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1 TSPO over-expression increases motility, transmigration

2 and proliferation properties of C6 rat glioma cells

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21	Proliferation; Ligand-binding; over-expression
22	

1 SUMMARY

2 Gliomas are one of the most malignant cancers. The molecular bases regulating the 3 onset of such tumors are still poorly understood. The translocator protein (TSPO), formerly known as the peripheral-type benzodiazepine receptor, is a mitochondrial 4 5 permeability transition (MPT)-pore protein robustly expressed in gliomas and involved in the regulation of apoptosis and cell proliferation. TSPO expression 6 levels have been correlated with tumor malignancy. Here we describe the production 7 8 of C6 rat glioma cells engineered to over-express the TSPO protein with the aim of 9 providing the first direct evidence of a correlation between TSPO expression level 10 and glioma cell aggressiveness. We observed that TSPO potentiates proliferation, 11 motility and transmigration capabilities as well as the ability to overcome contactinduced cell growth inhibition of glioma cells. On the whole, these data demonstrate 12 13 that TSPO density influences metastatic potential of glioma cells. Since several data 14 suggest that TSPO ligands may act as chemotherapeutic agents, in this paper we also 15 demonstrate that TSPO ligand-induced cell death is dependent on TSPO density. 16 These findings suggest that the use of TSPO ligands as chemotherapeutic agents 17 could be effective on aggressive tumor cells with a high TSPO expression level.

18

1 INTRODUCTION

2 The translocator protein (18 KDa) (TSPO), formerly known as the peripheral-type 3 benzodiazepine receptor (PBR) [1], is an abundant mitochondrial protein expressed 4 in peripheral tissues, as well as in astrocytes and microglial cells within the central 5 nervous system (CNS) [2] and [3]. Increased TSPO expression has been observed in cancer, including brain tumors [4], [5], [6], [7] and [8]. TSPO has been implicated in 6 many physio-pathological conditions including apoptosis, microglial activation 7 8 related to brain damage, steroidogenesis, and cell proliferation [9]. In mitochondria, 9 TSPO is associated with other proteins, including the voltage-dependent anion 10 channel (VDAC) and the adenine nucleotide carrier (ANT), making up the 11 mitochondrial permeability transition (MPT)-pore [10] whose modulation has been suggested as a critical event in the regulation of processes underlying cellular 12 13 survival/death [2], [10], [11] and [12]. It has been demonstrated that specific TSPO 14 ligands show pro-apoptotic activity and may act, through the modulation of the 15 MTP-pore opening, as potential drugs useful in cancer therapy [5], [13], [14], [15], 16 [16] and [17]. Among tumors, gliomas represent an elective target for TSPO ligand 17 mediated chemotherapy; this protein being virtually absent in the neurons and over-18 expressed in glioma with respect to normal glial cells [3], [18], [19], [20], [21] and 19 [22].

It is interesting to note that TSPO expression level positively correlates with cell malignancy in several tumors including gliomas [18], [23], [24] [25], [26], [27] and [28]. However, the role of TSPO up-regulation in brain tumor cells is still puzzling.

For these reasons over the last years we have been focussing our attention on the study of TSPO and TSPO ligand effects in glioma aggressivness. Up to now no studies have been performed by direct comparison of brain tumor cells engineered to over-express TSPO with wild type cells of the same cell line. Here, we describe the production of glioma cells engineered to over-express the TSPO and examine the relationship between TSPO up-regulation and growth, motility, transmigration and susceptibility to the TSPO ligand PK11195-induced apoptosis of glioma cells.

A CERCENT

1 MATERIALS AND METHODS

2 CELL CULTURES

Rat C6 glioma cells (a kindly gift from Professor Damir Janigro, Cleveland Clinic
Foundation, Cleveland, OH) were cultured as described in [15].

