these cells.

Problems to be addressed by further experiments
With the above experiment, besides establishing a reproducible experimental paradigm, we could provide additional experimental proof of the protective role of CaBPs in injury. However, since the experiment were performed using wild type animals and different pools of neurons, the question remained: whether the cellular resistance in these cells was provided by means of their increased CaBP level or local environmental factors had a decisive influence. To resolve this dilemma, the same cells with altered CaBPs have to be challenged in their
original environment, and, preferably the environmental change of the injured neurons has to be assayed, at the same time.

AIMS OF THE STUDY

- To demonstrate, whether the scale/profile of intracellular calcium increase after injury in a naturally vulnerably cell type (motor neurons in the hypoglossal nucleus) could be transformed to that of resistant motor neurons (oculomotor neurons) by PV upregulation
- To develop methods to quantify perineuronal (micro)glial reaction after injury
- To analyze the consequence of the reduced motoneuronal calcium increase on the local glial reaction after injury in PV upregulated mice

MATERIALS AND METHODS

Experimental animals

*Maintenance of the transgenic colony*

The description of the generation of the PV overexpressing (PV+/+) mice used in this study has been given elsewhere (Beers et al, 2001). Parental breeding between transgenic CaMKII-parvalbumin mice #4031x4034, 4035 obtained from Dr. S. H. Appel (The Methodist Hospital Neurological Institute, Houston, TX) was set up at the conventional animal facility of the Institute of Experimental Medicine, Budapest. All transgenic mice used in this study were derived from the original parental cross between transgenic male #4031 mated to 4034 and #4045 females of line Tg (parv 14-4). All litters obtained from this cross (mice #12-21) and subsequent sibling/sibling matings as well as the parents were homozygous for the transgene as proved by testcrosses to wild-type mice. To prove the presence of the transgene, DNA was isolated from tail biopsies and prepared according to a standard protocol published online on the Jackson Laboratory website http://www.jax.org/imr/tail_phenol.html). In general, 0.1 µg of genomic tail DNA was used to amplify the transgene using two different primer pairs: (A): primers derived from the 3d and 4th exon of the endogenous CaMKII gene as: forward 5’-CTGCAGACTCCTTCGACCAC-3’ and reverse 5’-GCCATCAGCGTCTTTGTTC-3’. The
sizes of the products were 200 bp (transgene) and 1400 bp (endogenous CamKII gene). (B) primers were: forward primer 5’-GCGAGCGAGTCGAGTGGTTGTCTG-3’ and reverse primer 5’-TTAGCTTTCGGCCACCAGAGTGGAGAATTC-3’ (Beers et al, 2001), size of the product 1200 bp. Amplification was as follows: denaturation at 94 °C for 4’, followed by 30 cycles of 30” at 94 °C, 45” at 57 °C and 45” at 72 °C, final extension 7’ at 72 °C. DNA was run on EtBr- containing agarose gels and photographed under UV light.

Fig.1. Polymerase chain reaction (PCR)-based genotyping of CamKII-parvalbumin transgenic mice. Transgenic litter from a homozygous cross between mice #4031x#4034, #4045 of line PV 14-4 (#12-21) were tailed-clipped, DNA from tail biopsies was purified and subjected to 30 rounds of PCR with two different primer pairs as detailed below. (A): primers resulting in a transgene-specific 200 bp and 1400 bp (endogenous CamKII gene) were used. This band is missing in the lanes of transgenic DNA because of highly preferential amplification of the transgene. (B): primer pair B was used resulting in a 1200 bp, transgene-specific band. All mice from this cross (Tg12-Tg21) were positive for the transgene.

Characterization of the motoneuron pools in PV+/+ mice

Besides the demonstration of the presence of the transgene in the PV+/+ animals by PCR, hypoglossal- and spinal motor neurons, which are devoid of parvalbumin in the wild type mice were checked for the presence of the PV by light microscopic immunocytochemistry.

In the hypoglossal nucleus, PV was demonstrated using the avidin biotin technique. In these sections, PV-positive motor neurons were identified according to their shape, size and distribution (Fig. 2). PV immunostaining was performed on the 30-µm thickness cryostat sections. Endogenous peroxidase activity was first blocked with H2O2, and then sections were immersed in Triton X-100 to enhance antibody penetration. Sections were pretreated with normal rabbit serum for one hour, followed by overnight incubation at 4°C with 1:60 000
dilution of polyclonal rabbit anti parvalbumin antibody (PV 28, Swant). After washing, 1:200 dilution of biotinylated goat anti rabbit secondary antibody (BA 9200, Vector) was placed on the samples and incubated for one hour. The antibodies were detected with Vectasatin Elite ABC Kit (PK-6100, Vector) and visualized by incubation in diaminobenzidine (DAB, 5% in PBS, Sigma).

Fig.2. Parvalbumin staining of the hypoglossal of PV+/+ (A) and B6/SJL (B) mice. In the hypoglossal nucleus of the PV+/+ animals, numerous stained cellular profiles are visible which cells, according to their size, shape and distribution can be considered as motor neurons (A). In the hypoglossal nucleus of the B6/SJL mice only background staining can be seen (B). Some stained cellular profiles at the ventro-medial and lateral border of the hypoglossal nucleus in both figures may belong to the nucleus of Roller and the intercalated nucleus of medulla, respectively. mlf: medial longitudinal fasciculus. Bar: 100 μm.

In the spinal cord of PV+/+ mice, double immunostaining, for cholinacetyltransferase (ChAT) and PV was performed, also using 30-μm thick cryostat sections. Endogenous peroxidase
activity was first blocked with H2O2, then sections were pretreated with normal donkey serum for one hour, followed by overnight incubation at 4 °C with a primary antibody cocktail, containing the same polyclonal primary antibody against PV as above (1:60 000) and a polyclonal goat anti ChAT antibody (Chemicon, AB1582; 1:250). After washing, a cocktail of Alexa Fluor 546 (anti rabbit) and Alexa Fluor 488 (anti goat) fluorescent secondary antibodies were placed (1:200) and incubated for one hour at room temperature. Sections were washed in PBS, mounted on silane-coated glass slides, covered with Gel/Mount (Biomed) and visualized under a laser confocal microscope (Olympus FV1000) (Fig. 3).

![Image](image_url)

**Fig. 3.** PV-positive motor neurons at the ventro-lateral region of the spinal cord in PV+/+ mice were identified according to a colocalized staining with PV (a) and ChAT (b). Bar: 30 µm.

**Axotomy/target deprivation of hypoglossal and oculomotor neurons**

Twelve weeks old male B6/SJL mice (BRC, Szeged, Hungary), with breeding pairs originally obtained from Charles River Laboratories (Budapest, Hungary), and PV+/+ mice (IEM, Budapest; BRC, Szeged) were selected for the study. To avoid multiple survival surgery, animals were assigned either for hypoglossal or oculomotor nerve cut experiments. A total of 60 mice were used either for hypoglossal axotomy (n=40) or for oculomotor surgery (n=20). Four animals were euthanized at 5 time points for both types of surgery and were processed for electron microscopic calcium histochemistry study. Surgical procedures were performed under deep anesthesia with Avertin (tribromoethanol; Fluka; 240 mg/kg body weight in a 0.3
ml volume) administered intraperitoneally. The hypoglossal nerve was transected lateral to the hyoid bone and a piece of 3-5 mm nerve segment was removed to prevent regeneration. For transection of left oculomotor nerve, animals were enucleated and the orbits were cleared of the remaining extraocular muscles and nerve segments.

Axotomy of the sciatic nerve

Twelve weeks old male B6/SJL, Balb/C mice (BRC, Szeged) and PV+/+ mice were selected for the study. Animals were assigned either for electron microscopic calcium detection study, or to light microscopic immunocytochemistry, or functional assay study. Surgical procedures were performed under deep anesthesia, as described above. The sciatic nerve was transected at the mid-thigh, and a small piece (3-5 mm) was removed to prevent reinnervation (Fig. 4). In these and in the above experiments the non-operated side served as an internal control. Following surgery animals were housed in a temperature controlled room with a 12 h light/dark cycle, with access to water and food ad libitum, and were allowed to survive for 1, 4, 7, 14 and 21 postoperative days. All the experiments were performed in accordance with the institutional guidelines for the use and care of experimental animals and the appropriate governmental laws for animal protection (protocol No. 72-45/b/2001 and No. 03876/0014/2006).

Electron microscopic detection of calcium

Specimen preparation

Mice anesthetized irreversibly with methoxyflurane were perfused transcardially, first with 90 mM potassium oxalate (Sigma) and then with 3% glutaraldehyde (Polysciences) containing 90 mM potassium oxalate (pH 7.4) (Borgers et al, 1977, 1981). Tissue blocks containing either the nuclei of cranial nerves III or XII, or the lumbar segment of the spinal cord were dissected and fixed overnight in the same fixative (4 °C). Specimens were rinsed in 7.5%
sucrose in 90 mM potassium oxalate (pH 7.4), and then postfixed for 2 h (4 °C) in 2% potassium pyroantimonate (Merck) and 1% osmic acid (Sigma), adjusted to pH 7.4 with acetic acid (Molar). Following postfixation, specimens were rinsed in distilled water (pH adjusted to 10 with KOH) for 10 minutes, dehydrated in graded series of ethanol, processed through propylene oxide (Sigma) and embedded in plastic (Durcupan ACM, Fluka). Blocks were polymerized for 2 days at 56 °C.

Semithin (0.3-μm) sections were cut from the blocks on an Ultracut UCT ultramicrotome (Leica), etched (Maxwell, 1978), stained (Richardson et al, 1960) and screened under the light microscope (Olympus Vanox-T) to identify the borders of nuclei III, XII and lamina IX in the spinal motor neurons. After trimming, 50-nm ultrathin sections were obtained, mounted on formvar-coated single-hole copper grids, contrasted with uranyl acetate and lead citrate and analyzed in the electron microscope (Zeiss CEM 902). The procedure resulted in electron-dense deposits (EDDs) due to the precipitation of tissue calcium by the fixative, which was easily recognized by conventional transmission electron microscopy.

Specificity control of the histochemical reaction for calcium

Time-to-time, the specificity of the fixation procedure for calcium was checked by electron spectroscopic imaging (ESI) (Bauer, 1988; Reimer et al, 1988). For such analysis, 15-25-nm sections were prepared, mounted on uncoated 700-mesh copper grids and without contrasting, examined under the electron microscope in ESI mode at Δe=250 eV, which setting was used for structural imaging, as well (Körtje and Körtje, 1992). Such images, resembling “dark-field” images, were then inverted for easier comparison. The analytical image displaying the net calcium distribution was obtained by subtracting the background image (Δe=310 eV) from the “edge” image (Δe=355 eV). The non-significant signal below the [mean + 2.5 × s.d.] level was deleted in the analytical image, and then, for easier comparison of the co-distribution of the EDDs with the true calcium, the significant calcium signal was false-colored according to the significance level in the [mean + 2.5 × s.d.] to [mean + 6 × s.d.] interval. This final image, representing the distribution of the precipitated calcium in the tissue, was compared visually with micrographs obtained by structural imaging of the same area, displaying the arrangement of the EDDs (Fig 5).
Fig. 5. Electron spectroscopic imaging (ESI) of electron-dense deposits (EDDs) of the histochemical reaction. A: Fine structural image of a typical intra-mitochondrial cluster of EDDs. B: The same area was also analyzed by ESI set to produce the net distribution of calcium at the significance level of <background + 2.5 s.d.>. The comparison of (A) with (B) demonstrates a good correlation of the EDDs with the tissue calcium.

**Semiquantitative evaluation of the intracellular amount of EDDs**

Ultrathin sections were screened under the electron microscope at low magnification (3 000 ×) to identify motor neurons, which were selected randomly for measurements. One region from each cell body, also selected at random, was photographed at a magnification of 12 000 ×, printed at 2.7 × enlargement, until 12-15 samples had been collected from both the operated and the non-operated sides in each animal. The amounts of calcium were determined by measuring the relative volume of the EDDs with respect to selected reference volumes. Three types of reference spaces were selected: either the whole perikaryal volume, or the mitochondrial, or the extra-mitochondrial compartments of the perikaryal space. The three-dimensional parameters were determined with a point counting technique (Mayhew, 1992). After random superimposition of a tessellation of sampling points on the prints, grid points overlying either the EDDs \( P_{\text{EDD(cyt)}} \) or \( P_{\text{EDD(mit)}} \) or the corresponding reference area \( P_{\text{(cyt)}} \) or \( P_{\text{(mit)}} \) were counted. The scores obtained on individual micrographs were pooled according to the operated and non-operated sides and the corresponding ratios expressing the “histochemical concentration of calcium” in the mitochondrial and cytoplasmic compartments were calculated for both sides \( \frac{\sum P_{\text{EDD(cyt)}}}{\sum P_{\text{(cyt)}}} \) and \( \frac{\sum P_{\text{EDD(mit)}}}{\sum P_{\text{(mit)}}} \). Finally, the
corresponding values obtained on the operated side were divided by those obtained on the non-operated side to express the changes (fold increase) induced by the axotomy in the mitochondrial and cytoplasmic calcium contents for each animal. The fold increases in calcium in the hypoglossal and oculomotor nuclei of the PV+/+ animals and in the hypoglossal nucleus of the B6/SJL mice were compared on postoperative days 1, 4, 7, 14 and 21 statistically (two-way ANOVA with Duncan post-hoc comparison; StatSoft Statistica). In the spinal cord, after sciatic axotomy, the fold-increase of calcium was determined only at postoperative day 7, when the glial reaction was at its maximum. The difference between the strains was compared statistically (one-way ANOVA with Duncan post-hoc comparison; StatSoft Statistica).

