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## Changes in antioxidant enzyme expression in response to hydrogen peroxide in rat astroglial cells

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**Abstract** Oxidative stress has been causally linked to a variety of neurodegenerative diseases. To clarify the role of the antioxidant enzyme (AOE) system in oxidative brain damage primary cultures of rat astroglial cells were exposed to hydrogen peroxide ( $H_2O_2$ ). Expression of AOE and several parameters for cell viability and functionality were measured. In our experiments astrocytes responded to low concentrations of  $H_2O_2$  exposure with a pronounced generation of ROS which ran parallel with induction of lipid peroxidation. This distinct oxidative stress was not reflected in cell viability or functionality parameters measured. Cytotoxicity, a decrease in glutathione content of astrocytes, and impairment of mitochondrial functions became obvious only for higher concentrations of  $H_2O_2$ . After  $H_2O_2$  exposure catalase, manganese superoxide dismutase, and glutathione peroxidase expression levels were found to be increased, whereas copper/zinc superoxide dismutase mRNA expression was not affected. These data indicate that the AOE system of astrocytes can be directly regulated by oxidative stress and may thus contribute to protection of cells against oxidative insults.

**Keywords** Catalase · Superoxide dismutase · Glutathione peroxidase · Hydrogen peroxide · Astrocytes

### Introduction

Oxidative stress is believed to be implicated in a wide variety of human degenerative disorders of the central nervous system (CNS), including Alzheimer's disease, Parkinson's disease, and amyotrophic lateral sclerosis, and conditions such as ischemia and excitotoxicity (Halliwell 1992). To prevent injury from reactive oxygen species (ROS), aerobic organisms are endowed with defense systems including both specific enzymes and scavenger molecules, such as glutathione or  $\alpha$ -tocopherol. Included among the antioxidant enzymes (AOEs) are catalase, copper/zinc superoxide dismutase (CuZnSOD), manganese superoxide dismutase (MnSOD), and the glutathione peroxidases (GPxs). Catalase and GPxs convert  $H_2O_2$  to  $H_2O$  and the SODs catalyze the dismutation of the superoxide radical anion (Aebi 1984; McCord 1979; Ursini et al. 1995).

The CNS is generally considered to be especially sensitive to free radical damage due to its high metabolic rate, high levels of lipids, and the comparatively low content of protective systems (Halliwell and Gutteridge 1989). The antioxidant defense via the AOE is particularly important for the brain. An increased activity of CuZnSOD in the brains of transgenic mice protects neurons from ischemia reperfusion injury and also lessens the cerebral edema caused by traumatic brain injury (Li et al. 1999). Conversely, a deficiency in CuZnSOD or MnSOD worsens the brain damage (Chan 1996; Kondo et al. 1996). Familial amyotrophic lateral sclerosis is associated with mutations in the CuZnSOD gene, but the mechanisms by which these mutations lead to amyotrophic lateral sclerosis are still unknown (Rabizadeh et al. 1995).

The expression of AOE can be regulated by oxidative stress itself. Exposure to  $H_2O_2$  or hyperoxia led to an increase in catalase mRNA expression in various epithelial cell lines as well as in the rat lung (Clerch et al. 1990; Shull et al. 1991; Tate et al. 1995). Previous results

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from our laboratory have shown differential regulation of AOE expression in rat hepatocyte cultures and hepatoma cells (Röhrdanz and Kahl 1998; Röhrdanz et al. 2000). Inflammatory mediators are able to induce selectively MnSOD mRNA expression in different cell types, among others in primary cortical neurons and astrocytes (Dougall and Nick 1991; Kifle et al. 1996; Mokuno et al. 1994; Wong and Goeddel 1984). Since inflammation is assumed to act via generation of ROS, induction of AOE is thus linked to oxidative stress. Pinteaux et al. (1996) found an increase of MnSOD activity and a decrease in CuZnSOD activity after exposure of primary glial cell cultures to ROS generated by a xanthine/xanthine oxidase mixture. However, if and how the mRNA expression of those enzymes was affected has not been shown.

Astroglial cells represent the major compartment of GPx in the human mesencephalon and hippocampus (Damier et al. 1993; Takizawa et al. 1994). For MnSOD and CuZnSOD a neuronal localization in the human brain is discussed (Akai et al. 1990; Bergeron et al. 1996; Ceballos et al. 1989; Zhang et al. 1994). Changes of the cellular localization of SODs were observed in cases of Alzheimer's disease and Down's syndrome (Ceballos-Picot 1997; Furuta et al. 1995; Pinteaux et al. 1998). In cerebral sites of degeneration MnSOD was more enriched in reactive astrocytes than in neurons. This suggests that MnSOD may be activated by oxidative stress as a defense mechanism against deleterious effects of ROS. Neuroprotection through astrocytes was also shown in primary cultures of astrocytes and neurons. Co-culture with astrocytes attenuated killing by H<sub>2</sub>O<sub>2</sub> of striatal neurons (Desagher et al. 1996). This neuroprotective role of astrocytes appears most evident following reactive gliosis, where different antioxidant mechanisms were found to be stimulated.

To elucidate the role of AOE in oxidative brain damage, we investigated the influence of oxidative stress on AOE expression. In our model system primary cultures of astroglial cells were treated with H<sub>2</sub>O<sub>2</sub> as a direct mediator of oxidative stress.