5 RNA ISOLATION, REVERSE TRANSCRIPTION-PCR, AND CLONING OF

6 **RAT** *TSPO*

Total RNA was extracted from 5×10^6 cells using the Nucleospin RNA II kit 7 (Macheray-Nagel) according to manufacturer's instructions. For RT-PCR, about 1µg 8 9 DNase-digested total RNA was reverse-transcribed using Superscript II reverse 10 transcriptase (Invitrogen) following the recommendations of the manufacturer. Full-11 length cDNA of TSPO was amplified using TSPO specific forward primer (5'CCCAAGCTTATGTCTCAATCCTGGGTACC3') containing a restriction site 12 13 for the endonuclease Hind III at the 5' end and a TSPO specific reverse primer (5'GCGAGGATCCAAAGCAGGATGACCTCAACC3') containing a restriction 14 15 site for the endonuclease BamH I at the 5' end. The condition for PCR were as 16 follows: 2 min at 95°C, 30 cycles of 30 sec at 95°C, 1 min at 55°C, and 30 sec at 17 72°C. A final extension was carried on at 72°C for 7 min. The amplified fragment 18 was Hind III and BamH I digested and then cloned into the vector pFLAG-cmv-2 19 (Sigma) in frame with a short flag epitope in the amminoterminal position. After transformation in competent E. coli cells, some clones were isolated and plasmid 20 21 DNAs were sequenced by automated fluorescent cycle sequencing (ABI). One of 22 these clones (TSPO-pFLAG-cmv-2) was used in the following applications.

23 TRANSFECTION

For transfection, 2µg of TSPO-pFLAG-cmv-2 or pFLAG-cmv-2 (control) were mixed with 10 µL Lipofectamine 2000 reagent (Invitrogen) in 0,5 mL of Optimem (Gibco). 90% confluent C6 glioma cells were transfected with the mixture and incubated for 6 hours according to manufacturer's instructions. Then the transfection medium was replaced with complete medium and the cells were used for experiments.

7 WESTERN BLOT ANALYSIS

Whole cell extract of TSPO-pFLAG-cmv-2 or pFLAG-cmv-2 transfected C6 cells 8 were used for Western blot analysis. 24, 48, 72 and 96 hours after transfection C6 9 10 cells were detached with 2mM EDTA, collected by centrifugation (1000g for 5 min) 11 and lysed by adding 80 µL of lysis buffer (1% Triton X100, 10% glycerol, 20mM Tris HCl pH 7.5, 150 mM NaCl, 10 mM EDTA, 1 mM phenylmethylsulphonyl 12 13 fluoride (PMSF), 0.5 µM aprotinin, 0.5 µM leupeptin) for 30 min on ice. Then samples were frozen-thawed for three times and centrifuged to collect the 14 15 supernatants. Protein concentration was measured by using the Micro BCA protein 16 assay kit (Pierce). 25µg of proteins were separated on a SDS-PAGE gel and 17 electrotransferred onto a nitrocellulose membrane. After blocking with 5% non-fat 18 dry milk (Biorad) in TPBS (50 nM Tris, 150 nM NaCl, 0.05% Tween 20) for 4 19 hours at room temperature, the FLAG-TSPO recombinant protein was detected by 20 adding mouse anti-FLAG antibody (Sigma, 1:1000) in 3% non-fat dry milk in TPBS 21 for 1 hours at room temperature. Filters were then washed in TPBS and incubated 22 for 1 hour at room temperature with 1:100000 dilution of POD-conjugated anti-23 mouse IgG antibody (Biorad). After several washes in TPBS, signal cross-

hybridization was revealed by chemoluminescence using the Supersignal West Dura
substrate (Pierce). To test that an equal amount of total protein were loaded in each
line, after FLAG-TSPO revelation, filters were stripped and re-probed with mouse
anti-ERK antibody (Santa cruz, 1:1000) and revealed as described above.

5 RADIOLIGAND BINDING ASSAYS

Membranes from transfected C6 glioma cells with TSPO-pFLAG-cmv-2 or pFLAG-6 7 cmv-2 (collected 24 hours after transfection) were prepared as previously described 8 by Gorman and Veenman [28] and [29] with minor modifications. Briefly, confluent 9 cells were washed with phosphate-buffered saline, harvested by scraping and 10 collected by centrifugation at 1000 x g for 10 min at 4°C. Cell pellet was suspended 11 in 5 mM Tris-HCl buffer, pH 7.4 containing 160 µg/ml benzamidine, 200 µg/ml 12 bacitracin and 20 µg/ml trypsin inhibitor, homogenized using Ultraturrax 13 homogenizer and centrifuged at 48000xg for 15 min at 4°C. Pellet was suspended in 14 50 mM Tris-HCl buffer, pH 7.4 (assay buffer), homogenized and centrifuged at 15 48000 x g for 15 min at 4°C. Protein content was estimated by the Bradford method 16 [30] using the Bio-Rad Protein Assay reagent, according to the manufacturer's 17 protocol, with bovine serum albumin (BSA) as standard.