**Immunohistochemical characterization of inflammatory markers in the spinal cord**

*CD11b staining*

Microglial activation after axotomy was followed by CD11b-staining using light microscopic immuycytochemistry. Mice under irreversible anesthesia were transcardially perfused with 10 mM PBS, followed by 3% paraformaldehyde in 10 mM PBS. The spinal cords were removed and immersed in the same fixative for 24 h (4 °C). To prevent the formation of freezing artifacts, tissue samples were placed in sucrose (30% in 10 mM PBS, pH 7.4) for at least 2 days at 4 °C. Sections with a nominal thickness of 30 µm were then cut on a cryostat at -18 °C, and collected in 10 mM PBS. To detect microglial cells with an IHC procedure, free-floating sections were stained according to the avidin-biotin technique (Cuello, 1993). Sections were rinsed first with 3 changes in PBS, which was followed by a 30-minute blockade of endogenous peroxidase activity with 0.3% hydrogen peroxide in 10 mM PBS containing 0.2% Triton X-100 (Sigma) (TPBS). Then, after washing (3 changes in PBS), to reduce the nonspecific staining, sections were incubated in 2% normal goat serum (Vector) in TPBS for 1 h. Microglial cells were labeled with anti-CD11b antibody (1:500 in TPBS; rat-anti-mouse antibody, Serotec) during incubation at 4°C overnight. After washing in PBS (3 changes), sections were incubated in biotin-labeled secondary antibody for 1 h (1:800 in TPBS, goat-anti-rat IgG, Vector), followed by washing in PBS (3 changes), and incubation in the avidin-biotin complex (1:1600 in PBS, Vector) for 1 h, at room temperature. Finally, after washing in PBS, the reaction was visualized by incubation in DAB (5% in PBS; Sigma)
for 15 min. Stained sections were thoroughly rinsed in PBS, mounted on silane-coated glass slides, dehydrated in graded series of ethanol, processed through xylane (Molar) and coverslipped with Entellan (Merck).

**MCP1 staining**

The immunostaining to detect the chemokine MCP-1 (monocyte chemotactic protein-1) was performed similarly, as above on free-floating 30-µm thickness cryostat sections, using the avidin-biotin method adapted to the primary antibody: polyclonal rabbit anti-murine JE antibody (PeproTech, 1:1000).

**Evaluation of the functional recovery after sciatic nerve cut**

Additional, twelve week old, male PV+/+, B6/SJL and Balb/C mice (4 animals in each group) were allocated for the study. Following sciatic nerve cut, animals were housed in a temperature controlled room with a 12h light/dark cycle, and had access to water and food *ad libitum*. The function of the operated hindlimb of animals was characterized daily at the same time. An arbitrary, but systematically applied scale was used to assess the recovery of the hindlimb function (Table 1).

<table>
<thead>
<tr>
<th>Grade</th>
<th>Definition</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Practically no function on the operated side, the animal does not rear</td>
</tr>
<tr>
<td>2</td>
<td>Drags paralyzed hindlimb, the overall behavior, mobility is sluggish</td>
</tr>
<tr>
<td>3</td>
<td>Occasionally uses operated limb, increased mobility, cannot climb on the grid</td>
</tr>
<tr>
<td>4</td>
<td>Walks actively using both hindlimbs, still cannot bend fingers on the operated side</td>
</tr>
<tr>
<td>5</td>
<td>Tries to bend fingers during walking, but cannot grasp grid during climbing</td>
</tr>
<tr>
<td>6</td>
<td>Time-to-time bends fingers during walking but still cannot do that during rearing</td>
</tr>
<tr>
<td>7</td>
<td>Walks normally but easily tired, still cannot grasp grid with the fingers</td>
</tr>
<tr>
<td>8</td>
<td>Normal walking, increased ability to grasp the grid with the fingers</td>
</tr>
<tr>
<td>9</td>
<td>Normal walking, only occasional problems during rearing and grasping the grid</td>
</tr>
<tr>
<td>10</td>
<td>Full recovery</td>
</tr>
</tbody>
</table>

The subjectivity in grading the hindlimb function was tried to be reduced by evaluating the animals in the different groups in a single session in each day by pinpointing even slight relative differences in the behavior. Furthermore, all the evaluation processes as well as the daily animal care during the 28-day course of the study was performed by the same person.
RESULTS

Calcium increase is reduced after injury in hypoglossal motor neurons of PV+/+ mice

Fig.6. Calcium in the hypoglossal motor neurons of B6/SJL mice. A: non-operated side, B-F: operated side at postoperative day 1, 4, 7, 14 and 21. Go: Golgi apparatus, mit: mitochondrion, arrow: EDD. Bar: 1 μm.
Fig. 7. Reduced calcium increase after axotomy in the hypoglossal motor neurons of PV+/+ mice compared to the B6/SJL strain. A: non-operated side, B-F: operated side at postoperative day 1, 4, 7, 14 and 21, respectively. Go: Golgi apparatus, mit: mitochondrion, arrow: EDD. Bar: 1 μm.
Qualitatively a gradual elevation of the intracellular calcium content in the axotomized hypoglossal motor neurons of the B6/SJL mice was seen with increasing postoperative time relative to the non-operated side, with a tendency to return to the control level by postoperative day 21 (Fig. 6). In the hypoglossal neurons of the PV+/+ animals, however, except for a slight increase in the mitochondrial calcium content on postoperative day 1, the calcium content was observed not to be elevated relative to the control side on any postoperative day (Fig. 7). In the oculomotor neurons of the PV+/+ mice, similarly as in the hypoglossal neurons in these animals, no calcium increase was detected in the injured neurons as compared with the intact cells at any time point after the surgery (not shown).

In both motor nuclei, the volume fraction of the EDDs, characterizing the distribution of calcium in the tissue, was expressed quantitatively. Statistically, the time dependences of the calcium levels in the two nuclei in the PV+/+ strain did not differ, regardless of whether the total calcium (p=0.21), or the cytoplasmic (p=0.43) and mitochondrial (p=0.20) calcium levels separately were compared (Fig. 8). However, the postoperative calcium increase in the motor neurons from the B6/SJL mice was significantly different from that of the PV+/+ animals (p=0.001).

Fig. 8. Quantification of the amount of EDDs in the perikaryal volume (A), in the mitochondria (B) and in the cytoplasm (C) of the hypoglossal motor neurons of B6/SJL (white bars) and PV+/+ mice (black bars) and of the oculomotor neurons of PV+/+ animals (gray bars). Data are expressed as the fold increases relative to the non-operated side on days 1, 4, 7, 14 and 21 postoperatively. The overall calcium increase in the motor neurons from the B6/SJL mice was significantly different from that for the PV+/+ animals (p=0.001, two-way ANOVA) with slight differences at day 4, 7 and 14 postoperative times in the different cellular compartments.
These data provide evidence for that – with regard to the calcium handling after injury – an increased calcium buffer may confer an “oculomotor-like resistant phenotype” to the naturally vulnerable hypoglossal motor neurons against injury. For more details see Paizs et al. (2010a).

**Inflammatory reaction can be quantified by systematic sampling and background corrected segmentation using immunohistochemistry**

Inflammatory responses in the CNS are commonly assayed by the reaction of the immune-competent microglial cells in the CNS, characterized by IHC. Obtaining reliable conclusions on the basis of IHC staining is hindered by the regular need of subjective judgments, due inconsistencies of the staining intensity from section-to-section or in repeated experiments. Our method, utilizing digital image analysis, is based on the use of an internal reference area on the analyzed sections, which makes the relative changes of IHC staining in different experiments comparable, independent of the examiner (Fig. 9). Another source of variability of staining could be attributed to the internal heterogeneity of the object to be characterized, which means that identical fields could never be analyzed. To compensate also for this variability, a systematic random sampling paradigm was designed, which provides data, expressed in single numbers, describing the dimension and strength of IHC staining in the entire volume to be characterized (Fig. 10). In this integrated approach, the figures are derived by pooling relative IHC staining intensities from all of the sections of the series from a particular animal. The procedure (1) eliminates the problem due to the personal assessment of the significance of IHC staining intensity, (2) does not depend on the precise dissection of the tissue on a gross scale, i.e. considerably reduces the consequences of a limited, arbitrary sampling of the region of interest, and finally (3) arrives at quantitative data characterizing the extent of inflammation in the animal. The procedure has been validated in a well documented experimental paradigm, selected to be analogous to our intended experimental plan. Thus, as a positive test, inflammatory reactions in the spinal cord, measured as microglial activation, were followed by IHC after axotomy of the sciatic nerve in mice, and, then, the effect of a known anti-inflammatory compound (minocycline) has been tested. These measurements proved that our quantification procedure is capable to measure inflammatory reactions light microscopically and sensitive enough to demonstrate the efficacy of anti-inflammatory interventions (Table 2). For more details see Paizs et al. (2009).
Fig. 9. Approximate area of the expected reaction is marked on the spinal cord sections (a: AOI), then mirrored to the non-operated (unaffected) side (b: AOI'). The background intensity has been determined using the unaffected side as reference (c-f), then the significant signal has been segmented automatically (g, h).
Fig.10. Expected dimension of the microglial reaction in the spinal cord after the lesion of the sciatic nerve (a: red shaded volume), which is assayed by systematic sectioning (at 300 µm) started at a random (x) position (b).

With this integrated sampling/calculation procedure the anti-inflammatory effect of minocycline could be demonstrated (Table 2; p < 0.05). Similar significant results could be obtained either by measuring the pooled relative staining intensity or the pooled relative area density covered by the stained cells (Table 2).

<table>
<thead>
<tr>
<th>animals</th>
<th>operated only</th>
<th>operated + minocycline</th>
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<tbody>
<tr>
<td></td>
<td>poled relative stained area [%]</td>
<td>pooled intensities [−ln(I(op)/I(co))]</td>
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<tr>
<td>#1</td>
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<td>1.67</td>
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<tr>
<td>#2</td>
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</tr>
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<td>#6</td>
<td>657.43</td>
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</tr>
<tr>
<td>mean</td>
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</tr>
<tr>
<td>S.E.M.</td>
<td>49.01</td>
<td>0.66</td>
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</table>

**Increased motoneuronal calcium buffer reduces local inflammation after acute lesion**

The relationship between the increased calcium buffering capacity of motor neurons and the altered local inflammatory reaction has been examined in the spinal cord of PV+/+ and control (B6/SJL and Balb/c) mice after the lesion of the sciatic nerve. The temporal change of the inflammatory markers has been followed in 21 days after the operation with IHC (total of 30 mice; 2 in each group) supplemented with the characterization of the recovery of the
hindlimb function (4 mice in each group). Calcium level of the motor neurons was determined at day 7, when the inflammatory reaction peaked (6 mice in each group).

Resting microglial cells (Fig. 11a) surrounding motor neurons, after the injury of these cells become activated, change their morphology: withdraw their processes, which, as well as their cell bodies turn to be hypertrophic (Fig. 11c). At the same time, the injured motor neurons release factors, such as MCP1 (Fig. 11b), attracting microglia to the site of injury.

![Fig. 11. Resting microglial cells (a) become activated (c; b: green color) around motor neurons after injury; motor neurons release chemokine MCP1 (b; red color) serving as a chemotactic signal.](image)

Indeed, by regular inspection of the whole cross sections of the spinal cords of the operated animals it could be documented, that both the microglial activation/reaction (except for the input region of the sensory fibers; c.f. Fig. 4) and the MCP1 release was restricted to, and around the ventro-lateral pool of motor neurons (Fig. 12).

![Fig. 12. MCP1 staining (a) and increased CD11b staining (b) is confined to the area occupied by the injured motor neurons. Microglial activation at the input of the sensory fibers has not been evaluated in this study.](image)
The inflammatory reaction commenced as early as day 1 after operation in the spinal cord, either the MCP1 chemotactic signal, or the altered CD11b expression is considered. These reactions lasted at least 3 weeks long, however with a different decay time: the decrease of the release of the MCP1 by the injured motor neurons preceded the attenuation of the microglial reaction at their vicinity (Fig. 13). Nevertheless, the decrease of both of these inflammatory markers was significantly faster in the PV+/+ mice compared to the control strains (Fig. 13).

**Fig. 13.** A difference has been demonstrated between the decay of the expression of the chemotactic signal, MCP1 (a), released by the injured motor neurons, and the CD11b (b) staining, characterizing the microglial reaction after axotomy. Most importantly the MCP1 release of the motor neurons from the PV+/+ mice (claret colored bars) returned to close to the baseline at day 14 after operation, significantly (*: p<0.008) faster than in the control mouse lines (blue: B6/SJL; yellow: Balb/c), followed by a similarly faster decrease of the microglial reaction, demonstrated at day 21 after the operation (**: p<0.04).

An important result of the present experiments is – similar to the hypoglossal motor neurons of the PV+/+ mice – that the PV upregulation attenuated the increase of the intracellular calcium after axotomy in the spinal motor neurons, as well. The calcium measurements have been performed at postoperative day 7, at the time when the calcium increase in the wild type hypoglossal neurons was significantly higher than in those of the PV+/+ mice. Thus, without performing a complete time series
analysis, it can be assumed, that the PV upregulation lends the same “oculomotor-type” resistance to the spinal motor neurons as to the hypoglossal neurons, which is most likely paralleled with a reduction of the calcium mediated injury. In accord with this assumption, the expression of the distress signal, represented by the chemokine MCP1, disappears in the PV+/+ motor neurons faster than in the control strains, followed by a similarly faster elimination of the microglial activation at the vicinity of these cells (Fig. 13). Thus, the reduction of calcium increase in motor neurons is indeed capable of reducing local inflammation, furthermore, according to the observed rate of improvement of the hindlimb function in these animals in the same period, leads to a faster functional recovery (Fig. 15).

![Graph](image)

Fig.15. Recovery of the function of the operated hindlimb of PV+/+ (claret color), B6/SJL (blue) and Balb/c (yellow) mice. The recovery was estimated on a 1-10 (10-100%) scale, as described in Table 1. Data are expressed as mean ± s.e.m.

