

S100b protects IMR-32 cells against Ab(1-42) induced neurotoxicity via modulation of apoptotic genes expression^{*}

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ABSTRACT

Amyloid beta (1-42) peptide is considered responsible for the formation of senile plaques that accumulate in the brain of patients with Alzheimer's disease (AD). In the past years considerable attention has been focused on identifying new protective substances that prevent or almost retard the appearance of amyloid beta (1-42)-related neurotoxic effects. In this study, human neuroblastoma cells (IMR-32) was used as system model to evaluate the protective role of S100b, a neurotrophic factor and neuronal survival protein, that is highly expressed by reactive astrocytes in close vicinity of beta-amyloid deposits, against amyloid beta (1-42)-dependent toxicity. Our results show that at nanomolar concentrations, S100b protects cells against A β mediated cytotoxicity, as assessed by MTS vitality test. The protective mechanism seems to be related to the effect on bcl-2 (an anti-apoptotic gene) expression, which is highly down-regulated by amyloid beta (1-42) treatment, while resulted more expressed in the presence of S100b. On the contrary, Bax, a pro-apoptotic gene, resulted down-regulated by the treatment with S100 compared with the results obtained in the presence of amyloid beta (1-42) peptide. However, at micromolar doses, S100b is toxic for IMR-32 cells and its toxicity adds to that of the A β peptide, suggesting that additional molecular mechanisms may be involved in the neuro-

toxic process.

Keywords: S100b; Neurodegeneration; Oxidate Methionine; Apoptotic Genes Expression

1. INTRODUCTION

S100b comes from the S100 family of EF-hand Ca²⁺ binding proteins, mainly expressed in astrocytes of the central nervous system, exerting intracellular and extracellular regulatory activities [1,2]. In particular the intracellular roles of S100 proteins are implicated in the regulation of protein phosphorylation, the dynamics of cytoskeleton constituents, Ca²⁺ homeostasis, enzyme activities, transcription factors, cell growth and differentiation, and the inflammatory response [3-5]. The extracellular implications are instead implicated in trophic maintenance, survival and outgrowth of neuronal cells, both during development of the nervous system and in neurodegenerative conditions [4].

In the brain, astrocytes release S100b constitutively but the level of protein is increased by a number of agents such as 5-HT_{1A} receptor agonists, glutamate, adenosine, lysophosphatidic acid [6-8] and by pathological circumstances such as Alzheimer's disease and Down's syndrome, where S100 levels in severely affected brain regions are greater than age-matched controls [9,10].

On this subject, recent hypotheses shows that in Alzheimer's disease, the astrocytes production of pro-inflammatory cytokines such as S100b and IL-1, regulates the early phases of neuronal degeneration [1,11-13]. In fact, although the hypothesis of amyloid and tau aggregation could explain the neuropathological alterations during the late stage of AD, strong evidences demon-

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strate that distress accumulated in cells, pro-inflammatory cytokines mediated, could initiate an early activation of the innate cellular immunity and the inflammatory cascade [14]. In this scenario, the S100b produced by astrocytes, seems to be one of the most important actors, since modulates both the phases.

In fact, in early triggering events of AD, astrocytes produce S100b at nM mount, exerting "*in vitro*" a pro-survival effect on neurons and a stimulation of neurite outgrowth. Instead, during the late phases of neurodegeneration, when levels of S100 increase (mRNA were elevated 10 - 20-fold in extracts of temporal lobe from autopsy samples of Alzheimer's disease patients) it prevails neuronal death via apoptosis [15-17]. Recent results [18-20], evidence that beta amyloide peptide, the major protein component of senile brain plaques in Alzheimer's disease, determines both an oxidative stress and a cascade mechanism apoptotic triggering that depend also on the oxidative state of Met-35. We investigated in the present study, the relationships between the S100b and A β (1-42) peptide. In particular we have studied, in human neuroblastoma (IMR-32) cells, the effect of low (nanomolar) and higher (micromolar) S100b concentration previous treated with A β peptides carrying Methionine 35 both in reduced and oxidative state, on cellular vitality and on the expression of major apoptotic gene expression. The obtained results indicate that an important defensive mechanism exists to preserve the A β (1-42) induced neuronal degeneration carried out by S100 protein at nanomolar concentration and this effect is more evidenced with methionine in reduced state. On the contrary, at micromolar doses S100 adds its toxicity to that induced by the A β peptide, suggesting that additional molecular mechanisms may be involved in the neuropathology of Alzheimer Disease.