18 [3 H]PK 11195 (specific activity 69.9 Ci/mmol) binding assays were conducted 19 incubating cell membranes (40 µg of protein), with 0.5-26 nM [3 H]PK 11195 in a 20 final volume of 500 µl of 50 mM Tris/HCl, pH 7.4 buffer. Non-specific binding was 21 determined in the presence of 1 µM unlabelled PK11195. Binding equilibrium was 22 reached following a 90 min incubation at 0°C. Samples were filtered rapidly under

1 vacuum through GF/C glass fiber filters, washed four times with 3 ml of assay 2 buffer. Finally, filters were placed in vials containing 4 ml of scintillation liquid and bound radioactivity was measured by a scintillation counter (Tri-Carb 2800 TR, 3 Perkin Elmer). 4

Scatchard analyses of saturation binding data, graphic presentation and statistical 5 analyses were performed using the non-linear multipurpose curve-fitting Graph-Pad 6 Prism computer program (Graph Pad Software, version 4.0; San Diego, CA). In 7 detail, for saturation binding experiments one-site binding curve fitting was used. 8

IMMUNOFLUORESCENCE MICROSCOPY 9

Approximately 2x10⁴ TSPO-pFLAG-cmv-2 or pFLAG-cmv-2 transfected C6 cells 10 11 were plated onto poly-D-lysine slides and incubated overnight. Cells were then fixed in paraformaldehyde (4%) for 30 min at room temperature, washed in PBS and 12 13 blocked in blocking solution (0.1% Triton X-100, 0,5% bovine serum albumin in 14 PBS) for 30 minutes. Then, cells were incubated for 1 hour at room temperature in 15 blocking solution containing rabbit anti-FLAG antibody (Sigma, 1:100) and mouse 16 anti-cytochrome c antibody (Santa Cruz, 1:50). After washing in PBST (0.1% Triton 17 in PBS), cells were incubated for 1 hour at room temperature in blocking solution 18 containing FITC-conjugate anti-rabbit secondary antibody, (1:200, Invitrogen) and 19 rhodamine-conjugate anti-mouse secondary antibody (1:200, Santa Cruz). Slides 20 were then mounted and analyzed under the Axioplan fluorescent microscope (Zeiss).

21

CELL PROLIFERATION ASSAYS

22 To determine the proliferation rate, the same number (4000 cells for well) of TSPOpFLAG-cmv-2 or pFLAG-cmv-2 transfected C6 cells, were seeded in 24-well plates, 23

1 immediately after transfection, and allowed to proliferate for 24, 48, 96 h. Both 2 attached and floating cells were taken for counting with a hemocytometer. We designed this experiment in way to maintain cells in the exponential phase of growth 3 (not confluent) up to about 120 hours. For each cell line the experiment was 4 5 performed on three independent transfections. The growth curves were confirmed repeating the entire experiment for three times. The number of TSPO-pFLAG-cmv-2 6 or pFLAG-cmv-2 transfected C6 cells were also evaluated after they had been 7 confluent for three days. For this last analysis, 90% confluent C6 glioma cells were 8 9 transfected with TSPO-pFLAG-cmv-2 or pFLAG-cmv-2 vectors as described above. 10 Three days after transfection the medium covering cells was collected, the adherent 11 cells trypsinized, added to the collected medium and counted with a hemocytometer. For each cell line the experiment was performed on ten independent transfections 12 13 To calculate the mitotic index of C6 cells overexpressing TSPO, 6-well plates containing 90% confluent C6 glioma cells were transfected with TSPO-pFLAG-14 15 cmv-2, pFLAG-cmv-2, or pEGFP-C1 (a vector that drives the expression of the 16 Green Fluorescent Protein in the cytoplasm). Three independent transection were 17 performed for each vector. Six hours after transfection, cells of each well were 18 detached and split in three 6-well plates (to reach 30% confluence) and allowed to 19 proliferate for 24 and 72 hours.