With regard to the rate of recovery, two further comments can be made: (1) even in an extended period of observation after the surgery (4 weeks) no full recovery could be noticed in either strains, (2) although no daily data are available about the microglial activation, the increase in the rate of recovery in the PV+/+ mice was in the same timeframe as the significant decrease of microglial activation compared to the control strains (cf. Fig. 15 and Fig. 13).
DISCUSSION

The role of calcium and CaBPs in neuronal degeneration

It is well documented, that abnormal and sustained increases in intracellular calcium induce a series of calcium-dependent enzymatic processes – not only in motor neurons – which participate in cellular destruction. Certainly, several catabolic enzymes, including protein kinase C, calpains, phospholipase A₂, nitric oxide synthase and endonucleases are directly involved (see e.g. Berridge, 1998; Berridge et al, 1998). Furthermore, excess calcium ions facilitate other destructive processes as well (see e.g. Marambaud et al, 2009). However, it has been debated for a long time whether the calcium buffering capacity plays a role in modifying the outcome of these processes.

CaBPs within the cells are commonly located at critical positions: in the vicinity of plasmalemmal calcium pumps and channels, around the mitochondria and endoplasmic reticulum, organelles known to play key roles in Ca²⁺ signaling and buffering (Brini, 2003). Thus, these proteins may slow down the diffusion of calcium ions away from the influx regions, or facilitate their recycling into internal stores, and thereby reduce the amplitude of calcium concentration changes in the cytosol (Müller et al, 2007; Vanselow and Keller, 2000; von Lewinski and Keller, 2005). The concept that CaBPs may exert a general neuroprotective effect was proposed on the basis of data acquired from a wide range of in vitro (Clementi et al, 1996; Mattson et al, 1991; McMahon et al, 1998) and in vivo experiments (Beck et al, 1994; Rami et al, 1992; Tortosa and Ferrer, 1993). The idea has received additional support from a variety of clinical observations, e.g. the age-dependent loss of CB from the cholinergic neurons of the basal forebrain in humans displayed a similar pattern to that of the loss of these cells in Alzheimer’s disease (Geula et al, 2003), while in Huntington’s disease those striatal neuronal groups which contained high levels of CB proved to have a decreased vulnerability (Mitchell and Griffiths, 2003). In Parkinson’s disease, the CB-positive dopaminergic neurons in the substantia nigra are relatively spared. Finally, in ALS, the most susceptible motor neurons are comparatively deficient in CB and/or PV (Alexianu et al, 1994; Ince et al, 1993).

However, not all studies lend support to the protective role of the CaBPs during injury. The reports range from documentation of the inability of a high CaBP content to confer resistance on the (Purkinje) cells in the cerebellum (Slemmer et al, 2005), through the lack of a
correlation between the CB and/or PV content and the neuronal vulnerability in the hippocampus (Freund et al, 1990), to the finding that PV actually enhances NMDA-mediated toxicity (Hartley et al, 1996), or to the notion that not an excess, but rather a deficiency in CB might be protective (Klapstein et al, 1998). The diversity of these results, however, might be attributed to the fact that the actual impact of the CaBPs on survival is affected by the local cellular environment, and influenced by the neuronal network properties of the injured cells. In an effort to eliminate such perturbations of the results, in the present study, we decided to assay the changes in calcium level after injury in motor neurons with an artificially elevated PV content, and to compare the results obtained from the same population of neurons with a naturally low PV content in wild-type animals.

**Lesion induced by axotomy**

Axonal transection is an experimental method for the study of neuronal responses to injury leading to degeneration, or resulting in plastic changes and recovery (Price et al, 1992), some of which resemble those seen in ALS (Price et al, 1994). Factors known to influence the outcome of axotomy-induced injury are the type of lesion (crush, cut, target deprivation or avulsion), the location of the trauma relative to the perikaryon, differences in species, type of the motor neurons and, most importantly, age of the animals (Koliatsos and Price, 1996). For example, approximately 70% of adult hypoglossal motor neurons survive target deprivation in rats, but only 40% survive if the surgery is made during the first postnatal week (Snider and Thanedar, 1989). Similarly, in the spinal cord of newborn animals, while avulsion or crush of the sciatic nerve produces a 73-80% loss of lumbar motor neurons (Koliatsos and Price, 1996), axotomy of the same nerve in adult animals generates no visible loss of motor neurons (Schmalbruch, 1984). Thus, in this study, to evoke sublethal injury, we performed the surgical interventions in animals 12 weeks of age, when the adult response was already present.

**Calcium level changes induced by axotomy**

The lesion of the hypoglossal nerve of the B6/SJL animals was followed by a transient increase in the intracellular calcium in the corresponding motor neurons, with a peak increase of approximately 1.5-fold around day 7-14. In the same hypoglossal motor neurons of the PV+/+ animals, apart from a slight, insignificant increase in the mitochondrial calcium on day 1 after surgery, no increase was detected relative to the calcium level in the motor neurons on the non-operated side at any time point postoperatively. Likewise, no calcium increase was
seen in the motor neurons of the oculomotor nuclei of the PV+/+ mice after axotomy, and no significant difference could be discerned statistically between the two examined motor nuclei of the PV+/+ animals. Based on these results, in the spinal cord, only one time point, postoperative day 7 was selected to check the calcium increase, when the maximum change in the calcium level could be expected in the motor neurons of the wild type animals. In spinal motor neurons of the PV+/+ mice, similarly to the hypoglossal neurons, no significant calcium increase could be demonstrated, while in the same motor neurons from both control strains a 1.5-2-fold calcium increase, comparable to that of the wild type hypoglossal motor neurons could be demonstrated.

In our earlier study, to compare the calcium response in the hypoglossal and oculomotor neurons of wild type animals, the Balb/c mouse strain was used, and an increased resistance of the oculomotor neurons was documented (Obá et al, 2006). Those experiments left open the question whether the differences in the cellular environment could have contributed to the different resistance. Since, in the present study, we could document a similarity between the responses to axotomy of the same (spinal) motor neurons from two control strains (Balb/c mice and the B6/SJL parental mice to the PV+/+ animals), as well as a similarity between the responses to axotomy of different (hypoglossal- and spinal) motor neurons from the same (PV+/+) strain, we may assume, that neither the diversity in the strains, nor the difference between the cellular environment, but the changed calcium buffering capacity is indeed responsible for the contrasting resistance of these cells, at least in acute lesion paradigms.

The role of inflammatory factors in injury of motor neurons
Microglial activation and proliferation is a nearly stereotypical response to any insult to neuronal tissue. Resting microglial cells normally sense neuronal damage, readily transform into an activated state (undergo morphologic changes of process retraction and cell body enlargement as well as functional changes of increased expression of immune/inflammatory markers) and respond to remove damaged cells by phagocytosis (Kreutzberg, 1996). Microglia may exert both positive and negative effects in the context of neuronal lesion (Kempermann and Neumann, 2003). The positive aspect, besides the removal of the potentially damaging debris left behind following neuronal death (i.e., macrophage function) may involve the production of certain growth factors. However, activated microglia can also
cause neuronal damage, even cell death, through release of a number of cytotoxic mediators, including cytokines, free (oxygen) radicals, nitric oxide etc.

In the present experiments we documented that increasing the calcium buffering capacity in motor neurons reduces the increase of intracellular calcium level after acute injury (Fig. 14), and anticipated that such a way the calcium mediated injurious processes could be attenuated, as well. In agreement with this assumption, the duration of the expression of the chemokine MCP1 by the motor neurons (considered as a distress signal) has been considerably and significantly reduced: it was confined to 14 days after operation, while at two weeks this signal in the control mouse strain was still significantly high (Fig. 13). Following the shorter emission of MCP1 in the PV+/+ mice, the microglial activation started to turn to the baseline also significantly earlier than in the control mouse strains (Fig. 13). In addition to the reduced inflammatory reaction of PV+/+ mice, a further support for the protective role of PV-upregulation in motor neurons was provided by the faster functional recovery data in these mice (Fig. 15). These results together provide evidence for the protective role of PV in acute injury and suggest that the inflammatory reaction might be graded after injury, determined by the magnitude (and/or duration) of the primary lesion.

**Quantification of the inflammatory markers by light microscopic IHC**

As a byproduct of (or necessity of our experiments) was the development of a reproducible, quantitative light microscopic procedure to assay the magnitude of inflammatory reaction after the neuronal lesion. The procedure practically eliminates the subjectivity of the operator in assessment of the significance (degree of positivity) of the immune-staining, and compensates for a frequently neglected effect of the so-called “random sampling” as illustrated in Fig. 16. Clearly, even using internal standards in the section, the calculated intensity (characterizing microglial activation) strongly depends on the position of the sampling, which cannot be easily reproduced. Our method,
bracketing the whole spinal cord with the expected region for analysis, and sampling it systematically, integrates the numbers obtained from the individual sections, thus considerably reduces the variability of the results. As a consequence, we could achieve a precision high enough for our conclusions using only two animals at each time point to record the temporal dependence of the inflammatory reaction.

Relevance of the results in relation to the mechanisms of ALS

The suggestion that CaBPs may protect motor neurons in ALS was based on the parallelism observed between the relatively spared motor functions in patients, such as the eye movement and voluntary sphincter functions. The most widely used model of ALS suitable for in vivo research is a variety of transgenic mice developed on the basis of mutations of the enzyme SOD1 detected in a subpopulation of ALS patients (Deng et al, 1993; Gurney et al, 1994; Rosen et al, 1993). Using transgenic animals based on the G93A mutation of SOD1, we earlier demonstrated not only that changes in the intracellular calcium level paralleled the degeneration of the motor neurons, but also that the survival of the motor neurons was associated with higher levels of CB and/or PV (Siklós et al, 1998). Since those experiments still involved the comparison of different motor neurons, (spinal versus oculomotor neurons), it was decided to develop a mouse line with an elevated PV content, suitable for measurement the protective effect of CaBPs in the same neuronal population in vivo (Beers et al, 2001).

When PV+/+ transgenic mice were bred to the mutant SOD1 animals, a significantly increased preservation of the spinal motor neurons and a delayed onset of the disease was noted as compared with the mice containing only the SOD1 transgene. Although the survival of the double transgenic mice was also improved, the animals could not be ultimately rescued. To test directly whether an elevated PV level attenuates the calcium increases in the motor neurons, in the present study, PV+/+ mice were subjected to axotomy, which is accepted as a standard model for the induction of an acute sublethal injury in adult-type neurons (Koliatsos and Price, 1996). As expected, a non-significant calcium level increase was demonstrated in the hypoglossal and spinal motor neurons of PV+/+ mice, which was similar to that measured in the oculomotor neurons. Thus, the lack of success in preventing the death of mutant SOD1 animals by the upregulation of PV in the motor neurons could be a consequence of the CaBPs acting solely as calcium buffers, which could be saturated if the stress signal endures (Siklós and Appel, 2005). This is in accord with the observation that the sphincter functions are also
impaired in the patients, just only at the late phase of the disease (Bergmann et al, 1995). On the basis of this concept, with increasing the calcium buffering capacity, the maximum goal would be a delay of the injury of the vulnerable motor neurons and a provision of a neuroprotection similar to that present in the relatively resistant motor neurons. Alternatively, we may assume that the neighboring cells contribute to the damage of the motor neurons (Boillée et al, 2006; Julien, 2007; Obál et al, 2001). In the present study we demonstrated that however, at least in acute injury, the magnitude of the primary lesion of the motor neurons is that which determines the extent/duration of the inflammatory reaction, thus it seems still a reasonable aim to concentrate the protection mostly on motor neurons, but more effectively than the improved calcium buffering capacity can afford. In this respect our current experiments are aimed at characterizing the effect on the calcium level and survival of motor neurons of antagonists of calcium channels (AMPA receptors), which are relevant in the pathomechanism of ALS in the G93A transgenic model of ALS, i.e. during chronic stress (Paizs et al, 2010b).

Finally, we wish to think that our results about the disturbed calcium levels in the motor neurons of ALS patients and in the motor neurons of certain ALS models contributed significantly to the determination of starting a clinical trial with Talampanel, an AMPA-receptor antagonist (Pascuzzi et al, 2009).
SUMMARY

- In hypoglossal motor neurons of B6/SJL mice (parental line of the PV+/+ animals), as large as 1.7-fold calcium increase after axotomy was demonstrated, compared to the non-operated conditions, which was significantly different from the calcium increase in the oculomotor neurons of the same animals.

- In the PV+/+ mice there was no calcium increase either in the hypoglossal- or oculomotor neurons, providing evidence for that PV upregulation lends an “oculomotor-like” resistance.

- In the spinal motor neurons of PV+/+ mice a similarly reduced calcium level was demonstrated after injury as in the hypoglossal motor neurons, suggesting that independent of the local neuronal environment, PV upregulation indeed contributes to the resistance against calcium increase.

- In the spinal cord of the PV+/+ mice a reduced and a faster resolving inflammatory reaction could be demonstrated after acute injury compared to the B6/SJL mice. These observations, together with the reduced calcium increase in PV+/+ spinal motor neurons suggest that the magnitude of the primary lesion in the motor neurons determines the surrounding microclial activation, and stabilization of the calcium homeostasis in motor neurons may be indeed protective.

- With regard to the inflammatory reactions, a computerized method has been developed, based on light microscopic IHC and image analysis, which is able to reproducibly quantify the expression of certain inflammatory markers with high precision. The method has a general applicability.

The results confirm that elevated intracellular calcium buffer could attenuate both the *intracellular calcium increase (thus calcium-mediated degenerative processes)* and the *local inflammatory reactions* after acute injury, but leaves open the possibility that such a buffer may exhaust if the stress endures. Under chronic stress conditions, such as in animals with pathogenic mutations, which start working as early as *in utero*, an alternative way of reducing calcium levels, based e.g. on special ion channel blockers, is suggested.
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APPENDIX

Reprints of the publications providing the basis of the thesis

I.
Melinda Paizs, József I. Engelhardt, László Siklós:
Quantitative assessment of relative changes of immunohistochemical staining by light microscopy in specified anatomical regions.