2. MATERIALS AND METHODS

2.1. Cell Culture and Treatments

Human neuroblastoma IMR-32 cells were grown in minimum essential medium (Biochrom KG, Berlin, Germany) supplemented with 10% heat inactivated fetal bovine serum (HyClone, Logan, UT), 100 IU/ml penicillin, and 100 μ g/ml streptomycin (Invitrogen, Grand Island, NY, USA), and cultured at 37°C in an atmosphere of 5% CO₂ in air. Cell differentiation was induced by 1 mM dibutyryl cAMP and 2.5 μ M 5-bromodeoxyuridine (Sigma, St. Louis, MO, USA), which were added to the culture medium three times per week, starting from the day after plating. After a week, the differentiated cells were plated at an appropriate density according to each experimental procedure.

A β peptides were obtained by Peptide Speciality Laboratories GmbH (Heidelberg, Germany). Analysis of the

peptide by reverse phase high performance chromatography (HPLC) and mass spectrometry revealed a purity >98%. Stock solution of A β peptide, 2.5 mM in DMSO were prepared according to the manufacturer's instructions and stored at -80°C (in previous studies [21,22] these conditions have been shown to lead to the predominance of the soluble monomeric form of this peptide). Human recombinant S100b was purchased by Sino Biological Inc; stock solutions of S100b was solved in minimum essential medium, filtered and stored at -20°C. S100b was added to the cultures 12 hr before exposure to A β peptide and to conform control experiments, DMSO concentration (<0.5%) was the same in all solutions used.

2.2. Cell Availability Measurement

For vitality determination, IMR-32 cells were plated in 96-well plates at a density of 10,000 cells/well and incubated for 24 and 48 h with and without (as control) 10 μ M A β and with different concentration of S100b protein. Cell survival was evaluated by the 3-[(4,5-dimethylthiazol-2-yl)-5,3-carboxymethoxyphenyl]-2-(4-sulfophenyl)-2H tetrazolium, inner salt (MTS) reduction assay (CellTiter 96 Aqueous One Solution Cell Proliferation Assay, Promega, Madison, WI, USA). The MTS assay is a sensitive measurement of normal cellular metabolic status, particularly that in mitochondria, which reflects early cellular redox changes [23]. After exposure to the different treatments, cells were incubated with MTS solution (2 mg/ml) for 4 h at 37°C in a 5% CO₂ incubator. The intracellular soluble formazan produced by cellular reduction of the MTS was determined by recording the absorbance of each 96-well plate using the automatic microplate photometer (SpectraCount, Packard Bioscience Company, Groningen, The Netherlands) at a wavelength of 490 nm. The morphological features of apoptotic degeneration observable afterwards 48 h of incubation with the different treatments (10 μ M amyloid and plus S100b) were analyzed and photographed by phase-contrast microscopy (Nikon Eclipse TS100) 40 \times .

2.3. Reverse Transcription and Polymerase Chain Reaction

Total RNA was isolated using SV total RNA isolation System (Promega) that includes the elimination of any genomic DNA by DNase treatment. The purity and quantity of the resulting RNA were determined via the measurement of the absorbance at 280 and 260 nm respectively. The A₂₆₀/A₂₈₀ ratio was about 1.8. The RNA was concentrated by precipitation and re-dissolved in water RNase free. Total RNA (1 μ g) from each sample was used for first strand cDNA synthesis using the M-MLV Reverse and Oligo-dT, as random primer (SIGMA).

The PCR was performed with 200 ng of cDNA using Gene-spin Taq-Polymerase with 2.0 mM MgCl₂, 0.4 μM primer, 0.4 μM dNTPs and 2 units polymerase for each reaction (25 μl) under primer specific conditions.