20 Cells were then detached, collected by centrifugation and resuspended in 40 μ l of 21 glycerol, acetic acid, PBS (1:1:13) solution containing 1x Hoechst. Two 5 μ l 22 aliquots of cell suspension for each sample were spotted on a glass slide and allowed 23 to dry. The number of mitotic figures was counted under the fluorescence

microscope Axiovert 25 (Zeiss). Two 10 µl aliquots for each sample were used to
count the number of total cells with a hemocytometer. For each cell line the mitotic
index [(mitotic figures/ total cells) x1000] was calculated in 9 replicate wells. Two
independent experiments were performed.

5 MOTILITY ASSAY

Equal amount of C6 cells were plated in 6-well plates (95% confluent), transfected 6 with TSPO-pFLAG-cmv-2, pFLAG-cmv-2 or pEGFP-C1 and a scratch (to remove 7 cells) was done along the diameter of all 6-well plates 6 hours after transfection. 8 9 Photographs were taken immediately after scratching (T0) in three different points along the scratch. 20, 25, 40 hours after the scratch the same points were 10 photographed at the same magnification to determine the colonization of the scratch. 11 The scratch of each picture was measured by using the Adobe Photoshop ruler. The 12 13 migration rate (cm/h) was calculated for each treatment considering the mean 14 difference between the scratch width at T0 and at each time of analysis in the three 15 photographed points. For each cell line the experiment was performed on six 16 independent transfections.

17 TRANSMIGRATION ASSAY

18 Cell ability to transmigrate was assayed in modified Boyden chambers (Neuro 19 Probe). 10% FBS in C6 growth medium supplemented with 0.1% BSA was placed 20 in the lower chamber as source of chemoattractant. Polyvinylpyrrolidone-free 8 µm-21 pore polycarbonate filters were coated with 5 mg/L porcine skin gelatin (Sigma) and 22 placed between upper and lower chambers. TSPO-pFLAG-cmv-2 or pFLAG-cmv-2 23 transfected C6 cells were trypsinized, resuspended in growth medium supplemented

with 0.1% BSA and then placed in the upper chamber at a density of 2×10⁵ cells for chamber. After 5 hours of incubation at 37°C in a humidified atmosphere, filters were fixed in absolute ethanol for 30s and stained with 2% toluidine blue (Sigma) for 10min at room temperature. Cells incapable of migrating were gently scraped from the upper side of the filter, while migrated cells were counted. Fifteen microscope fields (10X) for each filter were counted.

7 CELL VIABILITY ANALYSIS BY MTS CONVERSION ASSAY

TSPO-pFLAG-cmv-2 or pFLAG-cmv-2 transfected C6 cells were seeded in 96-well 8 plates at a density of 1×10^4 cells for well. The following day or 96 hours after 9 transfection, culture medium was replaced with fresh medium supplemented with 10 11 different PK 11195 concentrations (1-100µM). Each cell line has been also treated with the vehicle ethanol. 24 hours after the treatment the number of living cells was 12 13 measured by quantitative colorimetric MTS assay (Promega) using a microplate reader (Wallac Victor 2, 1420 Multilabel counter, Perkin-Elmer) as previously 14 described by [15]. 15

16 Each treatment was performed in duplicate and three independent experiments were17 performed.

Graphic presentation and data analysis were performed using the non-linear
multipurpose curve-fitting Graph-Pad Prism computer program (Graph Pad
Software, version 4.0; San Diego, CA).