II.
Melinda Paizs, József I. Engelhardt, Zója Katarova, László Siklós:
Hypoglossal motor neurons display a reduced calcium increase after axotomy in mice with upregulated parvalbumin.
Quantitative assessment of relative changes of immunohistochemical staining by light microscopy in specified anatomical regions

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Key words. Enzyme-based method, image analysis, immunohistochemistry, light microscopy, quantification, sampling paradigm.

Summary

Despite the advent of ever newer microscopic techniques for the study of the distribution of macromolecules in biological tissues, the enzyme-based immunohistochemical (IHC) methods are still used widely and routinely. However, the acquisition of reliable conclusions from the pattern of the reaction products of IHC procedures is hindered by the regular need for subjective judgments, in view of frequent inconsistencies in staining intensity from section to section or in repeated experiments. Consequently, when numerical comparisons are required, light microscopic morphological descriptions are commonly supplemented with analytical data (e.g. from Western blot analyses); however, these cannot be directly associated with accurate structural information and can easily be contaminated with data from outside the region of interest. Alternatively, to eliminate the more or less subjective evaluation of the results of IHC staining, procedures should be developed that correct for the variability of staining through the use of objective criteria. This paper describes a simple procedure, based on digital image analysis methods and the use of an internal reference area on the analyzed sections, that reduces the operator input and hence subjectivity, and makes the relative changes in IHC staining intensity in different experiments comparable. The reference area is situated at a position of the section that is not affected by the experimental treatment, or a disease condition, and that can therefore be used to specify the baseline of the IHC staining. Another source of staining variability is the internal heterogeneity of the object to be characterized, which means that identical fields can never be analyzed. To compensate for this variability, details are given of a systematic random sampling paradigm, which provides simple numerical data describing the extent and strength of IHC staining throughout the entire volume to be characterized. In this integrated approach, the figures are derived by pooling relative IHC staining intensities from all sections of the series from a particular animal. The procedure (1) eliminates the problem arising from the personal assessment of the significance of the IHC staining intensity, (2) does not depend on the precise dissection of the tissue on a gross scale and (3) considerably reduces the consequences of limited, arbitrary sampling of the region of interest for IHC analysis. The quantification procedure is illustrated by data from an experiment in which inflammatory reactions in the murine spinal cord, measured as microglial activation, were followed by IHC after the lesion of the sciatic nerve.

Introduction

There is a continuous need for accurate comparisons of the distributions of macromolecules in experimental biology or in medical diagnostics, and recurrent attempts are made to achieve the objective light microscopic characterization of the intensity and distribution of reaction products in immunohistochemically (IHC) stained sections. The primary obstacle during the quantification of staining is the frequent non-reproducibility of the pattern, strength and contrast of staining in repetitive investigations. This can originate from variations in the recording conditions (e.g. from uncontrolled, illumination-induced fading in fluorescent microscopy [Kirkeby & Thomsen, 2005], or from differences in the alignment and aperture setting in transmitted light microscopy), which can be at least partially corrected or compensated. However, even with identical microscopic conditions, certain variations still arise from unavoidable differences in the replication of the staining protocols in repeated experiments. Although this manuscript focuses on the problems associated with enzyme-based techniques,
some of the difficulties are common in both enzyme- and fluorescence-based methods. To overcome the difficulties of quantification, individualized classification criteria are often introduced. The methods applied may involve an entirely subjective approach, based on the observer’s estimation of staining intensity, stained cell density and cell morphology (Colburn et al., 1997; He et al., 1997). Alternatively, a semi-quantitative analysis may be applied, with the measurement of grey scale intensity, an operator-independent threshold method being used for determination of the percentage of the total area covered by stained cells (Blackbeard et al., 2007). Automated analysis of corrected immunoreactive areas in selected sections from the region of interest is also possible, likewise based on a subjective, visual selection of the threshold for image segmentation (Stuessel et al., 2000).

The other major cause of problems during quantification is the biological variability. The main source of such experimental error is the interindividual variability, which is responsible for about 70% of the observed experimental variance (Howard & Reed, 1998). Further variance stems from the hierarchical nature of microscopic investigations, that is, from the frequently neglected differences between blocks and sections: in other words, from the way the object is sampled. The contribution of this factor to the overall variance is in the range of 20–25% (Howard & Reed, 1998). Generally speaking, such statistical variability may be decreased by increasing the number of observations (Gundersen & Østerby, 1981). The first problem, however, is associated at least in part, with observer subjectivity. This type of error (bias) does not affect the precision of the measurements, but since the real values sought experimentally are normally not known, the systematic deviations from these values cannot be estimated (see e.g. West, 1999).

The present manuscript provides an integrated procedure that attempts to correct for both types of variability. The method consists of two modules: compensation for the changes in staining intensity at the section level, and the application of systematic sampling at the object level. The irregularity of staining in repeated experiments is taken into account by determining the staining intensity in reference regions in each section that are presumed to be unaffected by the experimental treatment. The density of IHC staining (or the percentage area covered by stained profiles) in the area of interest (AOI) will be related to that in the reference area (AOI′) in the same section. To determine the positivity (significance) of the immune reaction, objective criteria are used: the standard deviation (SD) and the background level of the staining intensity are determined for the AOI′ and a segmentation threshold (background + 2 × SD) is then applied to the whole section. Regions with staining intensity exceeding this limit are accepted as significantly stained profiles. To compensate for the non-homogeneous spatial distribution of the stained structures within the volume to be characterized, a robust systematic random sampling is applied (Gundersen & Jensen, 1987). A specimen containing the entire region of interest is dissected and sectioned equidistantly, with a starting position set at random. The parameters characterizing the IHC staining in each section, related to the AOI′, are then pooled to arrive at simple numerical data expressing the IHC staining of the object of interest.

To illustrate the procedure step by step, an experimental model of nerve injury is used. Nerve transection (axotomy), which is a well-established model of neuronal injury (Koliatsos & Price, 1996), leads to the activation of microglial cells at the site of injury in the central nervous system (Kreutzberg, 1996; Obal et al., 2001). This type of cellular response can be attenuated by treatment with minocycline, a member of the tetracycline class of antibiotics (Zemke & Majid, 2004). In our example, the microglial response in the ventrolateral horn of the murine spinal cord to axotomy of the sciatic nerve is compared with that after minocycline treatment. An enzyme-based IHC method is used to detect the microglia, and the proposed integrated evaluation/sampling procedure is applied to demonstrate the effect of treatment.

Material and methods

Animals

B6/SJL hybrid mice were bred in the Biological Research Center (Szeged, Hungary) from breeding pairs originally obtained from Charles River Laboratories (Budapest, Hungary). Twelve-week-old male mice were used for this study. All animal experiments were performed in accordance with the institutional guidelines for the use and care of animals and with Hungarian governmental laws relating to animal protection (protocol no. XVI./03876/001/2006). The animals were housed (at most four animals per cage) at room temperature under a 12:12 h light/dark cycle, with free access to food and water.

Surgical procedure and pharmacological treatment

Surgery was performed under Avertin (tribromoethanol: Fluka, Buchs, Switzerland) anaesthesia (240 mg kg⁻¹ body weight in a 0.3 mL volume, administered intraperitoneally). The left sciatic nerve was transected in the mid-thigh, and a 3–4-mm portion was removed to prevent regeneration. The incision was then sutured, and the animals were returned to their cages after they had recovered from the narcosis. One group of operated mice were injected intraperitoneally once a day with minocycline (an inhibitor of microglial activation: Sigma, Budapest, Hungary), dissolved in sterile saline solution (40 mg kg⁻¹ body weight in a 0.1 mL volume), first applied 1 h prior to the surgery. The animals in both groups (operated + minocycline-treated, or operated only) were sacrificed on day 7 after the sciatic nerve transection.
Fig. 1. Dissection and sampling of tissue for immunohistochemical (IHC) characterization. (A) A piece of tissue (with a characteristic length of 10–12 mm) from the lumbar portion of the spinal cord was dissected, post-fixed and embedded for sectioning. The dissection was made according to gross anatomical landmarks in such a way that the specimen encapsulated the entire region with the expected inflammatory reaction (red-shaded area). It is not necessary to know the exact location of the region of interest within the dissected tissue. (B) The dissected piece of the spinal cord was systematically sampled by transverse sectioning along the axis of the spinal cord. The starting position for sampling was randomly selected by determining the first location of the specimen where the entire cross-section of the spinal cord could be recognized (onset point; section no. 0); then, before collection of the sections, a random number of sections (between 1 and 10) with 30-μm thickness were cut and discarded. Thus, the sampling started with section no. 1 located at an arbitrary distance \( x \mu m \) (1 < \( x \) < 300 μm) from the border of the tissue. Next, a systematic set of sections was cut at 300 μm distances until the whole spinal cord sample was sectioned through. (C) A set of equidistantly spaced sections was prepared from each spinal cord for IHC staining and quantification, which means that each section in the series represented a 300-μm transverse slice of the spinal cord (15 slices in the figure). The region of interest within the sections was determined by means of microscopic landmarks in the sections. Since the size (length) of the dissected tissue was intentionally larger than that of the region with the expected inflammation (red-shaded area), the very first and the very last sections in the series may not display an increased number of microglial cells in the region of interest.

Tissue preparation and sampling

Under terminal anaesthesia, the animals were perfused transcardially with 30-mL phosphate-buffered saline (PBS; 10 mM) followed by 30 mL 3% paraformaldehyde (PFA; Sigma) in 10 mM PBS. The lumbar segments of the spinal cords throughout the entire region involving the expected inflammatory reaction (approximately 10–12 mm in length) were removed and post-fixed overnight in 3% PFA (in 10 mM PBS) at 4°C (Fig. 1(A)). Tissue blocks were then cryoprotected in 30% sucrose (Sigma) in 10 mM PBS for 3 days at 4°C, and embedded in OCT medium (Tissue-Tek, Zoeterwoude, the Netherlands) for transverse sectioning. Transverse sections at a thickness of 30 μm were cut from the spinal cord along its longitudinal axis on a cryostat (Kryostat 1720; Leitz, Wetzlar, Germany) in such a way that the starting positions were selected at random: the first N sections (N selected at random between 1 and 10) were cut and discarded; then, systematically, each section at a distance of 300 μm was harvested, until the segment of spinal cord had been sectioned.
Uneven illumination. The white balance was adjusted for each first. At the beginning of each microscopic session, a flatfield photographs were taken with the same exposure settings as the was made with automatic camera control, and the following grabbed image was necessary from a section, the first snapshot could fit into a single field of view. When more than one component. The automatic exposure option of the camera
1200 bit) resolution at 8-bit depth for each (RGB) colour recorded with a Spot RT CCD camera (Diagnostic Instruments, Sterling Heights, MI, USA) in colour mode, using full (1600
camera. The same magnification (objective lens) was used to record all images belonging in a particular series. To avoid any slight variability in reproducing aperture settings, both the field limiting and contrast apertures were kept at the fully open position during (digital) photography. Images were recorded with a Spot RT CCD camera (Diagnostic Instruments, Sterling Heights, MI, USA) in colour mode, using full (1600 × 1200 bit) resolution at 8-bit depth for each (RGB) colour component. The automatic exposure option of the camera was used if all the image components of interest in a section could fit into a single field of view. When more than one grabbed image was necessary from a section, the first snapshot was made with automatic camera control, and the following photographs were taken with the same exposure settings as the first. At the beginning of each microscopic session, a flatfield image was taken and used through the session to correct for uneven illumination. The white balance was adjusted for each individual slide. Recorded images were stored in colour mode in uncompressed files with tagged image file format.

**Immuno- and ultrastuctural staining of sections**

To detect microglial cells with an IHC procedure, free-floating sections were stained according to the avidin–biotin technique (Cuello, 1993). Sections were rinsed first with three changes in PBS, which was followed by a 30-min blockade of endogenous peroxidase activity with 0.3% hydrogen peroxide in 10 mm PBS containing 0.2% Triton X-100 (TPBS; Sigma). Then, after washing (three changes in PBS), to reduce the non-specific staining, sections were incubated in 2% normal goat serum (Vector, Burlingame, CA, USA) in TPBS for 1 h. Microglial cells were labelled with anti-CD11b antibody (1:500 in TPBS; rat-anti-mouse antibody, Serotec, Oxford, UK) during incubation at 4°C overnight. After washing in PBS (three changes), sections were incubated in biotin-labelled secondary antibody for 1 h (1:800 in TPBS, goat anti-rat IgG, Vector), followed by washing in PBS (three changes), and incubation in the avidin–biotin complex (1:1600 in PBS, Vector) for 1 h, at room temperature. Finally, after washing in PBS, the reaction was visualized by incubation in diaminobenzidine (DAB, 5% in PBS; Sigma) for 15 min. Stained sections were thoroughly rinsed in PBS, mounted on silane-coated glass slides, dehydrated in graded series of ethanol, processed through xylane (Molar, Budapest, Hungary) and cover-slipped with Entellan (Merck, Darmstadt, Germany).

**Light microscopic image recording**

IHC-stained sections were examined in an Olympus Vanox-T AH-2 light microscope (Olympus, Tokyo, Japan). The alignment of the microscope was checked at the beginning of each microscopic session with special attention to the coaxial position of the lenses, the apertures and the recording CCD camera. The same magnification (objective lens) was used to record all images belonging in a particular series. To avoid any slight variability in reproducing aperture settings, both the field limiting and contrast apertures were kept at the fully open position during (digital) photography. Images were recorded with a Spot RT CCD camera (Diagnostic Instruments, Sterling Heights, MI, USA) in colour mode, using full (1600 × 1200 bit) resolution at 8-bit depth for each (RGB) colour component. The automatic exposure option of the camera was used if all the image components of interest in a section could fit into a single field of view. When more than one grabbed image was necessary from a section, the first snapshot was made with automatic camera control, and the following photographs were taken with the same exposure settings as the first. At the beginning of each microscopic session, a flatfield image was taken and used through the session to correct for uneven illumination. The white balance was adjusted for each individual slide. Recorded images were stored in colour mode in uncompressed files with tagged image file format.