## Materials and methods

### Preparation of primary rat astroglial cultures

Astrocytes were isolated from the whole rat fetal brain (E19–20). The brains were filtered through two different Nylon meshes (212 and 135 µm pore diameter) as previously described (Schmuck and Haynes 2000). The cells were cultured in DMEM medium supplemented with 10% FCS on Primaria culture plates (Becton and Dickinson, Heidelberg, Germany) in a humidified incubator in 5% CO<sub>2</sub>/95% air. The astrocytes consisted of around 95% glial fibrillary acid protein (Roche, Mannheim, Germany)-positive cells, 1% microglia (CD11/B/C-positive), and 1–5% oligodendrocytes (galactocerebroside-positive). The cultures were grown for 10–14 days with two changes of medium in between at 37°C before use. When not stated otherwise astrocytes were treated for 24 h with different concentrations of H<sub>2</sub>O<sub>2</sub> to perform the experiments.

### Viability assay

To determine the viability of H<sub>2</sub>O<sub>2</sub>-treated cells, mitochondrial dehydrogenase activity was measured in the MTT assay as described elsewhere (Carmichael et al. 1985).

### Intracellular ATP concentration

The intracellular ATP concentration was determined according to a chemiluminescence reaction in a luciferin/luciferase system (ATP Lite-M; Packard, Groningen, The Netherlands). Astrocytes were treated for 24 h with H<sub>2</sub>O<sub>2</sub> before they were tested.

### Mitochondrial membrane potential

The membrane potential of the inner mitochondrial membrane was measured by the dye tetramethylrhodamine after treating the astrocyte cultures for 24 h with H<sub>2</sub>O<sub>2</sub>. The dye was added to the cell culture medium in a concentration of 3.3 µM for 30 min at the end of the incubation period with H<sub>2</sub>O<sub>2</sub>. The cells were washed twice and the fluorescence was determined with a fluorostar spectrophotometer (SLT, Crailsheim, Germany) at 530/580 nm.

### Detection of ROS

The determination of ROS was done according to Oyama et al. (1996). The cells were washed with PBS and loaded with 2,7-dichlorofluorescein diacetate (Sigma, Deisenhofen, Germany) at a concentration of 50 µM in phosphate-buffered saline (PBS) for 1 h. Dichlorofluorescein diacetate crosses cell membranes and then undergoes deacetylation by intracellular esterases. The resulting compound, dichlorofluorescein, is proposed to be trapped within the cell and susceptible to ROS-mediated oxidation to the fluorescent compound, dichlorofluorescein (Buxser et al. 1999). After 1 h the remaining dichlorofluorescein diacetate was removed, the cells were washed with PBS, and then treated with the indicated concentrations of H<sub>2</sub>O<sub>2</sub>. The fluorescence was determined up to 4 h in a fluorostar spectrophotometer (SLT) at 485/530 nm.

### Lipid peroxidation

Lipid peroxidation was measured by quantitation of malondialdehyde (MDA) release of treated cells into the medium. MDA was determined after reaction of cell culture supernatants with thiobarbituric acid and subsequent high-pressure liquid chromatography according to a modified method by Draper et al. (1993).

### Intracellular glutathione (GSH)

Intracellular GSH concentration was measured according to Mundi et al. (1997). The cells were washed with PBS and then treated with the indicated concentrations of H<sub>2</sub>O<sub>2</sub> for 1 h. Afterwards the cells were loaded with 5-methylchlorofluorescein diacetate (Molecular Probes, Eugene, Ore., USA) in a concentration of 20 µM in PBS for 1 h. Then the fluorescence was determined in a fluorostar spectrophotometer (SLT) at 485/530 nm.

### RNA isolation and RNA analysis

Total RNA was isolated from cells using Trizol reagent (Gibco BRL, Eggenstein, Germany). For northern blot analysis, 5 µg total RNA were resolved by electrophoresis in a 1% agarose, 2.25 mM formaldehyde gel in a running buffer containing 20 mM MOPS, pH 7, 0.5 mM sodium acetate, and 1 mM EDTA. RNA was transferred to nylon membrane (Amersham, Arlington Height, Ill., USA) according to Maniatis et al. (1982). Purified cDNAs were labeled with [ $\alpha$ -<sup>32</sup>P] dCTP (111 TBq/mmol; Hartmann Analytic, Braunschweig, Germany) by random hexamer priming (Roche

Diagnosics, Mannheim, Germany). Blots were prehybridized and hybridized with cDNAs as described previously (Röhrdanz and Kahl 1998). Autoradiographs were made by exposing blots to X-ray film (Kodak XAR) with an intensifying screen at  $-80^{\circ}\text{C}$ . Blots were stripped and reprobed with different cDNAs. Autoradiographs were analyzed by densitometric scanning using the Quantity One system from BioRad (Munich, Germany).