21

1 **RESULTS**

2 TSPO OVER-EXPRESSION IN C6 RAT GLIOMA CELLS

TSPO cDNA was isolated by RT-PCR using C6 total RNA as template, and cloned 3 in pFLAG-cmv-2 vector in frame with the nucleotide sequence encoding the eight 4 5 aminoacid FLAG peptide (at the N-terminus of TSPO). The cDNA sequence isolated from C6 cell was identical to that deposited in DDBJ/EMBL/GenBank 6 under accession number NM 012515. To assess the effectiveness of our expression 7 system, the expression of the FLAG-tagged TSPO was initially monitored by 8 9 immunoblotting on whole extract of TSPO-pFLAG-cmv-2 or pFLAG-cmv-2 cells 10 obtained 24, 48, 72 and 96 hours after transfection. Anti-FLAG antibody revealed 11 the presence of a cross-reactive band of the expected molecular weight (20 kDa) in TSPO-pFLAG-cmv-2 transfected samples. Interestingly, no SDS and beta-12 13 mercaptoethanol resistant polymers were found with anti FLAG antibodies. The 14 recombinant protein was expressed at high level in samples collected 24 hours after 15 transfection and its expression was significantly reduced in cell extracts collected 48 16 and 72 hours after transfection. Recombinant protein was no longer detectable in the 17 extracts collected 96 hours after transfection (Fig 1 A). No cross-reactive bands were 18 visible in whole extract obtained from pFLAG-cmv-2 transfected cells (control cells). 19

In order to verify if the introduction of the FLAG epitope at the N-terminus of TSPO could modify the binding affinity for the specific and selective TSPO ligand, PK11195, homogenate membranes of TSPO-pFLAG-cmv-2 transfected cells, collected 24 hours after transfection, were used for saturation studies of specific

[³H]PK11195 binding. Scatchard analysis of [³H]PK11195 binding saturation data
revealed a single population of binding sites with a dissociation constant (Kd)
value of 5.0 +/- 0.52 nM . C6 glioma cells transfected with TSPO-pFLAG-cmv-2
showed higher levels (about 2 folds) of [³H]PK specific binding sites than control
cells (Fig 1 B).

6 We also analyzed the sub-cellular localization of TSPO-pFLAG recombinant protein 7 by immunofluorescence microscopy. Anti-FLAG antibody revealed a cytoplasmic 8 localization of TSPO-FLAG protein that appeared accumulated in well-defined sub-9 cellular structures (Fig 2A). Co-localization experiments performed by using anti-10 citochrome c antibody as a mitochondrial marker (Fig 2B) revealed an exclusive 11 mitochondrial localization for TSPO-FLAG recombinant protein (Fig 2C).

12

13 TSPO OVER-EXPRESSION INCREASES THE PROLIFERATION 14 CAPABILITIES OF C6 GLIOMA CELLS

With the aim of analyzing the relationship between TSPO expression level and cell 15 growth (Fig. 3), we assessed TSPO-pFLAG-cmv-2, pFLAG-cmv-2 or pEGFP-C1 16 17 transfected cell mitotic index. As shown in Fig. 3C, TSPO-pFLAG-cmv-2 18 transfected cells showed a higher mitotic index then control cells one day after 19 transfection. This difference became less pronounced 3 days after transfection when, 20 according to western blot analysis, the expression of TSPO recombinant protein is 21 strongly reduced (data not shown). Further support for TSPO positive influence on 22 cell proliferation, derived from the analysis of TSPO-pFLAG-cmv-2, pFLAG-cmv-2 or pEGFP-C1 transfected cell proliferation curves. As reported in Fig 3A, TSPO-23

1 pFLAG-cmv-2 transfected C6 cells have a higher proliferation rate than control 2 cells. Significant differences appeared 72 hours after seeding (*p<0.05). Although 3 this data could appear in disagreement with the early effect of TSPO up-regulation on mitosis, it has to it has to be take in mind that we designed our experiments in 4 5 way to maintain cells in the exponential phase of growth up to about 120 hours. Thus, the difference in cell number (also if small) that is produced by TSPO 6 overexpression during the first days is amplified in the following days as long as one 7 of the cell line reach the confluence and reduces cell growth rate. 8

9 We also evaluated the ability of TSPO-pFLAG-cmv-2 or pFLAG-cmv-2 transfected 10 C6 cells to growth after having reached the confluence to test TSPO influence on 11 cell ability to overcome contact-induced cell growth inhibition. As reported in Fig 12 3B the number of TSPO-pFLAG-cmv-2 transfected C6 cells was 25% higher 13 $(p<1x10^{-5})$ than that of control cells.