**Image analytical procedures in single sections**

Digital images were analyzed by using the built-in functions of the Image-Pro-Plus image analysis software (Media Cybernetics, Silver Spring, MD, USA) running under the Windows XP operation system on a PC. (Most of the used or equivalent operations are also available in other free or commercial software.)

Although the original colour (digital) pictures were used for easier orientation and to specify the AOI in the sections, all operations were performed on images previously converted to 8-bit greyscale. The evaluation steps included (1) specification of the AOI and AOI , (2) calculation of the background level and variance of the staining intensity on the 0–255 greyscale. (3) segmentation of the significantly stained pixels and (4) calculation of the percentile area of the significantly stained profiles and/or their average density in the AOI, relative to the similar values in the AOI. A simplified version of such calculations to estimate relative changes in IHC staining has already been tested in another model of nerve injury (Obál et al., 2006).

**Specification of the AOI and the corresponding AOI**. The AOI in a single section could be determined by using characteristic anatomical landmarks, such as, in our case, the border of the grey and white matter, the position of the central channel and the perimeter of the pools of motor neurons (Fig. 2(A)). In complex structures, detailed atlases should be consulted, but in most cases the specific regions can be easily marked. In our study, a general histological stain (cremoy violet, which stains Nissl substance and cell bodies) was used in one set of sections to prove that correct orientation was achieved in the IHC-stained sections with no counterstaining, or without the regular need for reference material (Fig. 2(B)). When the region to be analyzed was successfully identified in a section, it was outlined with the pointer device and marked as an (irregular) AOI. For reference purposes, any indifferent region of the section would be suitable; however, exploiting the symmetric organization of the nervous system, we propose that an identical area on the contralateral side of the section should be used. To define this area, the AOI on the operated side was mirrored about the symmetry axis of the section; then, to correct for occasional slight distortions of the sections, it was manually positioned to the corresponding location on the non-operated side (Fig. 2(B)).

**Calculation of the background level and variance of the staining for segmentation**. The major anatomical landmarks were first identified in the original colour images of the sections, and the area where an increased immune reaction was expected on the operated side and the reference area on the contralateral side were then marked accordingly (Figs 3(A) & (B)). The colour
image was next converted to the greyscale with a range of 0–255 (Figs 3(C) & (D)). To determine the background level, a large format low-pass filter (512 × 512 pixels) was first applied to the greyscale image (Figs 3(E) & (F)), which smoothed out the variability in the image owing to differences in IHC staining by substituting the grey value of each pixel with the average grey value of its neighbourhood with a dimension of 512 × 512 pixels. Although the image obtained approximated the grey value of its neighbourhood with a dimension of 512 × 512 pixels, the variability in the image owing to differences in IHC staining was used for thresholding: all the image points outside the stained profiles (cf. Figs 3(D) & (F)), this background distribution is certainly contaminated with the signal from the heavily stained area, which is obvious from a comparison of Figs 3(C) and (E). Next, to correct for this effect, an adjusted (second) background level was calculated. To derive this number, the first background level within the AOI was used for thresholding: all the image points darker than this level (which, at this approximation, were regarded as the stained profiles) were excluded from the image, and an average grey value was calculated for the remainder of the pixels within the AOI (Figs 3(G) & (H)). This number was accepted as the final value representing the grey level of the background (BCKGND).

The variance of the staining, that is, the variance of the grey level distribution of the images, in addition to the BCKGND value, was used as an aid to determine which image point should be regarded as significantly stained. To calculate the standard deviation (SD) of the distribution of staining intensity, all the image points within the AOI' of the original greyscale images were taken into account. Finally, those pixels in the AOI and AOI', whose grey values were lower than BCKGND – 2 × SD (i.e. significantly darker than the background), were dissected and accepted as significantly stained image points. For illustration and for visual control of the results, image areas with dissected pixels were outlined, and these profiles and the perimeters of the AOI and AOI' were superimposed on the original colour images (Figs 3(I) & (J)).

Calculation of tissue-specific parameters from the relative changes in IHC staining in individual sections. Two types of (non-independent) parameters were derived: (i) the percentile areas of the significantly stained profiles within the AOI and AOI' and (ii) the intensities of average staining within these areas. The first parameter was readily obtained after segmentation of the greyscale images at the cut-off level, since this operation yields the total area of the AOI (equal to that of AOI') and the sum of the areas of the segmented profiles in both regions. The numerical difference of these percentile areas results in a relative number describing how much larger the coverage of the cellular staining on the operated site is than that of an unperturbed, control area (Δ(area)).

The attenuation law (Bouguer–Lambert–Beer law) states that

\[
I_1 = I_0 \times e^{-M \times T \times C},
\]

where \(I_0\) is the incident and \(I_1\) is the transmitted light intensity, \(M\) denotes the molar absorptivity (extinction coefficient), \(T\) is the section thickness and \(C\) is the local concentration of the absorbing species. Assuming homogeneous section thickness and similar extinction properties of the absorbing material in the AOI and AOI', analogous equations can be written for the operated and control sides as:

\[
I_{1(\text{operated})} = I_0 \times e^{-M \times T \times C(\text{operated})}
\]
\[
I_{1(\text{control})} = I_0 \times e^{-M \times T \times C(\text{control})}.
\]

Hence,

\[
-\ln(I_{1(\text{operated})}/I_{1(\text{control})})/M/T = C(\text{operated}) - C(\text{control}),
\]
Fig. 3. Step-by-step procedure for the segmentation of significantly stained profiles in IHC-stained sections. (A, B) According to the anatomical landmarks (shape of the section, border of the grey and the white matter [dotted line], etc.), the region where the increased staining is accepted is outlined on the operated side; then, symmetrically, the reference area is specified on colour images of the section (RGB, 8-bit depth). (C, D) The colour pictures are converted to greyscale images with a range of 0–255 (from black to white) and the AOI and AOI’ are next copied as overlay images. (E, F) Application of a large-scale low-pass filter (512 × 512 pixels) to the whole images resulted in approximated background images by smoothing (averaging) the details in the greyscale images. In this particular case, the filtering process resulted in an average greyscale (background) level of 233 in the reference area (AOI’). Comparison of the greyscale distribution within and outside the AOI on the operated side indicates how the recruitment of the stained cells...
Fig. 3 continued (Fig. 3E) affects the local grey level of the background. (G, H) When the segmentation threshold was set to the calculated background (a grey value of 233 in the presented case), all the pixels with grey values lower than this limit (black ‘holes’ in the image) were excluded from the image, and a new average background level (238) was then calculated for the remainder of the pixels within the reference area (AOI′). With this new value for the background (BCKGND), a standard deviation (SD) of the greyscale distribution was calculated for the pixels darker than this level (i.e. grey values lower than this number) within the AOI′ in the greyscale images (Fig. 3D), and a cut-off value for the segmentation of the significantly stained profiles was calculated at BCKGND – 2 × SD. (I, J) With the cut-off value, the segmented profiles within the AOI and the AOI′ were outlined and superimposed on the original colour images for visual control of the results representing the significantly stained profiles.

which means that the logarithm of the ratio of the light intensities (measured on a greyscale in this case) in the AOI and the AOI′ is proportional to the difference in concentration of the absorbing species in these areas, that is, the target of determination (Δ(concentration)).

Obtaining parameters which characterize the entire volume to be analyzed

In each equidistant section cut from the tissue, consistent individual parameters can be derived which are either proportional to the concentration of the IHC reaction products (Δ(concentration)), or to their dimensions (Δ(area)) on the operated side relative to the corresponding data on the control side. Summation of the corresponding data obtained from each section, furnished simple numerical data characterizing the dimension (ΣΔ(area)) or the extent of IHC staining (ΣΔ(concentration)) throughout the whole volume of the region of interest from a given animal (Figs 4(A) & (B)). The process of summation of data situated equidistantly along the third Cartesian axis is equivalent to a certain type of integration, which incorporates the third dimension; thus, the calculated numerical data should be regarded as having dimensions of volume and mass on an arbitrary scale.

Results

Six animals serving as controls underwent sciatic nerve transection, while four other mice received minocycline treatment in addition to the axotomy. The reducing effect of the minocycline treatment on the microglial activation in the spinal cord following axotomy was tested by comparing the levels of CD11b IHC staining in the ‘operated only’ and ‘operated + minocycline’ groups of animals with the methods presented earlier. The data in Table 1 reveal that for both the reduction in the volume occupied by the significantly stained cells (P = 0.022), and the reduction in the total density of staining (P = 0.026) in the ventrolateral pool of motor neurons, the effect of minocycline was significant (Student one-tailed t-test).

Discussion

Although snap freezing can reduce the tissue preparation time considerably, which may be critical if tissue bio-banking is associated with a rapid intra-operative diagnosis, perfusion or immersion fixation gives superior tissue preservation (Beltz & Burd, 1989); furthermore, even for diagnostic purposes, most of the tissue is still treated with formalin fixative (Steu et al., 2008). Admitting also that snap-frozen tissue can additionally

![Fig. 4](image-url). Charts showing relative area (A) and density (B) data obtained from the analysis of individual sections belonging in the same systematic random series. The category axis gives the serial numbers of consecutive sections. Obviously, the exact location of the starting point of the systematic sampling is indifferent, since the difference between the operated and non-operated sides fluctuates around zero outside the volume affected by the experiment, and is therefore cancelled out on a statistical basis during summation of the data. (A) The percentile area covered by the IHC products in the reference area (AOI) was subtracted from that in the region of interest (AOI′) in each section and plotted. The sum of the data (taking into account their signs) is 159.54, which characterizes the extent of the IHC-stained structures within the volume to be analyzed. (B) The calculated numbers from each section [–ln(I(op)/I(co))], which are directly proportional to the differences in concentration of the IHC-stained structures, are plotted against the section number. The result of summation (0.67) characterizes the accumulation of IHC-stained material in the volume of interest.
offer exceptional possibility of the combination of stereological, histological and molecular analyses of adjacent sections from the same tissue (Schmitz et al., 2000), the chemical fixation preparation methods for IHC are still more widespread than the physical freezing procedures. For this reason, we followed the traditional perfusion fixation protocol for tissue preparation in the present study.

The commonly experienced difficulties inherent in the interpretation of IHC-stained sections are mainly due to unavoidable variations arising from two main sources: the variability of the staining at the section level and the diversity of the structure at the tissue level. In consequence, subjective input from the operator is necessary at critical points in the evaluation, which influences the results to an incalculable degree. The approach presented here for the evaluation of IHC-stained sections are mainly due to the present study.

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### Procedures at the section level

**Operator input.** In the presented method, the input of the operator is reduced to one unavoidable step: specification of the boundary of the studied AOI. The use of an identical AOI in the same section for control purposes ensures that changes relative to an indifferent area are calculated in each case even if, by chance, the specification of the AOI is not absolutely precise. If the AOI is to some extent smaller than the area delineated by the boundary of the anatomical region to be analyzed, the area will be under-represented, but the calculated relative difference between the AOI and AOI will not be considerably affected. If the AOI is somewhat larger than the area to be analyzed, the measured difference from the control region will be only slightly ‘diluted’ with data from an unaffected area, that is, the method will merely lose some sensitivity. The major landmarks in the sections, however, allow easy and reliable delineation of the area AOI (see Fig. 2). Accordingly, all such inconsistencies would cause relatively minor changes in the results.

**Determination of the background staining.** In the present experiments, no counterstaining was performed, and there was therefore no need for spectral (colour) aided segmentation of the IHC-stained profiles from the counterstained background, which would have required complex operations (van der Laak et al., 2000; Brey et al., 2003; Pham et al., 2007). Thus, instead of the selection of a specific wavelength (or colour) for analysis, or decoupling of the intensity data from the red–green–blue colour model, all images were converted to greyscale images represented only by 8-bit data at each image point, and the image analysis procedures were performed in a single channel with simplified calculations.

In each section, the intensity of the background staining (determined in an indifferent reference area) is used to identify image points with a level of staining significantly above that of the surroundings both in the AOI and within the AOI. Thus, the procedure ensures that the variability of staining in repeated experiments or the heterogeneity of the staining from section to section in a given series will be compensated. The procedure, which automatically calculates the background intensity on the basis of the distribution of the staining intensity without the aid of the operator, consists of two steps. In the first step, an average background image is calculated by applying a large-format low-pass filter to the digital image of the section, which yields a greyscale value at each image point with the average for its 512 × 512 pixel neighbourhood (Figs 3(E) & (F)). Since the great majority of the image points show no evidence of staining (Figs 3(C) & (D)), the result of this filter approximates closely, though not perfectly to the true background distribution (Figs 3(E) & (F)). After low-pass filtering, an average grey value (approximate background level) can be calculated within the AOI (233; Fig. 3(F)). The deviation of this number from the true (unknown) background is obviously greater in those regions where stained profiles is notably accumulate (cf. Figs 3(C), (E) and (D), (F)). To correct for this ‘contamination’ of the background distribution from the stained cells, as a second approximation, all image points with grey values lower than the calculated first background value (233) are excluded from the calculation, and a new average background value (BCKGND = 238) is determined for the remainder of the pixels within the AOI. This procedure of successive approximation of the true average background

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**Table 1. Effect of minocycline treatment on CD11b IHC staining after sciatic nerve cut.**

<table>
<thead>
<tr>
<th>Animals</th>
<th>Operated only</th>
<th>Operated + minocycline</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Sum of relative stained area (%)</td>
<td>Sum of ( \ln(I_{\text{op}}/I_{\text{co}}) )</td>
</tr>
<tr>
<td>#1</td>
<td>374.28</td>
<td>1.67</td>
</tr>
<tr>
<td>#2</td>
<td>394.03</td>
<td>1.83</td>
</tr>
<tr>
<td>#3</td>
<td>549.73</td>
<td>2.38</td>
</tr>
<tr>
<td>#4</td>
<td>642.55</td>
<td>3.19</td>
</tr>
<tr>
<td>#5</td>
<td>541.82</td>
<td>2.47</td>
</tr>
<tr>
<td>#6</td>
<td>657.43</td>
<td>3.24</td>
</tr>
<tr>
<td>Mean</td>
<td>526.64</td>
<td>2.46</td>
</tr>
<tr>
<td>SEM</td>
<td>49.01</td>
<td>0.66</td>
</tr>
</tbody>
</table>
within the AOI should in principle be stopped when the difference between the results of successive iterations is smaller than a specified limit. In the presented experimental example, even in the third step, the change in the calculated average background greyscale value was less than one digit on the 0–255 scale, and accordingly no further iterations were performed.