Dr. Shuichi Furuta (Sinshu University, Department of Biochemistry, Nagano, Japan) provided the rat catalase cDNA containing plasmid, PMJ1010 (Furuta et al. 1986). We obtained a 1.4 kb rat MnSOD cDNA containing plasmid, pSP65-RMS (Ho and Crapo 1987a), and a 0.6 kb rat CuZnSOD cDNA containing plasmid, pUC-13-RCS (Ho and Crapo 1987b), from Dr. Ye-Shih Ho (Wayne State University, Institute of Chemical Toxicology, Detroit, Mich., USA). For 18 S rRNA detection we used the oligonucleotide probe 5'-GCCGTGCGTACTTAGACATGCATG-3' (Chan et al. 1984).

PCR following reverse transcription was performed for semi-quantitative determination of GPx mRNA. One microgram of total RNA was transcribed into cDNA in a 25- $\mu\text{l}$  final volume of reaction buffer (50 mM TRIS-HCL, 75 mM KCl, 3 mM  $\text{MgCl}_2$ , 10 mM DTT, 0.5 mM each dNTP) and 5  $\mu\text{M}$  oligo-d(T)<sub>16</sub>-primer, 1 U RNase inhibitor, and 2.5 U MLV reverse transcriptase by incubation for 1 h at  $42^{\circ}\text{C}$ . The reaction was stopped by incubation at  $99^{\circ}\text{C}$  for 5 min. For rat GPx PCR 2.5  $\mu\text{l}$  of the product of reverse transcription was added to a 22.5- $\mu\text{l}$  PCR mixture containing 10 mM TRIS-HCL, 50 mM KCl, 0.1% Triton X-100, 2 mM  $\text{MgCl}_2$ , 0.2 mM each dNTP, 1 U Taq DNA polymerase (Promega, Mannheim, Germany), and 0.2  $\mu\text{M}$  each PCR primer. The primers were designed for detection of the rat GPx according to the genomic sequence published by Ho and Howard (1992); upstream primer 5'-ATGTCTGCTGCTCGGCTCTC-3' and downstream primer 5'-GTTGCTAGGCTGCTTGGACAG-3'. Amplification was initiated with 4 min denaturation at  $94^{\circ}\text{C}$  followed by 27 cycles at  $94^{\circ}\text{C}$ ,  $61^{\circ}\text{C}$ , and  $72^{\circ}\text{C}$  for 60 s. Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) PCR was performed with 100 ng synthesized cDNA as described elsewhere (El-Bahay et al. 1999). The amplified PCR products were 602 bp for GPx mRNA and 450 bp for GAPDH mRNA. The conditions of PCR were set to be in the linear phase of amplification to allow for semiquantification of mRNA content. Five microliters from each PCR reaction were electrophoresed in a 1% agarose gel in TRIS-borate-EDTA buffer. The cDNA bands were visualized by UV illumination after staining the gels with ethidium bromide. Gels were photographed and scanned densitometrically.

#### Statistics

Statistical analysis was performed using Student's *t*-test (Sigma Stat; Scientific, Erkrath, Germany).

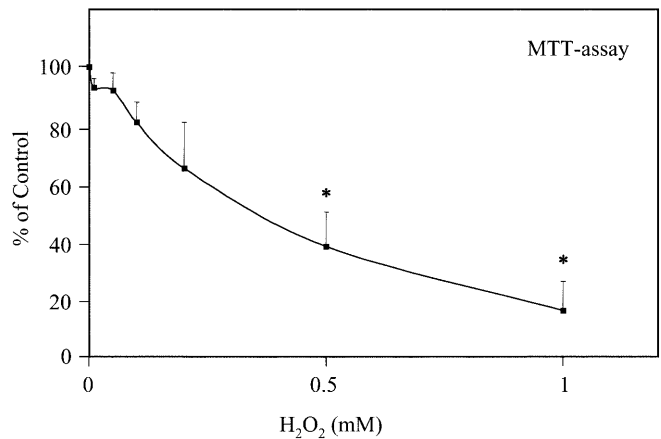
## Results

### Cytotoxicity

After  $\text{H}_2\text{O}_2$  exposure, viability of cells was determined via the MTT assay, which reflects mitochondrial dehydrogenase activity (Fig. 1). According to the MTT assay,  $\text{H}_2\text{O}_2$  toxicity started to show at concentrations of 0.05–0.1 mM  $\text{H}_2\text{O}_2$  with around 20% dead astrocytes, increasing to 80% dead cells at 1 mM  $\text{H}_2\text{O}_2$ .

### Mitochondrial function

Mitochondrial function was determined by the intracellular ATP concentration and the mitochondrial



**Fig. 1** Viability of astrocytes after treatment with  $\text{H}_2\text{O}_2$ . Astrocytes were incubated with the indicated concentrations of  $\text{H}_2\text{O}_2$  for 24 h and viability was determined via MTT assay. Values are expressed as percentage of controls (means  $\pm$  SEM from four to six different experiments). \* $P < 0.05$  vs controls

membrane potential (tetramethylrhodamine) (Fig. 2). There was a reduction in ATP content, which became obvious at a concentration of 0.5 mM  $\text{H}_2\text{O}_2$ . At 0.5 mM  $\text{H}_2\text{O}_2$  a 30% and at 1 mM  $\text{H}_2\text{O}_2$  a 40% reduction was detected. These results are confirmed by the loss in membrane potential with increasing concentrations of  $\text{H}_2\text{O}_2$ . Treatment of astrocytes with 0.5 mM  $\text{H}_2\text{O}_2$  lead to a loss of 50% and with 1 mM  $\text{H}_2\text{O}_2$  to a loss of 80% in membrane potential compared to controls, which ran parallel with the decrease in cell viability.