The primers used for amplification were:

GAPDH sense primer;

5'-AAGAAGATGCGGCTGACTGTCCGAGCCACAT-3',

GAPDH anti-sense primer;

5'-TCTCATGGTTCACACCCATGACGAACATG-3',

Bcl₂ sense primer;

5'-CGACGACTTCTCCCGCCGCTACCGC-3',

Bcl₂ anti-sense primer;

5'-CCGCATGCTGGGGCCGTACAGTTCC-3',

Bax sense primer;

5'-ACCAAGAAGCTGAGCGAGTGTC-3',

Bax anti-sense primer;

5'-ACAAAGATGGTACGGTCTGCC-3',

Caspase3 sense primer;

5'-AGAAGATCACAGCAAAGGAGC-3',

Caspase3 anti-sense primer;

5'-TCAAGCTTGTCCGTCATACTG-3'

The PCR-products had the following sizes: GAPDH 457 bp, Bcl₂ 319 bp, Bax 367 bp, Caspase3 378 bp. The following experimental protocol for PCR reaction (35 cycles, 45 cycles for HTert) was used: denaturation: 3 min; 95°C; amplification: denaturation: 30 s, 95°C; annealing: 20 s, at 60°C (GAPDH), 67°C (Bcl₂), 56°C (Bax) and 66°C (Caspase-3) respectively; elongation: 30 s, 72°C, final elongation: 72°C, 90 s. PCR products were analyzed by electrophoresis in agarose 1.8% with ethidium bromide (1 μg/ml) in TBE 1× buffer (Tris 40 mM, EDTA 1 mM, boric acid 44 mM) for 2 h at 80 V (constant voltage) with 123 bp ladder as molecular weight markers.

Images of gels were acquired (Bio-Rad Gel Doc 2000, Hercules, CA, USA) and scanned (Bio-Rad GS800, Hercules, CA, USA) using Bio-Rad Quantity One software. The density of the PCR bands were divided by that of the housekeeping gene and expressed as percent of the control band density.

2.4. Measurement of Caspase-3 Activity

Caspase-3 activity was measured by using a specific assay kit from Sigma Chemical Co. (St. Louis, MO, USA) following manufacture's instructions. DEVD-pNA was used as a colorimetric substrate. IMR32 cells were plated at a density of 2.9×10^6 cells/35 mm dish: after treatment with Aβ peptides and S100b, cells were harvested by centrifugation. The pellets were washed with PBS, lysed in 50 ml of chilled cell lysis buffer and left on ice for 10 min. Lysate was centrifuged at $10,000 \times g$ for 1 min at 4°C, and supernatant was used for the caspase-3 assay. The protein concentration was confirmed by the BCA assay. The protease activity was determined by the spectrophotometric detection at 405 nm of the chromophore

p-nitroanilide (pNA) after its cleavage by caspase-3 from the labeled caspase-3-specific substrate (DEVD-pNA). Additional control assays in the presence of specific caspase-3 inhibitor (DEVD-CHO) and in the absence of recombinant human caspase were performed for measuring the non-specific hydrolysis of the substrate (data not shown).

2.5. Statistical Analysis

The data were analysed using one-way ANOVA, followed by *post hoc* Newman-Keul test for multiple comparisons among group means, using a Prism TM computer program (Graph-Pad, San Diego, CA, USA), and differences were considered statistically significant if $P < 0.05$. All results are presented as the mean ± S.E.M. of at least three different experiments performed in triplicate, unless otherwise specified

3. RESULTS

The first point investigated in this study was to clarify the experimental conditions to evidence the exerted role by S100b on Aβ (with methionine both in reduced and oxidative state) induced cellular toxicity. Thus, the toxic effects of Aβ and Aβ-Met-35^{OX} in the presence and in the absence of different concentrations of S100b protein were evaluated on IMR-32 cells (a cellular line of human neuroblastoma).

Figure 1 shows the survival of IMR-32 cells (measured by the reduction of MTS activity), after the exposure to Aβ peptides with 5nM (panel (a)) and 5 μM (panel (b)) S100b. It appears evident that after 48 h of incubation, Aβ and Aβ-Met-35^{OX} peptides have a toxic effect on availability of IMR32 cells inducing a significant cellular death with respect to the control. It should also be noted that the major extent of cellular death is induced by Aβ where methionine-35 is in the reduced form according to our previous data [18]. About the S100b role, it is evident that the protein alone does not interfere at nanomolar dose with the IMR32 vitality but exerts an important protective effect on the cells treated with Aβ peptides, showing availability values very similar to the untreated cells. At micromolar concentration instead S100b is neurotoxic and its effect potentiate the toxic effect of beta peptides. Hence, for studying the molecular interaction between S100b and Aβ we used a concentration of 5 nM and 10 μM respectively on all experimental procedures [24].