**Segmentation of stained profiles and calculation of parameters characterizing the stained structures.** After determination of the BCKGND and the SD values of the greyscale distribution in the AOI, a cut-off level is determined to segment those image points that are considered significantly stained, that is, pixels with grey values significantly different from that of the background. This seemingly arbitrary decision is supported by statistical principles if the deviation of the cut-off level from the BCKGND value is set in units of SD, which determines the confidence of the results. In biological applications, this offset is normally set to a value in the range $2 \times SD$ to $3 \times SD$ implicating a probability between 67 and 99% for the segmented image to reflect the distribution of the authentic stained profiles. In our application example, the level BCKGND $- 2 \times SD$ was consistently applied.

After segmentation, different parameters were derived to characterize the extent and strength of the IHC staining in the AOI relative to the AOI. The difference between the numbers of pixels segmented in the AOI and the AOI (since these areas are identical) reflects the net increase in the stained cellular profiles due to the treatment. For easier comparison of the results obtained from different sections and different animals, these raw data were normalized to the total number of pixels within the AOI or the AOI, and expressed as the percentile area difference ($\Delta \text{(area)}$). The difference in staining intensity between the AOI and the AOI was determined directly by using the attenuation (Bouguer–Lambert–Beer) law, applied to the whole area of the AOI and the AOI. Through measurement of the average (optical) density in these areas, a number corresponding to the difference in the percentile stained area was obtained ($\Delta \text{(concentration)}$). The method is analogous to the classical procedures applied successfully in conventional (electron) microscopy to determine the local mass of a specimen by measuring the contrast of the micrographs (Halliday & Quinn, 1960; Zeitler & Bahr, 1965; Edie & Karlsson, 1977).

**Procedures at the tissue level**

Determination of the boundary of the AOI and the setting-up of the exact anatomical location along the direction perpendicular to the plane of the sections (unlike the same procedure within the plane of the section) is often hindered by the missing characteristic landmarks that can be recognized only in a set of consecutive sections (see Fig. 1). Thus, to overcome the need for serial sectioning in order to determine the exact anatomical location within the tissue, if probing of a tissue by means of microscopic sections is intended, arbitrary (random) sampling is often the choice. The paradigm, however, can lead to misleading results, due to the intra-individual variability of the tissue, an effect which is frequently neglected. However, as illustrated qualitatively in a composite figure assembled from the sections of our application example (Fig. 5), huge differences may be observed in the amount of staining, depending on the actual position of the section within the region to be analyzed. Quantitatively, as depicted in Fig. 4, fluctuations from section to section as large as 50–100% of the derived parameter values can be determined in the region of the tissue containing the expected reaction.

The proposed procedure of systematic sampling is not sensitive to the actual positions of the sections within the tissue, since the intra-individual variations average out during summation of the results obtained on the individual sections. The method is not sensitive to the actual starting point of
the sampling either, as long as the dissected tissue contains the entire region of interest, since the differences neutralize each other during the summation process outside the affected volume, due to the statistical nature of the variations between the operated and the control side (see Fig. 4).

Application of the procedure

In our application example, as a positive control, the effect of an antibiotic (minocycline) on the local microglial reaction on the spinal cord was tested experimentally after lesion of the sciatic nerve. This known anti-inflammatory compound has been shown to improve regeneration and reduce microglial activation in neuronal lesions (Blackbeard et al., 2007; Shankaran et al., 2007; Henry et al., 2008; Mishra & Basu 2008). In accordance with the literature data, our method to quantify the relative staining intensity of IHC-stained microglial cells after axotomy, demonstrated a significant decrease in microglial activation after minocycline treatment (Table 1). The presented procedure is of general applicability to quantify the relative staining intensity of IHC-stained microglial cells in the spinal cord of mice by AIS IgG. Neuroreport 12, 2449–2452.

References


Hypoglossal Motor Neurons Display a Reduced Calcium Increase After Axotomy in Mice with Upregulated Parvalbumin

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ABSTRACT
Motor neurons that exhibit differences in vulnerability to degeneration have been identified in motor neuron disease and in its animal models. The oculomotor and hypoglossal neurons are regarded as the prototypes of the resistant and susceptible cell types, respectively. Because an increase in the level of intracellular calcium has been proposed as a feature amplifying degenerative processes, we earlier studied the calcium increase in these motor neurons after axotomy in Balb/c mice and demonstrated a correlation between the susceptibility to degeneration and the intracellular calcium increase, with an inverse relation with the calcium buffering capacity, characterized by the parvalbumin or calbindin-D28k content. Because the differential susceptibility of the cells might also be attributed to their different cellular environments, in the present experiments, with the aim of verifying directly that a higher calcium buffering capacity is indeed responsible for the enhanced resistance, motor neurons were studied in their original milieu in mice with a genetically increased parvalbumin level. The changes in intracellular calcium level of the hypoglossal and oculomotor neurons after axotomy were studied electron microscopically at a 21-day interval after axotomy, during which time no significant calcium increase was detected in the hypoglossal motor neurons, the response being similar to that of the oculomotor neurons. The hypoglossal motor neurons of the parental mice, used as positive controls, exhibited a transient, significant elevation of calcium. These data provide more direct evidence of the protective role of parvalbumin against the degeneration mediated by a calcium increase in the acute injury of motor neurons.

INDEXING TERMS: calcium; calcium buffering; parvalbumin; axotomy; neuroprotection

Motor neuron disease (or amyotrophic lateral sclerosis [ALS]) is a fatal degenerative disorder of the nervous system that affects primarily the upper and lower motor neurons (Rowland and Shneider, 2001). Its etiology is basically unknown, although several mechanisms have been proposed, which probably contribute synergistically to the destruction of the motor neurons (Boillée et al., 2006; Bendotti and Carri, 2009; Rothstein, 2009). The sequence of events during these pathologic processes is mostly known, and some aspects of their reciprocal contribution to the propagation of the injury cascade have been well described (Mattson, 1998).

Our research focuses on the role of the increase in the intracellular calcium level of the motor neurons during degeneration. We hypothesize that the inability of the motor neurons to cope successfully with a nonphysiologic calcium load after or during injury may be one of the key events maintaining the complex destructive machinery through a positive feedback of the elementary steps, and an elevated calcium buffering capacity may attenuate the degree of the lesion during stress conditions (Appel et al., 2001; Siklós and Appel, 2005). Support for this hypothesis has emerged from clinical observations and experimental studies. Patients with ALS present an increased

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level of calcium in the affected motor nerve terminals (Siklós et al., 1996), with a correlation between the preservation of specific motor (external oculomotor and voluntary sphincter) functions and the intracellular levels of certain calcium binding proteins ([CaBPs] calbindin-D_{28k} [CB] and parvalbumin [PV]) in the corresponding motor neurons (Ince et al., 1993; Alexianu et al., 1994).

In accordance with these observations, in transgenic animals containing the G93A mutation of the Cu/Zn superoxide dismutase (SOD1) gene, which serve as a model for the familial form of ALS (Gurney et al., 1994), the viability of the motor neurons paralleled their PV content; in contrast, in accord with our hypothesis, decreases are observed in the intracellular calcium, with structural abnormalities in these cells (Siklós et al., 1998). On the basis of in vitro physiology measurements in oculomotor and hypoglossal neurons, the prototypes of motor neurons with high and low CaBP content, respectively, a correlation was proposed between the CaBP content and the ability of the motor neurons to resist calcium-mediated injury (Lips and Keller, 1998; Vanselow and Keller, 2000). Our earlier electron microscopic study following axotomy/target deprivation in these same motor nuclei in vivo provided further support for this concept: a lower increase in calcium level and less significant degeneration were demonstrated in the oculomotor neurons in which the PV content was higher than that in the hypoglossal motor neurons from mice of the parental line (Ince et al., 1993; Alexianu et al., 1994).

The results of the above experiments provided indirect evidence of the calcium buffer hypothesis, because different pools of motor neurons were compared in each case. In fact, the embedding of these motor neurons into different neuronal networks and different microenvironments may contribute to their different susceptibilities to degeneration. In order to test the hypothesis of the protective role of CaBPs more directly, the same population of motor neurons with high and low CaBP content, respectively, a correlation was proposed between the CaBP content and the ability of the motor neurons to resist calcium-mediated injury (Lips and Keller, 1998; Vanselow and Keller, 2000). Our earlier electron microscopic study following axotomy/target deprivation in these same motor nuclei in vivo provided further support for this concept: a lower increase in calcium level and less significant degeneration were demonstrated in the oculomotor neurons in which the PV content was higher than that in the hypoglossal motor neurons, in which PV and CB could not be detected (Obál et al., 2006).

The description of the generation of the PV overexpressing mice used in this study has been given elsewhere (Beers et al., 2001). Parental breeding between transgenic CaMKII-parvalbumin mice #4031 × 4034, 4035 obtained from Dr. S.H. Appel (The Methodist Hospital Neurological Institute, Houston, TX) was set up at the conventional animal facility of the Institute of Experimental Medicine, Hungarian Academy of Sciences, Budapest. All transgenic mice used in this study were derived from the original parental cross between transgenic male #4031 mated to 4034 and #4045 females of line Tg (parv14-4). All litters obtained from this cross (mice #12–21) and subsequent sibling/sibling matings as well as the parents were homozygous for the transgene as proved by test crosses to wild-type mice.

To prove the presence of the transgene, DNA was isolated from tail biopsies and prepared according to a standard protocol published online on the Jackson Laboratory website (http://www.jax.org/imr/tail_phenol.html). In general, 0.1 µg of genomic tail DNA was used to amplify the transgene by using two different primer pairs:

1. Primers derived from the third and fourth exons of the endogenous CaMKII gene as follows: forward: 5’-CTGCAGACTCCTTGGACCC-3’ and reverse 5’-GCCATACGCTTTGTTC-3’. The sizes of the products were 200 bp (transgene) and 1,400 bp (endogenous CamKII gene).
2. Primers were: forward primer 5’-GCGAGGAGTCGAGTGGTTGTCTG-3’ and reverse primer 5’-TTCGGCCACCAGAGTGGAGAATTC-3’ (Beers et al., 2001), size of the product 1,200 bp.

Amplification was as follows: denaturation at 94°C for 4 minutes, followed by 30 cycles of 30 seconds at 94°C, 45 seconds at 57°C and 45 seconds at 72°C, final extension 7 minutes at 72°C. DNA was run on EtBr-containing agarose gels and photographed under UV light.

**Axotomy experiments**

Twelve-week-old male B6/SJL mice and PV+/+ mice were selected for the study. To avoid multiple survival surgery, animals were assigned either to hypoglossal or to oculomotor nerve cut experiments. A total of 60 mice were used either for hypoglossal axotomy (n = 40) or for oculomotor surgery (n = 20). Four animals were euthanized at five time points for each type of surgery and were processed for electron microscopic calcium histochemistry study. Surgical procedures were performed under deep anesthesia with Avertin (tribromoethanol,
Fluka, Buchs, Switzerland; 240 mg/kg body weight in a 0.3-ml volume) administered intraperitoneally. The hypoglossal nerve was transected lateral to the hyoid bone and a 3–5-mm nerve segment was removed to prevent regeneration. For transection of the left oculomotor nerve, the animals were enucleated and the orbits were cleared of the remaining extraocular muscles and nerve segments. In both cases, the nonoperated side served as an internal control.

Following surgery, the animals were housed in a temperature-controlled room with a 12-hour light/dark cycle, with access to water and food ad libitum, and were allowed to survive for 1, 4, 7, 14, or 21 postoperative days. All the experiments were performed in accordance with the institutional guidelines for the use and care of experimental animals and the appropriate governmental laws for animal protection (protocols 72-45/b/2001 and 03876/0014/2006).

**Immunocytochemical staining procedures**

A detailed characterization of the PV+/+ mice has been presented in our earlier research paper (Beers et al., 2001). In addition, in the present study, PV was demonstrated in nerve nuclei III and XII in selected B6/SJL (n = 2) and PV+/+ (n = 2) mice by means of light microscopic immunocytochemistry using a polyclonal primary antibody against PV and the avidin-biotin detection technique. This antibody, obtained from Swant (Bellinzona, Switzerland; code no. PV-28, lot no. 5.5) as lyophilized antisera, was produced in rabbits by using rat muscle PV as the antigen. According to the manufacturer’s data sheet, the PV-28 antisera labels a subpopulation of neurons in the normal mouse brain with high efficiency, but does not stain the brain of PV knockout mice. In our titration experiment, a 1:60,000 dilution of the antibody was selected for subsequent regular use.

As a positive control, sections obtained from the brainstem and cerebellum of normal mice were stained and a pattern of immunoreactivity was obtained that was similar to that described as PV immunoreactivity in the literature (Celio, 1990) (data not shown). For immunocytochemical staining, mice under irreversible anesthesia with methoxyflurane were perfused transcardially with phosphate-buffered saline (PBS; 10 mM, pH 7.4), followed by 3% paraformaldehyde in 10 mM PBS (pH 7.4). The brains were removed and immersed in the same fixative for 24 hours (4°C), and tissue blocks containing either the entire hypoglossal or the oculomotor nuclei were then prepared. To prevent the formation of freezing artifacts, tissue samples were placed in sucrose (30% in 10 mM PBS, pH 7.4) for at least 2 days at 4°C. Sections with a nominal thickness of 30 μm were then cut on a cryostat at –18°C (Krioostat 1720; Leitz, Wetzlar, Germany), and collected in PBS.