### Generation of ROS

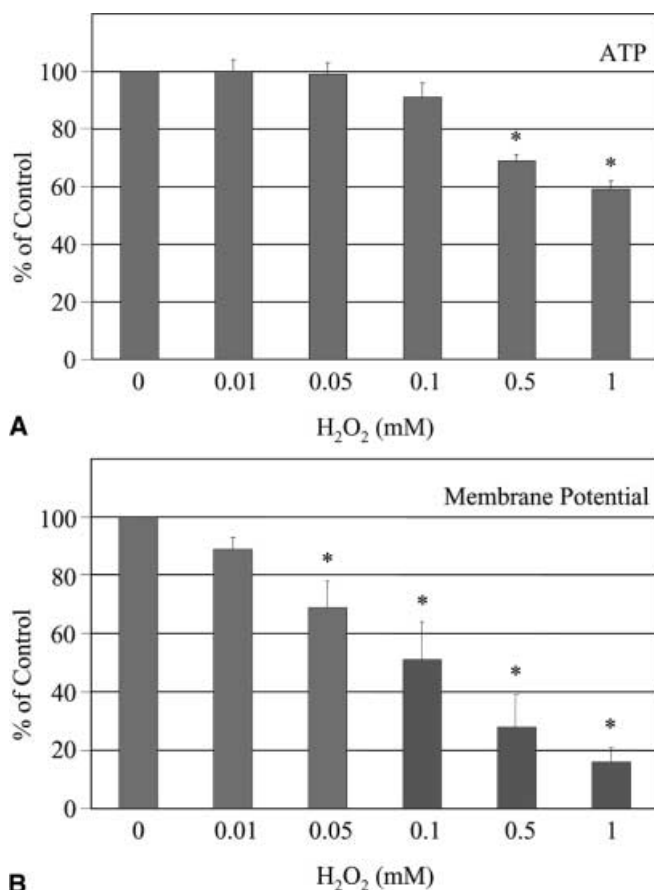
Upon treatment with  $\text{H}_2\text{O}_2$  a massive production of ROS could be detected in astrocytes (Fig. 3). A seven-fold increase in ROS compared to controls could already be detected for low concentrations of  $\text{H}_2\text{O}_2$  ( $\leq 0.1$  mM) after treatment for 60 and 120 min, respectively.

### Lipid peroxidation

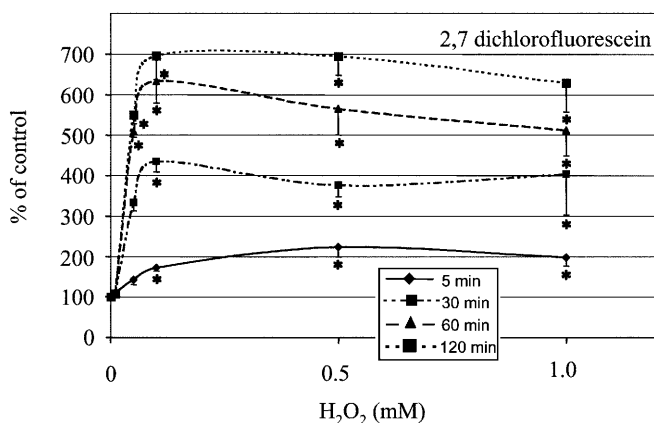
The extent of lipid peroxidation was assessed by measuring the release of the breakdown product MDA. In astrocytes an increase by the factor of 6–7 in MDA levels was obtained after treatment with  $\text{H}_2\text{O}_2$ . This drastic increase in MDA levels already started to show with  $\text{H}_2\text{O}_2$  concentrations of 0.05 mM and corresponded to the ROS production data (Fig. 4).

### Intracellular GSH

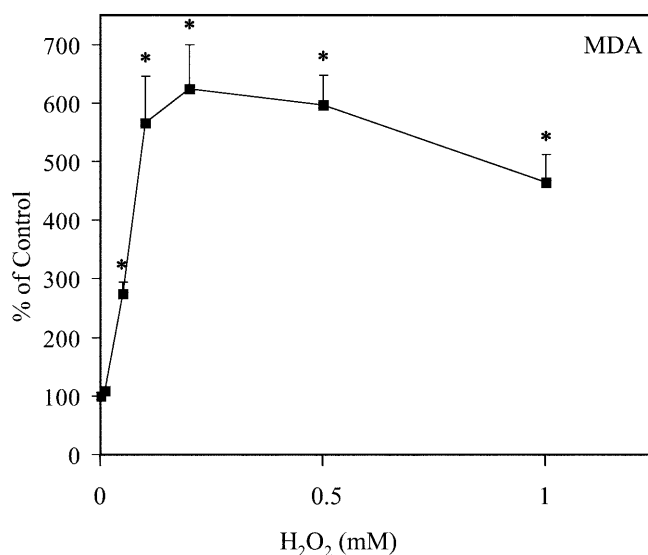
With increasing concentrations of  $\text{H}_2\text{O}_2$ , the internal stores of GSH were depleted in astrocytes (Fig. 5). A nearly 20% depletion of GSH was observed after