14

15 TSPO OVER-EXPRESSION INCREASE MIGRATION RATE OF C6 16 GLIOMA CELLS

With the aim of analyzing the relationship between TSPO expression level and cell migration capabilities, we assessed the rate at which controls (pFLAG-cmv-2 and pEGFP-C1) or TSPO-pFLAG-cmv-2 transfected cells recovered a scratch done on a layer of confluent cells. Colonization of the scratch was monitored 20, 25 and 40 hours after the scratching. As reported in Fig 4, TSPO-pFLAG-cmv-2 transfected C6 cells reduced the clear area (Fig 4 A, B) and migrated into the scratch at a significantly higher rate than that of controls cells (p<0.05) (Fig 4 C).

1TSPOOVER-EXPRESSIONINCREASESTRANSMIGRATION2CAPABILITIES OF C6 GLIOMA CELLS

One of the most important characteristics of aggressive tumoral cells is the ability to 3 penetrate anatomic barriers and diffuse in cell matrix. Therefore, we assessed C6 4 5 tumor cell transmigration across artificial barriers in vitro to evaluate tumor cell invasion properties of TSPO-pFLAG-cmv-2 transfected cells and control cells. We 6 separated populations of cells able to invade a gelatin layer and counted the number 7 of cells that reached the lower side of a polycarbonate filter. As reported in Fig 5, the 8 number of cells that more actively invade the gelatin layer was significantly higher 9 10 in TSPO-pFLAG-cmv-2 transfected C6 than control cells (p<0.05).

11 TSPO OVER-EXPRESSION SENSITIZES C6 GLIOMA CELLS TO TSPO 12 SPECIFIC LIGAND TREATMENT

13 To assess whether the TSPO over-expression affect C6 cell sensitivity to the pro-14 apoptotic activity of its specific ligand PK 11195, we evaluated cell viability of 15 control and TSPO-pFLAG-cmv-2 transfected cells following 24 hours exposure to 16 micromolar increasing PK 11195 concentrations. The 24 hour treatment was performed starting from the 24th hour or from the 96th hour after transfection. As 17 18 shown in Fig 6, PK 11195 caused a concentration-dependent inhibition of C6 cell 19 viability in both TSPO-pFLAG-cmv-2 transfected and control cells treated starting from the 24th hour after transfection. Interestingly, we observed that TSPO-pFLAG-20 21 cmv-2 transfected C6 cells showed a higher sensitivity to the antiproliferative activity of PK 11195 in comparison with control cells. These differences were 22 23 observed for each PK 11195 concentration. Accordingly the strong reduction in

recombinant TSPO expression that occurs 96 hours after transfection, no more
 differences in sensitivity to PK11195 was observable when the treatment was
 initiated 96 hours after transfection.

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1 **DISCUSSION**

2 In recent years, attention has been focused on TSPO role in the regulation of cell proliferation, as several literature data report that TSPO is strongly up-regulated in 3 tumoral tissues or tumor cell lines with respect to untransformed control tissues [4], 4 5 [6], [7] and [8]. Much research has been focused on TSPO as a promising protein target for chemotherapy, suggesting its ligands as potential therapeutic drugs [13], 6 [15], [16], [31], [32] and [33]. However, only indirect data are available to elucidate 7 the meaning of TSPO up-regulation on tumor cells. TSPO is highly expressed in 8 glioma cells and to understand the role of its up-regulation in glioma aggressiveness 9 10 we produced, for the first time, genetically modified glioma cells over-expressing 11 TSPO protein and we evaluated the influence of TSPO density on the typical features that give an invasive/aggressive phenotype (i.e. proliferation, motility and 12 13 transmigration capabilities) to a tumor cell. Among glioma cell lines we selected C6 14 cells because we previously characterized in this cell line TSPO expression and sub-15 cellular localization and established cell response to TSPO ligand treatment [15] and 16 [16].

By western blot analysis, TSPO recombinant protein expression was detectable 24 hours after transfection and some of it remained up to 72 hours, a sufficiently long period to carry out functional studies. Radioligand binding assays demonstrated that the FLAG epitope addition to the amino terminus of TSPO does not change the binding affinity for this ligand. Indeed, the [³H]PK11195 binding affinity for FLAGtagged TSPO resulted comparable to Kd value previously reported for native rat C6 glioma cells [29]. Moreover, radioligand binding experiments showed an increase of

the [³H]PK11195 specific binding sites in cells expressing TSPO recombinant protein with respect to control cells, confirming western blot results. By immunofluorescence assays we also demonstrated that TSPO recombinant protein accumulated at mitochondrion level accordingly with the main distribution of TSPO in these cells, as previously demonstrated by [16].