In phase-contrast observation (**Figure 2**), control cells (panel (a)) and cells treated with 5 nM S100b (panel (b)) appear generally healthy, with round cell bodies and a well developed network of neurites throughout the experiments. Aβ treatment was performed in the presence of 10 μM Aβ peptides, both with reduced (panel (c)) and

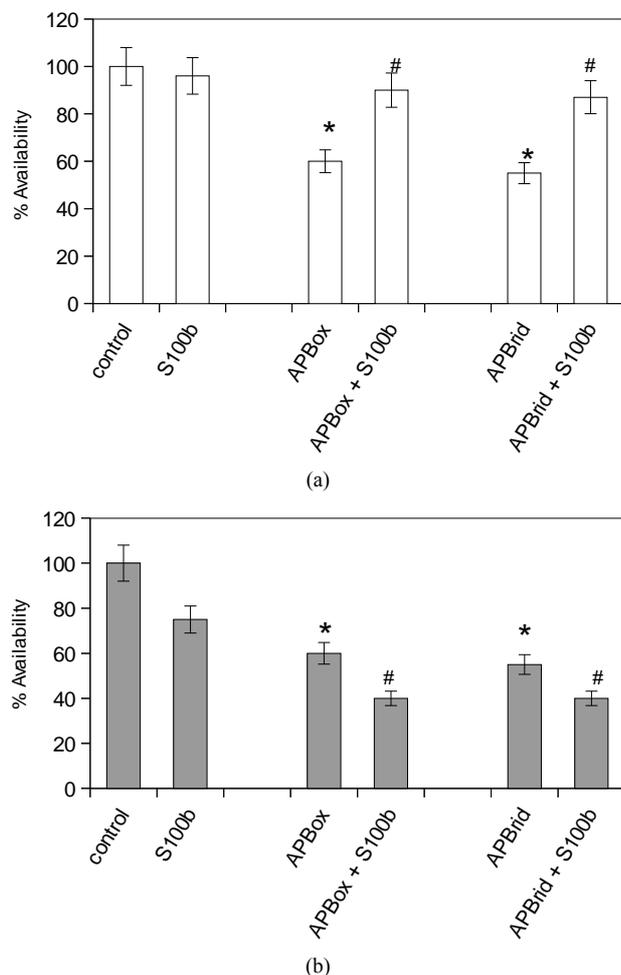


Figure 1. Panel (a): Effects of S100b 5 nM (Panel (a)) and 5 μ M (Panel (b)) in the absence and in the presence of 10 μ M A β ² (with methionine in reduced and oxidative state) on IMR-32 cells after 48 h of incubation. The cell survival is expressed as percent of cells untreated. Cells (10,000 cells/well) were cultured with substances under analysis (experimental conditions are reported materials and methods), and the availability of cells was measured by MTS assay. All values indicate means \pm S.E. of seven independent experiments. Significantly different from cells untreated: *P < 0.01. Significantly different from cells treated with beta amyloid peptides: #P < 0.01.

with oxidative 35-Methionine (panel (d)). After 48-h of treatment, a significant number reduction of living cells is evident and the cell bodies of a large amount of neurons were diminished, with condensed nuclei and retracted processes especially with the treatment with oxidative peptide. On the contrary, cells incubated with A β peptides, in the presence of S100b (panel (e) and (f)) were very similar to the control.

As was demonstrated in previous work S100b promotes cellular survival by increasing the expression of the anti-apoptotic factor [25,26], we analyzed the gene expression of pro-apoptotic gene Bax and anti-apoptotic genes Bcl-2 in untreated neuroblastoma cells (as control) and

exposed for 48 h to 10 μ M with both A β reduced and oxidative and with 5 nM S100b (Figure 3). It is evident that A β treatment is able to bring about the synthesis of mRNA for pro-apoptotic gene Bax particularly when the peptide has methionine-35 in reduced state. The simultaneous treatment with beta amyloid peptides and S100b determines a strong decreasing of Bax expression, according to the cellular availability but mostly an over-expression of anti apoptotic gene Bcl2. In fact, while the RNA for Bcl2 is blank for cells treated with beta amyloid peptides such as in the controls, the presence of S100b stimulates significantly the expression of this gene demonstrating an ability to modulate both Bax and Bcl2 genes.

Finally, as Bcl-2 has been shown to prevent caspase activation and subsequently cell death also the expression of caspase 3 gene and its enzymatic activity were investigated. As reported in Figure 4, the treatment with A β peptides increased caspase-3 expression and activity in IMR 32 cells and these outcomes are strongly prevented by the protective effect of S100b.