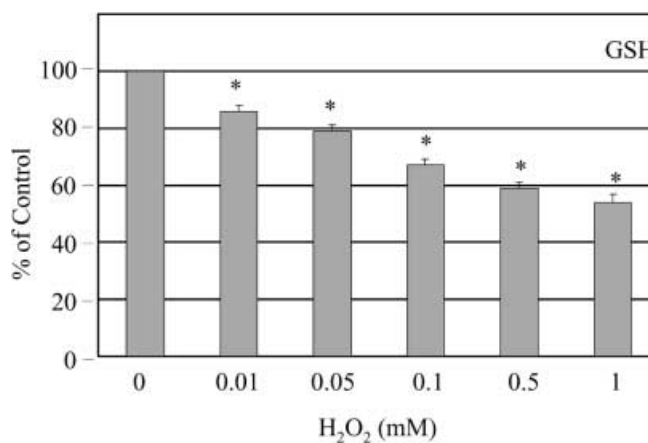
**Fig. 2A, B** Determination of effects on mitochondrial functions by H<sub>2</sub>O<sub>2</sub> in astrocyte cell cultures. Mitochondrial functions were determined after treatment of astrocytes with H<sub>2</sub>O<sub>2</sub> for 24 h. **A** Intracellular ATP assessed via chemiluminescence reaction. **B** Mitochondrial membrane potential measured by the dye tetramethylrhodamine. Values are expressed as percentage of controls (means  $\pm$  SEM from four different experiments). \* $P$  < 0.05 vs controls



**Fig. 3** Production of reactive oxygen species (ROS) by H<sub>2</sub>O<sub>2</sub> in astrocyte cell cultures. Astrocytes were exposed to the indicated concentrations of H<sub>2</sub>O<sub>2</sub> and ROS production was measured at different times thereafter. Values are expressed as percentage of controls (means  $\pm$  SEM from four to six different experiments). \* $P$  < 0.05 vs controls



**Fig. 4** Formation of malondialdehyde (MDA) after treatment with H<sub>2</sub>O<sub>2</sub>. Astrocyte cultures were incubated with the indicated concentrations of H<sub>2</sub>O<sub>2</sub> for 24 h and formation of MDA was assessed. The results are expressed as percentage of controls (means  $\pm$  SEM from five or six different experiments). \* $P$  < 0.05 vs controls

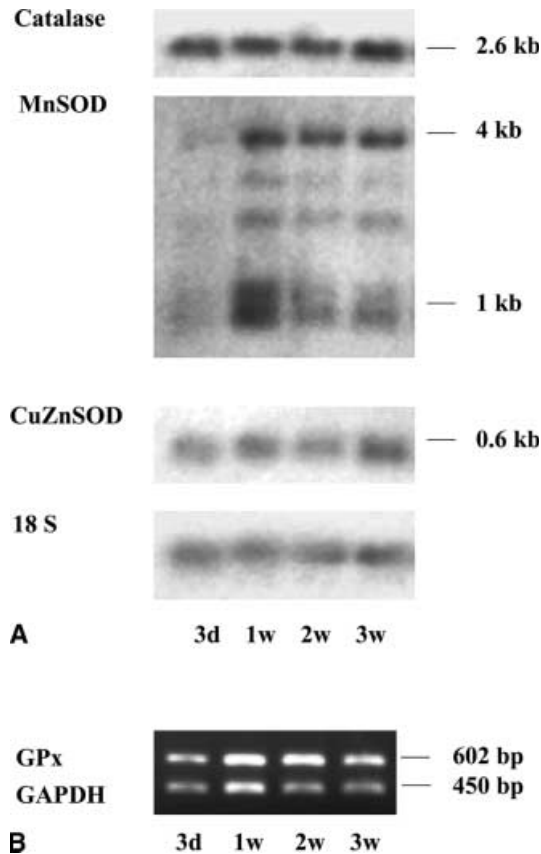


**Fig. 5** Determination of intracellular glutathione (GSH) concentration. Rat astrocytes were treated with the indicated concentrations of H<sub>2</sub>O<sub>2</sub> for 1 h. GSH concentrations were determined using a fluorescent dye. The results are expressed as percentage of controls (means  $\pm$  SEM from four different experiments). \* $P$  < 0.05 vs controls

treatment with 0.01 mM H<sub>2</sub>O<sub>2</sub> increasing to nearly 50% of GSH depletion at 1 mM H<sub>2</sub>O<sub>2</sub>.

#### Expression of AOE in astrocytes during cell culture

The basal expression levels of catalase, MnSOD, and CuZnSOD during culturing are shown in Fig. 6A. The amount of 18 S rRNA serves as internal loading control. mRNA expression of GPx was determined via semi-quantitative PCR with GAPDH mRNA expression as



**Fig. 6A, B** Basal expression levels of catalase, manganese superoxide dismutase (MnSOD), copper/zinc superoxide dismutase (CuZnSOD), and glutathione peroxidase (GPx) during culturing in primary astrocyte cell cultures. **A** Shown is a representative northern blot analysis for catalase, MnSOD, and CuZnSOD, and as internal loading control the 18 S rRNA expression. **B** Shown is a representative PCR analysis for GPx expression with glyceraldehyde-3-phosphate dehydrogenase (GAPDH) expression as internal standard

internal standard (Fig. 6B). In astrocytes expression of catalase, CuZnSOD, and GPx did not change during cell culture. Expression of MnSOD showed some changes during culturing; whereas at day 3 MnSOD expression was barely detectable, it shows a peak after 1 week of culturing and a stable expression pattern thereafter. Accordingly astrocytes were used for experiments at 2 weeks of culturing when a stable expression pattern of all AOE was reached.