Immunostaining was performed on free-floating sections. Endogenous peroxidase activity was first blocked with H2O2 (Sigma, St. Louis, MO; 0.3% in 10 mM PBS) for 20 minutes, followed by incubation in Triton X-100 (Sigma; 0.2% in 10 mM PBS) for 30 minutes to enhance antibody penetration. Sections were pretreated with normal goat serum (Vector [Burlingame, CA] S-1000; 2% in 10 mM PBS) for 1 hour, followed by an overnight incubation at 4°C with the primary antibody. After washing, sections were incubated in biotinylated goat-anti-rabbit secondary antibody (Vector BA-9200; 1:200) for 1.5 hours at room temperature. The bound antibodies were detected with the Vectastain Elite ABC Kit (Vector PK-6100) and visualized with diaminobenzidine (DAB; Sigma). After washing, sections were finally mounted on silane-coated glass slides, dehydrated, and coverslipped with Entellan (Merck, Darmstadt, Germany). Parvalbumin-positive motor neurons in nerve nuclei III and XII were identified according to their shape, size, and distribution.

**Light microscopic evaluation of parvalbumin immunostaining**

Stained sections were examined in an Olympus Vanox-T AH-2 light microscope. Images were recorded with a Spot RT CCD camera (Diagnostic Instruments, Sterling Heights, MI) at 1,600 × 1,200 pixel resolution and stored as 8-bit uncompressed grayscale images in tagged image file format (TIFF). Digital Images were analyzed by using the Image-Pro-Plus image analysis software (IPP; Media Cybernetics, Silver Spring, MD). The staining intensity was assessed by a macro module developed for the IPP program in our laboratory to quantify light microscopic immunohistochemical staining (Paizs et al., 2009).

Briefly, after the specification of the region of interest (I) and a control region (C; for internal reference) in each section, the program calculates the logarithm of the ratio of the transmitted light intensities at these regions (−ln(intensity(I)/intensity(C))], which is, according to the attenuation law (Bouguer-Lambert-Beer law), proportional to the difference between the concentrations of the light-absorbing material (immunohistochemical staining) at areas I and C. These values, representing the background-corrected staining intensities, were pooled, to form four groups according to the strains (B6/SJL and PV+/+) and motor nerve nuclei (III and XII). Eight fields were analyzed in each group, and the corresponding average values were compared by one-way ANOVA with Duncan post hoc comparison (StatSoft, Statistica, Tulsa, OK).

Parvalbumin-positive motor neurons in nerve nuclei III and XII were identified according to their shape, size, and distribution. The size distribution of the PV-positive cellular profiles was determined by using the appropriate
built-in function of the IPP program, and then frequency histograms were constructed in each group that were compared with those of motor neurons in the corresponding nuclei available in the literature (Eberhorn et al., 2006; Schwarz et al., 2009).

**Specimen preparation for electron microscopic detection of calcium**

Mice anesthetized irreversibly with methoxyflurane were perfused transcardially, first with 90 mM potassium oxalate (Sigma; pH adjusted to 7.4 with KOH) and then with 3% glutaraldehyde (Polysciences, Eppelheim, Germany) containing 90 mM potassium oxalate (pH 7.4) (Borgers et al., 1977, 1981). Tissue blocks containing the nuclei of either cranial nerve III or XII were dissected and fixed overnight in the same fixative (4°C). Specimens were rinsed in 7.5% sucrose in 90 mM potassium oxalate (pH 7.4), and then postfixed for 2 hours (4°C) in 2% potassium pyroantimonate (Mercck) and 1% osmic acid (Sigma), adjusted to pH 7.4 with acetic acid (Molar). Following postfixation, specimens were rinsed in distilled water (pH adjusted to 10 with KOH) for 10 minutes, dehydrated in graded series of ethanol, processed through propylene oxide (Sigma), and embedded in plastic (Durcupan ACM, Fluka). Blocks were polymerized for 2 days at 56°C.

Semithin (0.3-μm) sections were cut from the blocks on an Ultracut UCT ultramicrotome (Leica), etched (Maxwell, 1978), stained (Richardson et al., 1960), and screened under the light microscope (Olympus Vanox-T) to identify the borders of nuclei III and XII. After trimming of the blocks, 50-nm ultrathin sections were obtained, mounted on formvar-coated single-hole copper grids, contrasted with uranyl acetate and lead citrate (Reynolds, 1963), and analyzed under the electron microscope (Zeiss CEM 902). The oxalate-pyroantimonate procedure applied for the demonstration of calcium at the ultrastructural level resulted in electron-dense deposits (EDDs) due to the precipitation of tissue calcium by the fixative, which was easily recognized by conventional transmission electron microscopy.

**Specificity control of the histochemical reaction for calcium**

The specificity of the fixation procedure for calcium was checked by electron spectroscopic imaging (Bauer, 1988; Reimer et al., 1988). For such analysis, 15–25-nm sections were prepared, mounted on uncoated 700-mesh copper grids, and, without contrasting, examined under the electron microscope in spectroscopic imaging mode at ΔE = 250 eV. During the analysis, the accelerating voltage was set to 80 kV; 90-μm objective and a 650-μm spectrometer entrance apertures were used, with the slit width adjusted to 8–10 eV. By using the attached image analysis system (MTI-66 DAGE high-sensitivity camera; IBAS 2.0 image analysis program, Kontron, Munich, Germany), two-dimensional spectroscopic analysis was carried out at 20,000× instrumental magnification by recording element-specific and background images at energy loss values of the absorption edge of calcium (355 eV) and below (310 eV). To produce low-noise images for calculations, 500–600 video images were averaged for each energy loss value. The analytical image displaying the net calcium distribution was obtained by subtracting the background image from the “edge” image. The non-significant signal below the (mean ± 2.5 × s.d.) level was deleted in the analytical image, and then, for easier comparison of the co-distribution of the EDDs with the true calcium, the significant calcium signal was false-colored according to the significance level in the [mean ± 2.5 × s.d.] to [mean + 6 × s.d.] interval.

This final image, representing the distribution of the precipitated calcium in the tissue, was compared visually with micrographs obtained by structural imaging of the same area, displaying the arrangement of the EDDs. To match the resolution of the structural image to the digital analytical images (512 × 512 pixel), fine structural images were obtained in the spectroscopic mode of the analyzer as well, tuned to the carbon absorption edge (250 eV), which resulted in good-quality structural images of the sections (Körtje and Körtje, 1992). Such images, resembling “darkfield” images, were then inverted for easier comparison.

**Semiquantitative evaluation of the intracellular amount of EDDs**

Ultrathin sections prepared from the motor nuclei were screened under the electron microscope at low magnification (3,000×) to identify motor neurons, which were selected randomly for measurements. One region from each cell body, also selected at random, was photographed at a magnification of 12,000×, until 12–15 samples had been collected from both the operated and the nonoperated side of each nucleus. The negatives were printed at 2.7× enlargement. The amounts of calcium in the tissue compartments were determined by measuring the volume of the EDDs, and were expressed as volume-to-volume ratios (volume fractions) with respect to a selected reference volume. Three types of reference spaces were selected: either the whole perikaryal volume, or the mitochondrial, or the extramitochondrial compartments of the perikaryal space. For brevity, the latter was called the cytoplasmic volume. The three-dimensional parameters, the relative perikaryal, mitochondrial, and cytoplasmic volumes occupied by the
EDDs were determined on two-dimensional electron microscopic prints with a point counting technique (Mayhew, 1992).

After random superimposition of a tessellation of sampling points on the prints, grid points overlying either the EDDs \( P_{\text{EDD(cyt)}} \) or \( P_{\text{EDD(mit)}} \) or the corresponding reference area \( \Sigma P_{\text{cyt}} \) or \( \Sigma P_{\text{mit}} \) were counted. The scores obtained on individual micrographs for both compartments were pooled according to the operated and nonoperated sides of the examined nucleus of each animal, and the corresponding ratios expressing the “histochemical concentration of calcium” in the mitochondrial and cytoplasmic compartments were calculated for both sides \( \Sigma P_{\text{EDD(cyt)}}/\Sigma P_{\text{cyt}} \) and \( \Sigma P_{\text{EDD(mit)}}/\Sigma P_{\text{mit}} \). Finally, the corresponding values obtained on the operated side were divided by those obtained on the nonoperated side to express the changes (fold increase) induced by the axotomy in the mitochondrial and cytoplasmic calcium contents for each animal.

The fold increases in calcium in the hypoglossal and oculomotor nuclei of the PV+/- animals and in the hypoglossal nucleus of the B6/SJL mice were statistically compared on postoperative days 1, 4, 7, 14, and 21 (two-way ANOVA with Duncan post hoc comparison; StatSoft Statistica). The differences were accepted as significant at \( P < 0.05 \).

To produce mounted sets of figures for the illustration of the results, electron micrographs and light microscopic images were selected and processed digitally with the Photoshop program (Adobe Systems, San Jose, CA, version 8.0). Electron microscopic negatives were scanned with a transillumination scanner (HP ScanJet 7400c) at 300 dpi resolution as grayscale images, inverted, and imported to the Photoshop program. Light microscopic images were directly transferred to the Photoshop program at the original resolution of the CCD camera (1,600 × 1,200 pixel). Images were cropped, resized, and mounted to produce tables of figures at the final size. Shading corrections (contrast/brightness and gamma corrections) were applied to compensate for the occasional slight differences in the appearance of the individual pictures to harmonize the overall appearance of all the figures.

**RESULTS**

Although the PV+/- mouse line #14 used in the present study was bred to homozygosity, from time to time the integrity of the colony was confirmed with polymerase chain reaction (PCR), by using tail samples from the animals (Fig. 1).

For characterization of the distribution of calcium precipitated during fixation, the specificity of the method was checked periodically in the present study by two-dimensional analytical electron microscopy. These measurements proved that the pattern of EDDs visible in conventional transmission mode of the microscope can be identified with the distribution of calcium in the tissue (Fig. 2). Qualitatively, a gradual elevation of the intracellular calcium content in the axotomized hypoglossal motor neurons of the B6/SJL mice was seen with increasing postoperative time relative to the nonoperated side, with a tendency to return to the control level by postoperative day 21 (Fig. 3). In the hypoglossal neurons of the PV+/- animals, however, except for a slight increase in the mitochondrial calcium content on postoperative day 1, the calcium content was observed not to be elevated relative to the control side on any postoperative day (Fig. 4). In the oculomotor neurons of the PV+/- mice, similarly as in the hypoglossal neurons in these animals, no calcium increase was detected in the injured neurons as compared with the intact cells at any time point after the surgery (Fig. 5).

In both motor nuclei, the volume fraction of the EDDs, characterizing the distribution of calcium in the tissue, was expressed quantitatively. The results are presented graphically in Figure 6. The visual evaluation of the charts depicting the temporal changes in the total (Fig. 6A), mitochondrial (Fig. 6B), and cytoplasmic calcium (Fig. 6C) was consistent with the qualitative findings on the electron microscopic prints, suggesting that the calcium level in the hypoglossal neurons of the B6/SJL animals gradually increased after axotomy, whereas in the PV+/- animals there was no significant increase in calcium in either the oculomotor or the hypoglossal motor neurons at any time point. According to the raw data, the calcium increase in the mitochondria of the motor neurons of the
B6/SJL mice commenced earlier (day 1, Fig. 6B) and lasted longer (day 14, Fig. 6B) than in the cytoplasmic compartment (day 4 and day 7, Fig. 6B), a temporal difference that was not, however, significant ($P = 0.55$).

Statistically, the time dependences of the calcium levels in the two nuclei in the PV$^+/+$ strain did not differ, regardless of whether the total calcium ($P = 0.21$) or the cytoplasmic ($P = 0.43$) and mitochondrial ($P = 0.20$) calcium levels separately were compared (two-way ANOVA). However, the postoperative calcium increase in the motor neurons from the B6/SJL mice was significantly different from the corresponding situation in the PV$^+/+$ animals ($P = 0.001$, two-way ANOVA). The subsequent multiple comparison procedure revealed that this difference could be attributed to an approximately 1.5-fold increase in the total calcium in the hypoglossal neurons of the B6/SJL animals on day 7 ($P = 0.009$) and day 14 ($P = 0.002$) after the operation. The separate analysis of the mitochondrial and cytoplasmic calcium levels revealed similar significant differences after injury on postoperative days 7 ($P = 0.018$) and 14 ($P = 0.016$), and postoperative days 4 ($P = 0.018$) and 7 ($P = 0.0005$), respectively.

The PV contents in the hypoglossal and oculomotor nuclei were visualized by means of light microscopic immunocytochemistry. Whereas numerous cellular profiles with positive staining were observed in the hypoglossal nucleus of PV$^+/+$ animals (Fig. 7A), only background staining was seen in the same nucleus from the B6/SJL mice (Fig. 7B). The oculomotor nuclei from both strains contained stained cells and displayed strong PV staining in the neuropil (Fig. 7C, D). The intensity of the staining was expressed quantitatively on an arbitrary scale in each motor nucleus by determining the average net (background-corrected) amount of the chromophore (DAB). Compared with the control hypoglossal nucleus, a significant, 1.6-fold increase of immunostaining was detected in nucleus XII of the PV$^+/+$ mice (Table 1). Similar comparison of the oculomotor nuclei did not yield a significant difference (Table 1).

The increased intensity of PV immunostaining in the hypoglossal nucleus of PV$^+/+$ animals was attributed to the presence of PV-positive cellular profiles, which were missing in the nerve XII nucleus of B6/SJL mice. A frequency histogram of the profile diameter of the stained cells was constructed (Fig. 8A), which showed a range of 15–33 μm and a mean value of 21.8 ± 3.8 μm (mean ± s.e.m.) of the profile diameters. These numbers coincided well with the corresponding data available in the literature characterizing hypoglossal motor neurons (range: 11–34 μm; mean ± s.e.m.: 22.7 ± 0.68 μm) (Schwarz et al., 2009). Similar histograms have been constructed from the stained profiles visible in the oculomotor nuclei.
of both strains (Fig. 8B,C), which showed somewhat smaller mean diameter values (PV+/+: 15.7 ± 2.1 μm; B6/SJL: 16.4 ± 2.8 μm; mean ± s.d.), but were in agreement with the corresponding values from the literature describing oculomotor neurons (18.0 ± 3.3 μm; mean ± s.d.) (Eberhorn et al., 2006).