4. DISCUSSION

The molecular mechanisms of Alzheimer's disease (AD), one of the most common causes of dementia, is as yet not completely clear. AD has been identified as a protein misfolding disease due to the accumulation of abnormally folded amyloid beta protein, a peptide of 1-42 aminoacid, in the brain of Alzheimer's patients [27,28]. Amyloid beta is an abnormal proteolytic byproduct of the transmembrane protein amyloid precursor protein (APP), whose function is unclear but thought to be involved in neuronal development. Amyloid beta peptide is a monomeric soluble molecule which contains short regions of beta sheet and polyproline II helix secondary structures in solution [29], though they are largely alpha helical in membranes [30]; however, at sufficiently high concentration, they undergo a dramatic conformational change to form a beta sheet-rich tertiary structure which aggregates to form amyloid fibrils [31]. These fibrils deposit external neurons in dense formations known as senile plaques or neuritic plaques, in less dense aggregates as diffuse plaques, and sometimes in the walls of small blood vessels in the brain.

Another important pathogenetic event in AD is a relevant oxidative stress: the "A β cascade hypothesis" of Butterfield [32] in fact suggests that A β as small oligomers can insert into the lipid bilayer and initiate lipid peroxidation and oxidative damage to proteins and other molecules. A consequence of this event is the influx of Ca²⁺ into the neuron, resulting in the loss of ion homeostasis, mitochondrial dysfunction, synapse loss and finally cell death [32-35].

Methionine-35 (Met-35) side chain of A β P appears to play a critical role in peptide's neurotoxicity; indeed, this