#### Effect of H<sub>2</sub>O<sub>2</sub> on AOE expression in astrocytes

Astrocytes were cultured for 2 weeks and then exposed to different concentrations of H<sub>2</sub>O<sub>2</sub>. Figure 7A shows the northern analysis and Fig. 7B the densitometric data relative to 18 S rRNA. In Fig. 8A the PCR for detection of GPx mRNA expression is depicted and in Fig. 8B the densitometric data relative to GAPDH mRNA expression. To obtain densitometric data for MnSOD mRNA expression all five bands of MnSOD detected by

northern blotting were quantitated together, because no changes in between this expression pattern was observed (Hurt et al. 1992). There was an increase in MnSOD, catalase, and GPx mRNA expression levels after treatment with H<sub>2</sub>O<sub>2</sub>. A twofold increase of MnSOD expression levels could be shown starting at a concentration of 0.05 mM H<sub>2</sub>O<sub>2</sub>. Catalase mRNA expression was elevated around 2 times only at the highest concentration of 0.2 mM H<sub>2</sub>O<sub>2</sub>. The same was found for GPx mRNA expression, a slight but significant increase of 1.5 times after treatment with 0.2 mM H<sub>2</sub>O<sub>2</sub>. No changes in CuZnSOD mRNA expression was detected after H<sub>2</sub>O<sub>2</sub> exposure.