These results demonstrate the success in producing C6 cells over-expressing a 6 recombinant TSPO protein with peculiarities similar to those of the endogenous 7 TSPO protein and pushed us to undertake functional experiments to test the 8 9 proliferation, motility and transmigration properties of C6 cells over-expressing the 10 TSPO protein. We found that TSPO over-expression increased C6 cell proliferation rate and the mitotic index as well as the cell ability to overcome the contact-11 dependent inhibition of cell growth that is linked to tumor invasion capabilities. 12 13 Moreover we found that TSPO over-expression improved the motility rate and trasmigration capability of C6 cells, demonstrating that the increase of TSPO 14 15 expression level may contribute to the acquisition of an invasive/aggressive 16 phenotype in gliomas. This is the first direct evidence that TSPO density influences 17 glioma cell aggressiveness and is consistent with previous reports that correlate 18 TSPO expression level and tumor malignancy grade [18], [23], [24] and [28].

Our data agree with recent findings, obtained in breast cancer cells, demonstrating that a siRNA mediated reduction of TSPO endogenous level and TSPO overexpression induce a decrease and an increase in cell proliferation, respectively [34]. However, all these data are in contrast with a previous report describing that TSPO knockdown by stable antisense transfection induces an increase of *in vitro*

1 tumorigenicity and a reduction of apoptosis in C6 cells [35]. These authors have 2 suggested that increased cell proliferation in TSPO knockdown cells was due to a 3 loss of TSPO-related apoptosis. According to these interpretations, we should have found an increase in apoptotic cell death and a reduced proliferation rate in TSPO 4 5 over-expressing cells. Contrarily, in our experimental condition, we did not find an increase in the number of died cells and, in addition, TSPO transfected cells 6 proliferated at higher rate than controls. Although apparently contrasting, these 7 findings could be explained by the idea that TSPO could be found, inside the outer 8 9 mitochondrial membrane, as a monomer, heteropolymer or SDS and beta-10 mercaptoethanol resistant homopolymer. In the heteropolymers, TSPO is bound to 11 the mitochondrial permeability transition pore proteins [36] and [9] and the heteropolymerization of TSPO in the pore complex is limited by the density of 12 13 VDAC [37]. The silencing of TSPO expression affects the monomeric, homopolymeric and the heteropolymeric forms. On the contrary, the over-expression 14 15 of the FLAG-TSPO in the C6 cell mitochondrial membranes preferentially acts by 16 increasing the monomeric TSPO. Indeed, the amount of the heteromeric TSPO is 17 limited by VDAC density and unlikely the number of hetheropolymers could 18 increase without a parallel increase in VDAC expression. Moreover, our western 19 blot analysis exclude that the majority FLAG-TSPO protein is expressed as a SDS 20 and beta-mercaptoethanol resistant homopolymer. This explains why the silencing 21 and up-regulation have similar effects on the cell growth. Indeed, the reduction of 22 TSPO inside the pore makes the cells more resistant to endogenous apoptotic 23 stimuli, as suggested by [35] while the increase of the monomeric form promote

cholesterol transport [36] and, as a consequence, the biogenesis of the inner
 mitochondrial membrane that positively correlate with cell proliferation, motility
 and invasiveness.

Finally, it is well known that micromolar PK 11195 concentrations induce apoptosis in C6 glioma cells [18]. In the present study we tested the PK 11195 effects on cell viability of C6 glioma cells expressing different TSPO levels. We found that TSPO over-expression causes an increased susceptibility of C6 glioma cells to the antiproliferative activity of PK 11195, suggesting that this ligand may be useful also for the therapeutic treatment of gliomas with aggressive phenotype and high TSPO density.