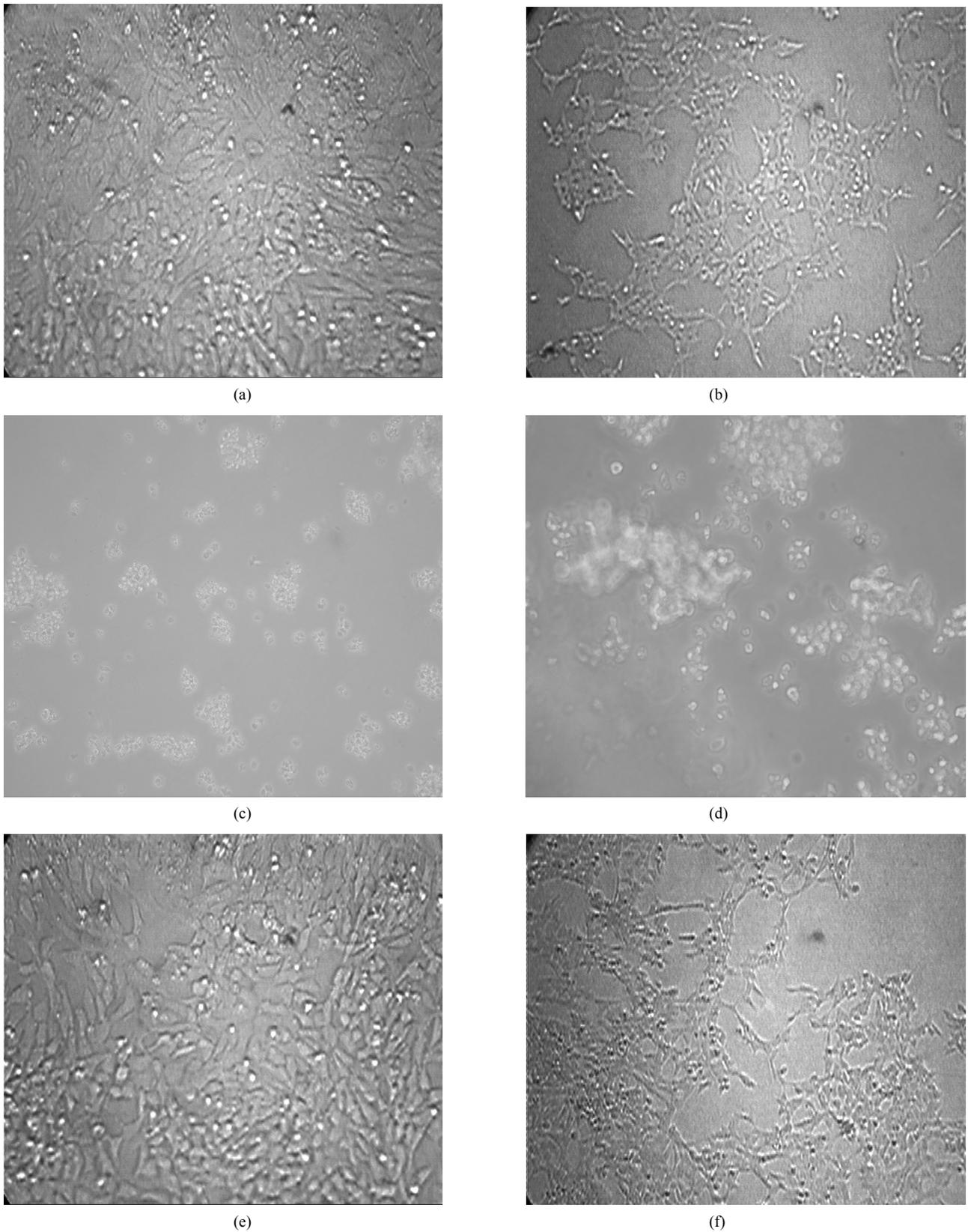


Figure 2. Phase-contrast micrography shows IMR-32 cells in medium (panel (a)); treated with 5 nM S100b (panel (b)); with 10 μM Aβ with reduced (panel (c)) and oxidative 35-Methionine (panel (d)) and cells incubated simultaneously with S100b in presence of Aβ (panel (e)) and Aβ-Met-35^{OX} (panel (f)). All treatments were performed for 48 h.

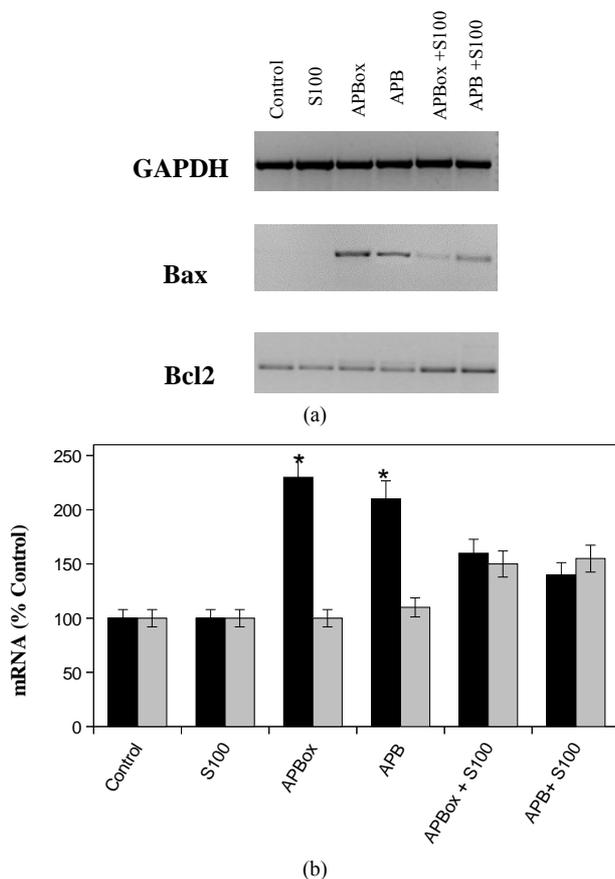


Figure 3. Panel (a): Expression of Bax and Bcl-2 genes in IMR-32 cells. Cells were incubated for 48 h with medium (control), with 5 nM S100, with 10 μ M A β with Methionine 35 oxidated (APBox) and reduced (APB) and with the simultaneous presence of S100 and A β peptides. GAPDH gene expression was used as housekeeping. Panel (b): The intensities of the bands were quantized by densitometric scanning of agarose gel bands and expressed as percent of Bax (black bars) and Bcl-2 (grey bars) mRNA expression respect to the control. Results are from eight independent experiments. Significantly different from controls *P < 0.01.

residue is mostly susceptible to oxidation in vivo [20, 36,37], and A β bearing oxidized Met-35 is found in large quantities in post-mortem AD plaques [38,39], while oxidative stress does not occur in the brain in vivo if the Methionine 35 residue of A β (1-42) is not present [40, 41].