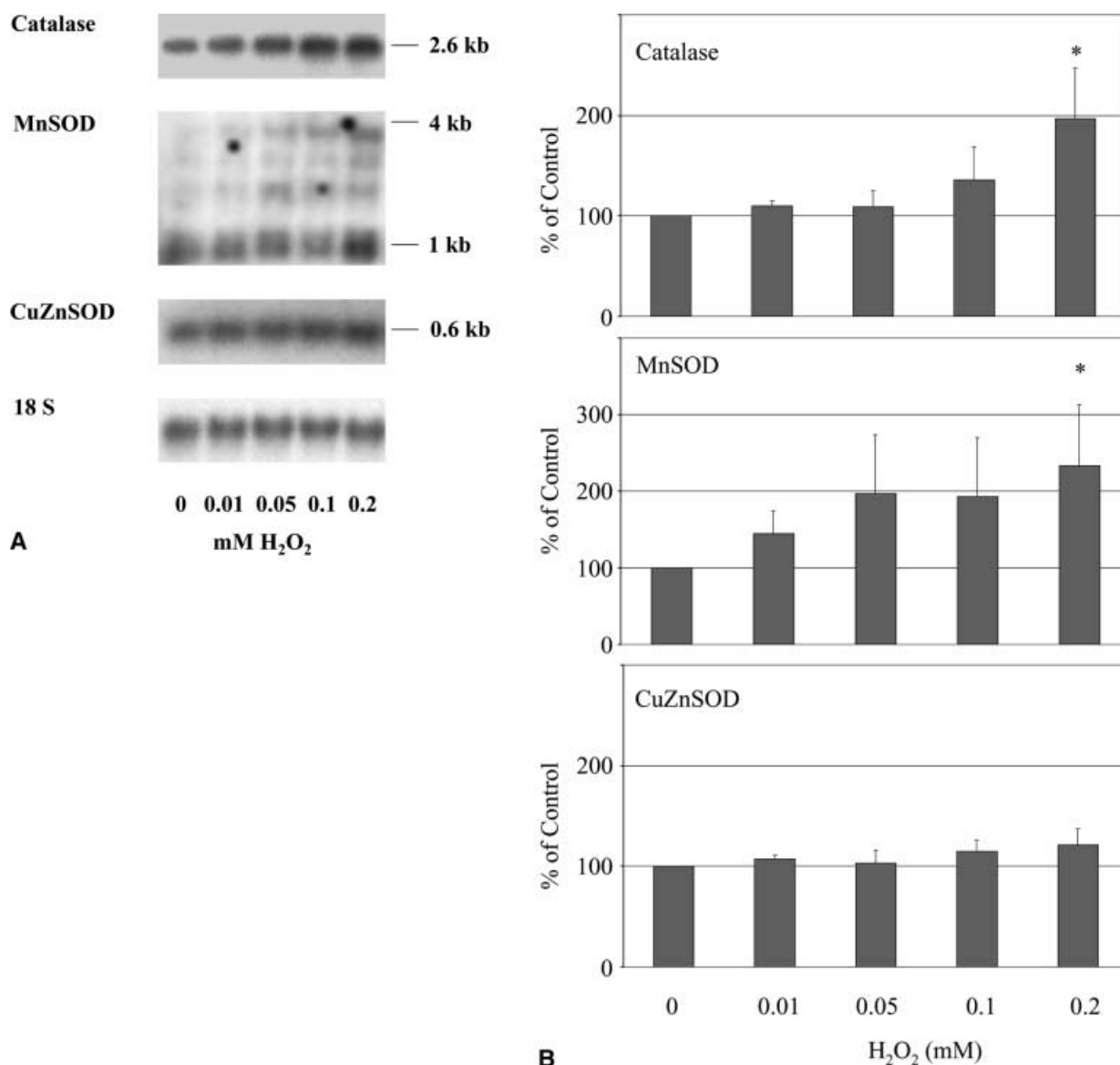
#### Discussion

The brain is particularly vulnerable to oxidative stress because of its high rate of oxygen consumption, the high levels of polyunsaturated fatty acids as substrates for lipid peroxidation, and the essentially non-regenerative nature of neurons (Halliwell 1992; Halliwell and Gutteridge 1989). The AOE system is very important for the defense against such oxidative damage. Changes in AOE expression caused by ROS may be crucial for cell survival.

After treatment of astrocytes with H<sub>2</sub>O<sub>2</sub>, a massive increase in intracellular oxidation paralleled by lipid peroxidation could be detected. The ability of astrocytes to respond toward an oxidative insult by the production of MDA was also described by Schroeter et al. (1999), who detected an increase in MDA levels after hyperoxia/reoxygenation. Caf e et al. (1995) found lipid peroxidation products in astroglial cell cultures after iron-stimulated oxidative stress. We detected high levels of ROS and MDA already at low concentrations of H<sub>2</sub>O<sub>2</sub>.

Viability was only slightly impaired at a concentration of 0.1 mM H<sub>2</sub>O<sub>2</sub>, when ROS concentration and MDA release were increased sevenfold. Cell damage became more severe at much higher concentrations. Treatment with 1 mM H<sub>2</sub>O<sub>2</sub> resulted in 60% dead cells. Several groups tested the neurotoxic potential of H<sub>2</sub>O<sub>2</sub>, mostly with emphasis on differences in cytotoxicity between neuronal and astroglial cultures (Abe and Saito 1998; Desagher et al. 1996; Langeveld et al. 1995). Astrocytes were always detected as the more resistant cell type toward H<sub>2</sub>O<sub>2</sub> toxicity. Therefore a neuroprotective role was attributed to astrocytes (Desagher et al. 1996; Wilson 1997). Similar toxicity data like ours were obtained in those studies. Abe and Saito (1996) found a steeper dose response curve probably resulting from different conditions in cell culturing.

Mitochondrial functions, as measured by ATP consumption and the mitochondrial membrane potential, decreased in parallel with cell viability. The same was observed for GSH depletion in astrocytes after H<sub>2</sub>O<sub>2</sub> exposure. Impairment of mitochondrial functions result in deficiencies of energy metabolism thus contributing to cell death. The importance of the GSH/GSSG ratio in

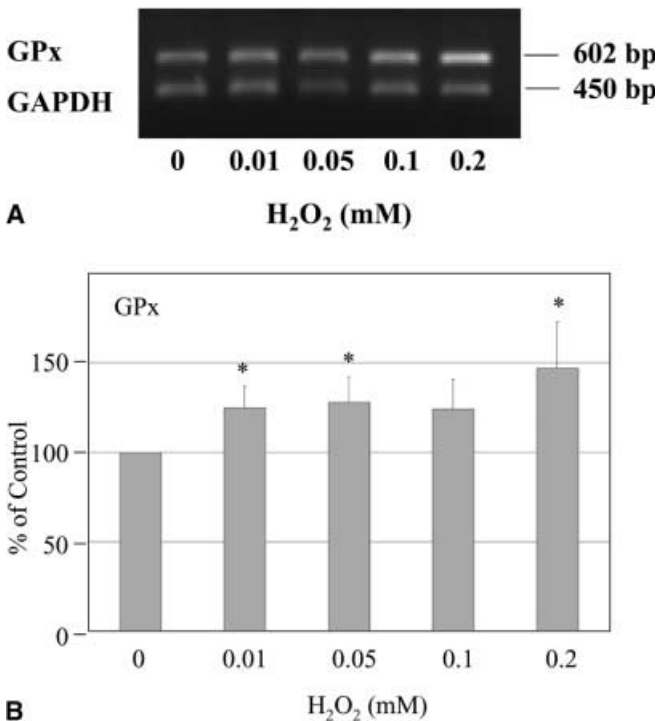


**Fig. 7A, B** Gene expression of catalase, MnSOD, and CuZnSOD in primary astrocyte cultures after treatment with H<sub>2</sub>O<sub>2</sub>. **A** Northern blot analysis of catalase, MnSOD, and CuZnSOD mRNAs. Two-week cultures of astrocytes were exposed to the indicated concentrations of H<sub>2</sub>O<sub>2</sub> for 24 h. Northern analysis was carried out using rat catalase, MnSOD, or CuZnSOD cDNA as a probe. The northern analysis shown is representative of three to six individual experiments. **B** Quantitative densitometric data for catalase, MnSOD, and CuZnSOD mRNA expression levels. mRNA levels of catalase, MnSOD, and CuZnSOD were quantitated relative to 18 S rRNA levels shown in the *lowest panel* of **A**. Values are expressed as percentage of controls (means ± SEM from three to six different experiments). \**P* < 0.05 vs controls