**DISCUSSION**

**Neuroprotective role of CaBPs**

CaBPs within the cells are also located at critical positions: in the vicinity of plasmalemmal calcium pumps and channels, around the mitochondria and endoplasmic reticulum, organelles known to play key roles in Ca²⁺

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**Figure 3.** A–E: Calcium distribution in motor neurons from the hypoglossal nucleus of B6/SJL animals. Arrows indicate electron-dense deposits reflecting the distribution of calcium. Motor neurons in cytoplasmic and mitochondrial locations from the nonoperated side (A) contain few deposits. The number of deposits gradually increases at postoperative day 1 (B) to days 4 (C), 7 (D), and 14 (E) and is lower on day 21 (F). ER, rough endoplasmic reticulum; Go, Golgi apparatus; mit, mitochondrion. Scale bar = 1 μm in F (applies to A–F).
signaling and buffering (Brini, 2003). Thus, these proteins may slow down the diffusion of calcium ions away from the influx regions, or facilitate their recycling into internal stores, and thereby reduce the amplitude of calcium concentration changes in the cytosol (Vanselow and Keller, 2000; von Lewinski and Keller, 2005; Müller et al., 2007). Because an elevated calcium level (Berridge et al., 1998; Arundine and Tymianski, 2003) and calcium-dependent mechanisms (Artal-Sanz and Tavernarakis, 2005) have often been implicated in acute injuries of the nervous system.

Figure 4. A–F: Calcium distribution in motor neurons from the hypoglossal nucleus of PV+/+ animals. Arrows indicate the reaction product of the histochemical procedure reflecting the distribution of the intracellular calcium. A small number of deposits can be seen in neurons from the nonoperated side (A) that, mainly within the mitochondria, were increased slightly on postoperative day 1 (B). At the subsequent time points, on postoperative days 4 (C), 7 (D), 14 (E), and 21 (F), the numbers of deposits are comparable with the control. ER, rough endoplasmic reticulum; Go, Golgi apparatus; mit, mitochondrion. Scale bar = 1 µm in F (applies to A–F).
system (Choi, 1995; Lipton, 1999; Raghupathi, 2004; Sullivan et al., 2005) or in neurodegenerative disorders (Mattson, 2007), as exemplified by Huntington’s disease (Bezprozvanny and Hayden, 2004), Alzheimer’s disease (Mattson and Chan, 2003), and ALS (Appel et al., 2001), it is a reasonable assumption that a larger buffering capacity of the intracellular calcium may confer a higher resistance against injury on the neurons via stabilization of the calcium homeostasis.

The concept that CaBPs may exert a general neuroprotective effect was proposed on the basis of data acquired from a wide range of in vitro (Mattson et al., 1991; Paizs et al., 1994) and in vivo models (Figure 5. A–F: Calcium distribution in motor neurons from the oculomotor nucleus of PV+/+ animals. Arrows indicate the electron-dense deposits that reflect the distribution of the intracellular calcium. A low number of deposits can be seen in the control condition (A), this pattern remaining practically the same on postoperative days 1 (B), 4 (C), 7 (D), 14 (E), and 21 (F). Go, Golgi apparatus; mit, mitochondrion. Scale bar = 1 μm in F (applies to A–F).)

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additional support from a variety of clinical observations, e.g., the age-dependent loss of CB from the cholinergic neurons of the basal forebrain in humans displayed a similar pattern to that of the loss of these cells in Alzheimer’s disease (Geula et al., 2003), whereas in Huntington’s disease those striatal neuronal groups that contained high levels of CB proved to have a decreased vulnerability (Mitchell and Griffiths, 2003). In Parkinson’s disease, the CB-positive dopaminergic neurons in the substantia nigra are relatively spared (Yamada et al., 1990). Finally, in ALS, the most susceptible motor neurons are comparatively deficient in CB and/or PV (Ince et al., 1993; Alexianu et al., 1994), which leads to the concise statement that “PV is a marker of ALS-resistant motor neurons” (Elliott and Snider, 1995).

However, not all studies lend support to the protective role of the CaBPs during injury. The reports range from documentation of the inability of a high CaBP content to confer resistance on the (Purkinje) cells in the cerebellum (Slemmer et al., 2005), through the lack of a correlation between the CB and/or PV content and the neuronal vulnerability in the hippocampus (Freund et al., 1990), to the finding that PV actually enhances N-methyl-D-aspartate (NMDA)-mediated toxicity (Hartley et al., 1996), or to the notion that not an excess, but rather a deficiency in CB might be protective (Klapstein et al., 1998). The diversity of these results, however, might be attributed to the fact that the actual impact of the CaBPs on survival is affected by the developmental state of the neuron (Lema Tome´ et al., 2006, 2008), shaped by the local cellular environment, and influenced by the neuronal network properties of the injured cells. In an effort to eliminate such perturbations of the results, in the present study, we decided to assay the changes in calcium level after injury in motor neurons with an artificially elevated PV content, and to compare the results obtained from the same population of neurons with a naturally low PV content in wild-type animals.

**Lesion induced by axotomy**

Axonal transection is an experimental method for the study of neuronal responses to injury leading to degeneration, or resulting in plastic changes and recovery (Price et al., 1992; Koliatsos and Price, 1996). The lesion of motor axons induces retrograde changes in neurons (Lieberman, 1971), some of which resemble those seen in ALS (Price et al., 1994). Factors known to influence the outcome of axotomy-induced injury are the type of lesion (crush, cut, target deprivation, or avulsion), the location of the trauma relative to the perikaryon, differences in species, type of the motor neurons, and age of the animals (Snider et al., 1992; Koliatsos and Price, 1996).

For example, approximately 70% of adult hypoglossal motor neurons survive target deprivation in rats, but only...
40% survive if the surgery is made during the first postnatal week (Snider and Thanedar, 1989). In the spinal cord of newborn animals, whereas avulsion or crush of the sciatic nerve produces a 73–80% loss of lumbar motor neurons (Schmalbruch, 1984; Koliatsos and Price, 1996; Dekkers et al., 2004), axotomy of the same nerve

**Figure 7.** A–D: Parvalbumin staining of the hypoglossal (A,B) and oculomotor nuclei (C,D) of PV+/+ (A,C) and B6/SJL (B,D) mice. In the hypoglossal nucleus of the PV+/+ animals, numerous stained cellular profiles are visible, which, according to their size, shape, and distribution, can be considered as motor neurons (A). In the hypoglossal nucleus of the B6/SJL mice only background staining can be seen (B). The neuropil staining and some stained cellular profiles at the ventromedial and lateral border of the hypoglossal nucleus in both figures may belong to the nucleus of Roller and the intercalated nucleus of medulla, respectively. The parvalbumin staining patterns of the oculomotor nuclei of the PV+/+ (C) and B6/SJL (D) mice are similar: in the heavily stained neuropil, stained cellular profiles are present. mlf, medial longitudinal fasciculus; ml, midline. Scale bar = 100 μm in A–D.
induces the complete loss of motor neurons (Schmalbruch, 1984). In contrast, sciatic nerve cut in adult animals generates no visible loss of motor neurons (Schmalbruch, 1984). It seems to be a general rule, however, that motor neurons become less vulnerable to axotomy by the time the animals have attained the age of 1 month: whereas the majority of motor neurons in adults can survive target deprivation for prolonged periods, in postnatal animals the proportion of the surviving population is merely 25–35%, or less (Koliatsos and Price, 1996).

In view of the fact that the majority of motor neurons in newborn animals do not survive injury to their axons, this model appears to be sufficiently attractive and sensitive to assay the effect of PV upregulation on the altered vulnerability of these cells. Indeed, in a recent publication, although the calcium level changes were not monitored, the higher PV level was documented to have a positive effect on the survival of neonatal spinal motor neurons after a sciatic nerve crush (Dekkers et al., 2004). However, in that experiment no re-establishment of the motor functions was observed, which might be attributed to the surviving population persisting in some altered form (Martin et al., 1998), as interference with the developmental regulation of CaBP expression during the early postnatal phase may have adverse effects (Clowry et al., 1997). In the present experiments, therefore, we performed the surgical interventions in animals 12 weeks of age, when the adult response was already present.

Calcium level changes induced by axotomy

The lesion of the hypoglossal nerve of the B6/SJL animals was followed by a transient increase in the intracellular calcium in the corresponding motor neurons, with a peak increase of approximately 1.5-fold around days 7–14 (Fig. 6A). In the same hypoglossal motor neurons of the PV+/+ animals, apart from a slight, insignificant increase in the mitochondrial calcium on day 1 after surgery, no increase was detected relative to the calcium level in the motor neurons on the nonoperated side at any time point postoperatively (Fig. 6A–C). Likewise, no calcium increase was seen in the motor neurons of the oculomotor nuclei of the PV+/+ mice after axotomy, and no significant difference could be discerned statistically between the two examined motor nuclei of the PV+/+ animals. Thus, as concerns the intracellular calcium increase after injury, the PV upregulation in these animals lends resistance of “oculomotor type” to the hypoglossal motor neurons.

Demonstration of tissue calcium by electron microscopy

The empirical approach used to gain information on the fine structural distribution of calcium is based on a cytochemical reaction to precipitate and visualize calcium as electron-dense deposits at the electron microscope. Such techniques, based on either the oxalate-pyroantimonate- (Borgers et al., 1977, 1981), the bichromate-
Relevance of CaBPs in ALS

The suggestion that CaBPs may protect motor neurons in ALS was based on the parallelism observed between the relatively spared motor functions in patients, such as the eye movement and voluntary sphincter functions (Bergmann et al., 1995), and the elevated CaBP contents of the corresponding motor neurons (Ince et al., 1993; Alexianu et al., 1994; Elliott and Snider, 1995). These clinical and pathological data led to the potential of CaBPs to rescue motor neurons being extensively studied in vitro in experiments mimicking stress conditions known to contribute to their degeneration during ALS, such as immunemediated (Ho et al., 1996), excitotoxicity-induced (Van Den Bosch et al., 2002), and oxidative injury (Roy et al., 1998). Although cultured cells do not possess the cellular milieu that is present in the nervous tissue, which is known to affect the degree of the lesion, in these experiments the protective role of the CaBPs was verified directly.

Ex vivo arrangements (either acute or organotypic slices, or whole mount neuromuscular preparations) integrate at least some of the natural environment of the motor neurons, with the consequence that the outcome of such experiments can be translated more reliably to protective strategies, or to conclusions relevant to the pathomechanism of the disease. The results of electrophysiology experiments on tissue slices containing either the oculomotor (Vanselow and Keller, 2000) or the hypoglossal nucleus (Lips and Keller, 1998; Ladewig et al., 2003) or spinal motor neurons (Palecek et al., 1999) have provided further support for the idea that a higher calcium buffer capacity, manifested by a higher CaBP content, affords resistance against calcium-mediated injury in the motor neurons (von Lewinski and Keller, 2005). These data accorded with the results of miniature end-plate potential measurements on whole mount neuromuscular preparations made from muscles containing vulnerable and resistant types of synapses (Mosier et al., 2000). The drawback of these experiments, however, is the fact that they do not sample motor neurons from the same location and, thus the results do not reflect the effects of the CaBPs alone, but may be influenced by the environmental differences.

The most widely used model of ALS suitable for in vivo research is a variety of transgenic mice developed on the basis of mutations of the enzyme SOD1 detected in a...
subpopulation of ALS patients (Deng et al., 1993; Rosen et al., 1993; Gurney et al., 1994). By using transgenic animals based on the G93A mutation of SOD1, we earlier demonstrated not only that changes in the intracellular calcium level paralleled the degeneration of the motor neurons, but also that the survival of the motor neurons was associated with higher levels of CB and/or PV (Siklós et al., 1998). Because those experiments still involved the comparison of different motor neurons, (spinal versus ocu-
lomotor neurons), it was decided to develop a mouse line with an elevated PV content, suitable for measuring the protective effect of CaBPs in the same neuronal population in vivo (Beers et al., 2001). When PV+/+ transgenic mice were bred to the mutant SOD1 animals, a significantly increased preservation of the spinal motor neurons and a delayed onset of the disease was noted as compared with the mice containing only the SOD1 transgene.

Although the survival of the double transgenic mice was also considerably improved, the animals could not be ultimately rescued. The lack of success in preventing the death of mutant SOD1 animals by the upregulation of PV in the motor neurons could be a consequence either of the CaBPs acting solely as calcium buffers, which could be saturated if the stress signal endures (Siklós and Appel, 2005), or of the effect of the neighboring cells contributing to the damage (Boilée et al., 2006; Julien, 2007; Obal et al., 2001). To test directly whether an elevated PV level attenuates the calcium increases in the motor neurons (Koliatsos and Price, 1996). As expected, a non-
significant calcium level increase was demonstrated in the hypoglossal motor neurons of PV+/+ mice, which was similar to that measured in the oculomotor neurons.

On the basis of this concept, the maximum goal would be a delay of the injury of the vulnerable motor neurons and a provision of a neuroprotection similar to that present in the relatively resistant motor neurons (Bergmann et al., 1995). The modest protective effect of the PV upregulation in the PV × SOD1 motor neurons was unexpected. Because an increased intracellular calcium level appears to be a key factor in the coupling of different pathways to cell destruction in models of ALS (Appel et al., 2001), the failure to prevent motor neuron death completely in the animals could be a consequence of fail-
ing to address all the cellular processes, or critical molecular steps of the destructive processes (Moisse and Strong, 2006; Cozzolino et al., 2008; Rothstein, 2009). This clearly warrants further studies in PV+/+ animals after the induction of similar lesions in order to assess the contributions of nonmotor neuronal compartments to survival or degeneration.

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