In this complex pathogenetic network, numerous exogenous and endogenous substances are been tested to individuate possible molecules able to oppose the neurodegenerative mechanisms generated by A β (1-42) [42-44]. In this context the observation that during the early stages of AD, microglia produces numerous proinflammatory molecules such as S100b which expression is highly modulated by reactive astrocytes in close vicinity of beta-amyloid deposits, induced us to investigate the interplay between A β , both with Met-35 oxidate and

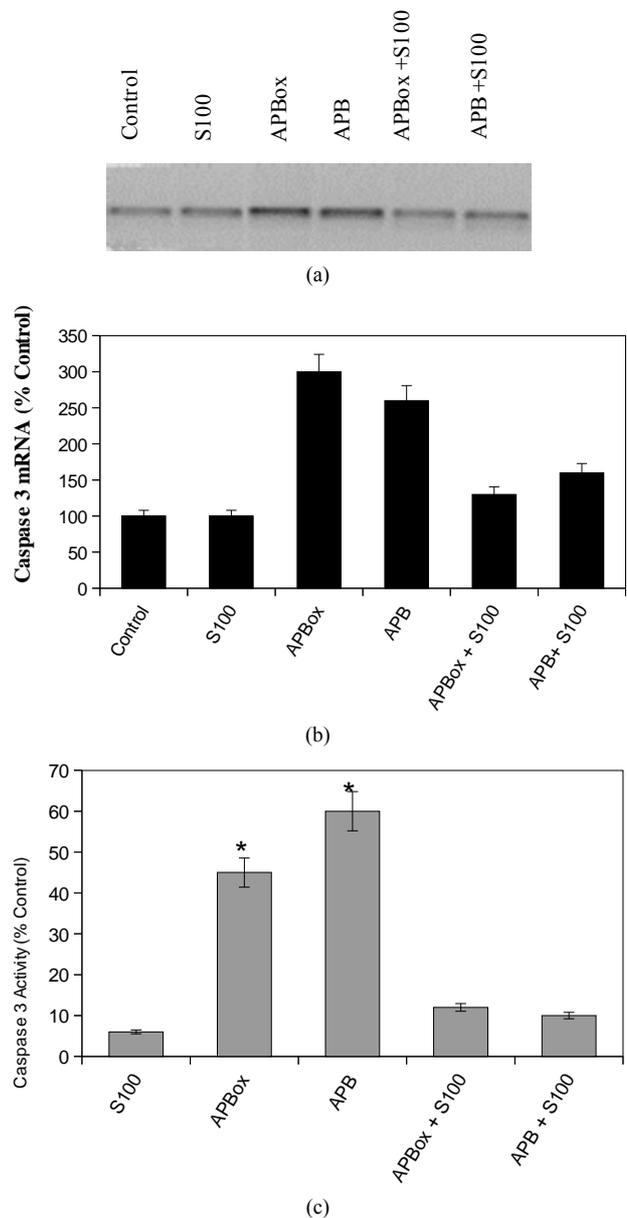


Figure 4. Panel (a): Expression of caspase 3 gene in IMR-32 cells. Cells were incubated for 48 h with medium (control), with 5 nM S100b, with 10 μ M A β with Methionine 35 oxidated (APBox) and reduced (APB) and with the simultaneous presence of S100b and A β peptides. GAPDH gene expression was used as housekeeping (data not shown). Panel (b): The intensities of the bands were quantized by densitometric scanning of agarose gel bands and expressed as percent of caspase 3 mRNA expression respect to the control. Panel (c) Caspase-3 activity (expressed as percent of control) in IMR-32 cells incubated for 48 h with medium (control), with 5 nM S100b, with 10 μ M A β with Methionine 35 oxidated (APBox) and reduced (APB) and with the simultaneous presence of S100b and A β peptides. Results are from eight independent experiments. Significantly different from controls *P < 0.01.

reduced, and S100b. In particular the S100b is carefully modulated during AD evolution, coming by nanomolar

values, during the first phases of pathology, to micromolar concentration [11-13]. In this study we examined the possible participation of this protein, at different concentrations, to contrast the AD progression.

In this regard, it has been evidenced that S100b at nanomolar concentrations is able to protect human neuroblastoma cells (IMR-32) against $A\beta$ -mediated toxicity. Since $A\beta$ -mediated mechanism of toxicity seems to be mediated by an apoptotic pathway [45], we investigated the expression of Bax, Bcl-2 and caspase 3 genes, three of the main important apoptosis-related genes [46,47], in $A\beta$ -treated cells both with oxidated and reduced methionine 35 with and without S100b.