astrocytes for detoxification of H<sub>2</sub>O<sub>2</sub> was investigated by Dringen and Hamprecht (1997). Immediately after administration of H<sub>2</sub>O<sub>2</sub> to astroglial cells GSH was found to be rapidly oxidized indicating that GPx and GSH are involved in astroglial detoxification of exogenously applied H<sub>2</sub>O<sub>2</sub>. The pentose phosphate pathway seemed to be responsible for re-establishing the high ratio of GSH

and GSSG levels, probably by the provision of NADPH for GSH regeneration. Ben-Yoseph et al. (1994) showed that the activity of the pentose phosphate pathway can directly be stimulated by H<sub>2</sub>O<sub>2</sub> in a concentration-dependent fashion. As long as glucose is not the limiting factor, astrocytes should thus be able to cope with GSH depletion by H<sub>2</sub>O<sub>2</sub>. It can be assumed that at least for the lower concentrations of H<sub>2</sub>O<sub>2</sub> in our experiments GSH is probably restored later on.

The AOE system confers protection against oxidative damage. Indeed, overexpression of catalase in immortalized neural cells protected against H<sub>2</sub>O<sub>2</sub> toxicity (Mann et al. 1997). On the other hand overexpression of MnSOD or CuZnSOD in cells of neuronal origin afforded no protection (Mann et al. 1997; Zhong et al. 1996) against oxidative damage caused by radiation or BCNU (1,3-bis(2-chloroethyl)-1-nitrosourea) treatment. Probably overexpression of MnSOD can cause an imbalance of AOE, which results in an elevation of



**Fig. 8A, B** Gene expression of GPx in primary astrocyte cell cultures after treatment with H<sub>2</sub>O<sub>2</sub>. **A** PCR blot of GPx and GAPDH mRNAs obtained by RT-PCR after exposure of 2-week cultures of astrocytes to the indicated concentrations of H<sub>2</sub>O<sub>2</sub> for 24 h. **B** GPx/GAPDH quotient after densitometric scanning of PCR blots obtained after RT-PCR shown in **A**. Values are expressed as percentage of controls (means  $\pm$  SEM from three different experiments). \* $P < 0.05$  vs controls

intracellular H<sub>2</sub>O<sub>2</sub>. If the expression levels of the H<sub>2</sub>O<sub>2</sub> scavenging enzymes, catalase and GPx, are not elevated accordingly oxidative damage will be the consequence. We found basal mRNA expression of catalase, MnSOD, CuZnSOD, and GPx in astrocyte cell cultures. Astrocytes were treated with H<sub>2</sub>O<sub>2</sub> after 2 weeks of culturing, when a stable expression pattern of AOE was observed. H<sub>2</sub>O<sub>2</sub> exposure lead to a 2–3 times increase in catalase and MnSOD mRNA expression in astrocytes. GPx mRNA levels increased about 1.5 times, whereas Cu-ZnSOD mRNA expression was not influenced by H<sub>2</sub>O<sub>2</sub> treatment. Obviously the upregulation of AOE in our system strongly depends on the presence of ROS. The mechanisms residing behind the induction of catalase and MnSOD by oxidants are still under investigation. In the promoter region of the MnSOD gene consensus sequences for NF- $\kappa$ B and AP1 binding sites can be found by computer analysis (Darville et al. 2000; Ho et al. 1991). The transcription factor NF- $\kappa$ B as well as AP1 have been extensively studied as transcriptional activators of genes in stress response (Baeuerle 1991; Curran and Franz 1988). In accordance, the involvement of NF- $\kappa$ B and AP1 in activating MnSOD genes in response to TNF- $\alpha$ , TPA, H<sub>2</sub>O<sub>2</sub>, or thiol-reducing agents has been suggested in human and rat cells (Darville et al. 2000; Kuo et al. 1999; Yoshioka et al. 1994). Recently induc-

tion of the rat MnSOD gene by inflammatory mediators was also found to be regulated through some intronic sequences (Darville et al. 2000; Kuo et al. 1999; Rogers et al. 2000). While MnSOD seems to be regulated on the transcriptional level, for catalase a post-transcriptional regulation can be assumed. Stabilization of catalase mRNA could be shown through a redox-sensitive mRNA binding protein in lungs of neonatal rats exposed to hyperoxia (Clerch and Massaro 1992). A similar mechanism may be important for upregulation of catalase gene expression in astrocytes.

In our experiments also GPx mRNA expression was influenced by H<sub>2</sub>O<sub>2</sub> treatment. In mice with a homozygous null mutation for Gpx1 an increased susceptibility to the oxidative stress inducing agent paraquat and H<sub>2</sub>O<sub>2</sub> could be shown (de Haan et al. 1998). Emphasis was laid on the importance of Gpx for protection of neurons against those oxidative stressors. In the 5' flanking region of the human Gpx gene two cis acting elements were characterized which are of importance for gene regulation in an oxidative environment (Cowan et al. 1993). It can be assumed that the rat gene is regulated similarly.

Our data show, that astrocytes are able to respond to oxidative stress via their AOE system. This upregulation of the AOE system may be crucial for cell survival after oxidative brain damage and may contribute to the important role for astrocytes in the antioxidant defense of the brain.

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