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1 FIGURE LEGENDS

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3 FIGURE 1: TSPO-FLAG recombinant protein expression. (A) Western blot 4 analysis of TSPO-FLAG recombinant protein in TSPO-pFLAG-cmv-2 and pFLAG-5 cmv-2 transfected cells. The molecular weight ais indicated on the right, anti ERK 6 has been used as loading control. (B) Saturation curves and Scatchard plots (inset) of specific [³H]PK 11195 binding to membranes of C6 glioma cells transfected with 7 TSPO-pFLAG-cmv-2 (♥) or with pFLAG-cmv-2 (■). Membranes were incubated 8 with increasing concentrations of [³H]PK 11195 ranging from 0.5 to 26 nM. Non 9 specific binding was measured in the presence of 1 µM PK 11195. Data are from a 10 11 single experiment carried out in triplicate. For C6 glioma cells transfected with 12 TSPO-pFLAG-cmv-2 or with pFLAG-cmv-2 the Kd values were, 6.0 and 5.0 nM and mean Bmax values were, 10.98 and 5.72 pmol/mg of protein, respectively. 13 14 FIGURE 2: Assessment of the subcellular distribution of TSPO-FLAG recombinant

FIGURE 2: Assessment of the subcellular distribution of TSPO-FLAG recombinant protein in TSPO-pFLAG-cmv-2 transfected C6 cell by fluorescent microscopy. (A) Distribution of TSPO-FLAG recombinant protein as visualized by using anti-FLAG antibody. (B) Subcellular distribution of mitochondria as revealed by using anticytochrome c antibody. (C) Merged panels A and B show that in TSPO-pFLAGcmv-2 transfected cell (arrow), TSPO-FLAG recombinant protein is expressed in the mitochondria.

FIGURE 3: Assessment of TSPO transfected cell proliferation capabilities. (A) Cell proliferation curves of pFLAG-cmv-2 and TSPO-pFLAG-cmv-2 transfected cells. Each value is expressed as mean \pm SD of three independent experiments done in triplicate. (* p< 0.05, calculated by student *t* test). (B) number of TSPO-pFLAG-

1 cmv-2 or pFLAG-cmv-2 transfected C6 cells after they had been confluent for three 2 days. Each value is expressed as mean \pm SD of the cell number counted in ten 3 independent transfections (* p<1x10⁻⁵, calculated by student *t* test). (C) Mitotic 4 index of pFLAG-cmv-2, pEGFP-C1 and TSPO-pFLAG-cmv-2 transfected cells 24 5 hours after transfection. (* p< 0.001, calculated by student *t* test between TSPO-6 pFLAG-cmv-2 and pFLAG-cmv-2)

FIGURE 4: Effect of TSPO over-expression on cell motility. (A) A representative 7 example of TSPO-pFLAG-cmv-2 transfected C6 cells recovering the scratch 8 surface. (B) Reduction of the scratch area 20, 25 and 40 hours after scratching in 9 pFLAG-cmv-2, pEGFP-C1 or TSPO-pFLAG-cmv-2 transfected cells. Mean values 10 \pm SD were obtained normalizing the scratch area at each time of analysis versus the 11 scratch area measured immediately after scratching (* p<0.05, ** p<0.01, calculated 12 13 by student t test between TSPO-pFLAG-cmv-2 and pFLAG-cmv-2). (C) Migration rate (cm/h) of TSPO-pFLAG-cmv-2, pEGFP-C1 and pFLAG-cmv-2 transfected 14 15 cells. Each bar shows the rate as mean \pm SD calculated in three point of six 16 independent transfections at 20, 25 and 40 hours (* p < 0.05, calculated by student t 17 test between TSPO-pFLAG-cmv-2 and pFLAG-cmv-2).

FIGURE 5: Effect of TSPO over expression on transmigration of C6 cells through an artificial barrier. (A) Cells were allowed to migrate through gelatin-coated filters in response to chemoattraction for 5 hours. (B) Number of cells passed through the filter has been counted in 15 microscope fields (10 X). Each bar shows the mean number \pm S.D of cells counted in three independent experiments (* p<0.05, calculated by student *t* test).

FIGURE 6: TSPO-pFLAG-cmv-2 and pFLAG-cmv-2 transfected C6 cell viability inhibition by PK 11195. Cells were treated with increasing concentrations (ranging from 1 to 100 μ M) of PK 11195 for 24 hours. After incubation, cell viability was measured by MTS conversion assay. The results are expressed as a percentage of viable cells observed after treatment with PK 11195 *vs* vehicle treated cells (100%) and shown as mean ± SD. The data are obtained from three independent experiments done in duplicate.

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