Bax is a gene encoding for a protein which accelerates programmed cell death by its translocation to the mitochondrion membrane, leading to the release of cytochrome *c*. When translocated to the mitochondrial membrane, Bax can in fact homodimerize and triggers the activation of terminal caspases by alteration of mitochondrial functions, which results in the release of apoptosis-promoting factors into the cytoplasm [46].

Bcl-2 product is instead a protein localized in the cytoplasmic face of the mitochondrial outer membrane, endoplasmic reticulum, and nuclear envelope [45,46]. Bcl-2 has been shown to prevent cytochrome *c* release, caspase activation, and cell death.

The results of present study evidenced that $A\beta$ treatments resulted in increased Bax expression in IMR-32 cells: but when $A\beta$ -treated cells were incubated in the presence of S100b, Bax expression results up-regulated. On the other hand, our results show that the expression of the anti-apoptotic gene Bcl-2 is very low in control and in IMR-32 cells treated with $A\beta$ peptides while treatment with S100b considerably increases the expression of this gene. This finding indicates that a significant shift of Bax/Bcl-2 *ratio* in favour of Bcl-2 occurs in $A\beta$ /S100b-treated cells with respect to that showed by $A\beta$ -treated cells. Another important apoptotic event is characterized by the caspase 3 activation. In particular, mRNA accumulation of Bax may be sufficient to initiate the intrinsic pathway of caspase activation and therefore to lead to cell death [47,48]. Indeed, Bax can form homodimers/heterodimers with other members of the Bcl-2 family upon translocation to the mitochondrial membrane, and activate a cascade of events including increased mitochondrial membrane permeability, release of cytochrome *c* into the cytoplasm and activation of caspases. Our results, evidenced that the treatment of IMR-32 with $A\beta$ peptides increases the expression of caspase 3 and its enzymatic activity: also in this case, the simultaneous presence of S100b determines a protective effect, inhibiting the production and activity of this pro-apoptotic enzyme. As expected and described previously in literature [18], the reduced peptide was far more effective than

the oxidized form in determining the apoptotic cascade but the protective effect of S100b is equally effective demonstrating a common interplay with the $A\beta$ peptides, independently by the oxidative state of methionine 35. To summarize, our results indicate that $A\beta$ -mediated degeneration in IMR-32 cells is due to a large shift of Bax/Bcl-2 ratio in favour of the pro-apoptotic Bax with subsequent cell death. The presence of nanomolar concentrations of S100b significantly lowers the toxicity induced by $A\beta$ peptides through the reduction of the pro-apoptotic Bax/Bax homodimers and the increase of antiapoptotic Bcl-2/Bcl-2 homodimers. Moreover Bcl-2 has been shown to prevent caspase activation and cell death; therefore, the effect of S100b on caspase 3 is mediated both by a direct effect on RNAm expression and indirectly through enzyme activity post-transcriptionally regulated by Bcl-2/Bcl-2 homodimers.

Another important question is the increase of S100b observed in late phases of Alzheimer: the levels of S100 beta protein and mRNA, were elevated 10 - 20-fold in extracts of temporal lobe from autopsy samples of Alzheimer's disease patients compared to those of aged control patients. Our data evidenced that the treatment with S100b at micromolar concentration on IMR-32 results strongly toxic potentiating the neurotoxicity by beta amyloide.

The synergical toxic effect exerted by S100 and amyloide peptide could be discussed on the basis of RAGE (advanced glycation end-product receptor) expression. RAGE is an inducible receptor, which exists on neurons and microglial cells, responsible of extracellular elimination of neurotoxic advanced glycation end-products (AGEs) [49]. Indeed, RAGE is multi-ligand receptors: able to accept more than one ligand concomitantly including $A\beta$ peptides, amphotericins, and also S100b [49-51]. Hence, S100b concentration, modulating also the expression of RAGE, could underlie the progression of AD and of a range of other chronic disorders [25,52-53].

5. CONCLUSION

In conclusion, our results evidenced that during early AD pathogenesis, a network between astrocytes and neurons could govern the $A\beta$ toxicity through apoptotic signal modulation. Instead, in advanced phases of pathology, the (over)-expression of S100b is so high to accelerate AD-like pathology. In this light the biosynthesis modulation of pro-inflammatory cytokines such as S100b, may be a promising therapeutic strategy to delay AD progression.

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