



GLT-1 promoter activity in astrocytes and neurons of mouse hippocampus and somatic sensory cortex

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GLT-1 eGFP BAC reporter transgenic adult mice were used to detect GLT-1 gene expression in individual cells of CA1, CA3 and SI, and eGFP fluorescence was measured to analyze quantitatively GLT-1 promoter activity in different cells of neocortex and hippocampus. Virtually all GFAP+ astrocytes were eGFP+; we also found that about 80% of neurons in CA3 pyramidal layer, 10–70% of neurons in I–VI layers of SI and rare neurons in all strata of CA1 and in strata oriens and radiatum of CA3 were eGFP+. Analysis of eGFP intensity showed that astrocytes had a higher GLT-1 promoter activity in SI than in CA1 and CA3, and that neurons had the highest levels of GLT-1 promoter activity in CA3 stratum pyramidale and in layer VI of SI. Finally, we observed that the intensity of GLT-1 promoter activity in neurons is 1–20% of that measured in astrocytes. These results showed that in the hippocampus and neocortex GLT-1 promoter activity is observed in astrocytes and neurons, detailed the distribution of GLT-1 expressing neurons, and indicated that GLT-1 promoter activity in both astrocytes and neurons varies in different brain regions.

Keywords: glutamate transporters, GLT-1/EAAT2, neurons, astrocytes, hippocampus, neocortex

INTRODUCTION

Glutamate is the main excitatory neurotransmitter in the mammalian CNS, where it plays a key role in numerous functions (Conti and Weinberg, 1999; Danbolt, 2001). Extracellular glutamate concentration is regulated by five high-affinity plasma membrane transporters (EAATs), which belong to SLC1 family: GLAST (EAAT1, SLC1A3), GLT-1 (EAAT2, SLC1A2), EAAC1 (EAAT3, SLC1A1), EAAT4 (SLC1A6) and EAAT5 (SLC1A7) (Kanai and Hediger, 1992, 2004; Pines et al., 1992; Storck et al., 1992; Fairman et al., 1995; Kanner, 2006; Torres and Amara, 2007). GLT-1 is responsible for the largest proportion of total glutamate uptake in the forebrain (Rothstein et al., 1996; Tanaka et al., 1997); accordingly, the lack of GLT-1 determines excitotoxicity, whereas its increase is neuroprotective (Tanaka et al., 1997; Guo et al., 2003; Rothstein et al., 2005).

Early localization studies showed that GLT-1 is primarily localized to astroglial membranes (Danbolt et al., 1992; Levy et al., 1993; Rothstein et al., 1994; Torp et al., 1994; Chaudhry et al., 1995; Lehre et al., 1995; Minelli et al., 2001). However, several studies reported the presence of GLT-1 in neurons, e.g. in pyramidal cells of hippocampal CA3 region and parietal cortex (Torp et al., 1994, 1997; Schmitt et al., 1996, 2002; Berger and Hediger, 1998; Brooks-Kayal et al., 1998; Mennerick et al., 1998; Plachez et al., 2000; Chen et al., 2004; Berger et al., 2005). In addition, GLT-1 protein has been documented in a significant percentage of axon terminals in different brain regions (Chen et al., 2004; Furness et al., 2008; Melone et al., 2009). Recently, Danbolt and co-workers re-evaluated the percentage of axon terminals expressing GLT-1 in CA1 stratum radiatum, and found that about 50% of them are GLT-1-positive (GLT-1+); they also showed that about 80% of terminals exhibit GLT-1-mediated

D-aspartate uptake, even though the density of GLT-1 molecules in neurons appears about ten times lower than in astrocytes (Furness et al., 2008). The expression of GLT-1 in such an elevated percentage of neurons may have important functional effects, and could change our understanding of the physiology of glutamatergic synapses.

In this context, it would be of some interest to know whether the promoter of the GLT-1 gene is active in neurons and its degree of activation. Here, we used GLT-1 BAC promoter reporter transgenic mice that express enhanced green fluorescent protein (eGFP) under the transcriptional control of the GLT-1 promoter (Regan et al., 2007) to quantify the distribution of cells displaying GLT-1 promoter activation in CA1, CA3 and SI cells of adult mice, and the relative levels of GLT-1 promoter activity.

MATERIALS AND METHODS

ANIMALS

Adult heterozygous GLT-1 eGFP BAC promoter reporter transgenic mice were used in this study. Heterozygous offsprings were obtained from homozygous GLT-1 eGFP BAC transgenic mice bred with C57BL/6 wild-type mice (Regan et al., 2007). Animals care and handling were approved by the Ethical Committee for Animal Research of Università Politecnica delle Marche.

Mice were anesthetized with an intraperitoneal injection of 12% chloral hydrate and perfused transcardially with a flush of saline solution, followed by 4% freshly depolymerised paraformaldehyde in 0.1 M phosphate buffer (PB). Brains were removed, post-fixed in the same fixative for 3–7 days, and cut in the parasagittal plane in 50- μ m thick sections with a Vibratome. Sections were mounted, air dried and coverslipped with Vectashield mounting medium (H-1000; Vector).

IMMUNOCYTOCHEMISTRY

Sections were first incubated in normal goat serum (NGS) (10% in PB; 1 h) and then overnight at room temperature in a solution containing primary antibodies directed to glial fibrillary acidic protein (GFAP; 1:800; Sigma; six mice, two sections/animal) or neuronal specific nuclear protein (NeuN; 1:200; Chemicon; six mice, two sections/animal). The next day, after several rinses in PB, sections were incubated in NGS (10% in PB; 20 min) and then in a solution containing an anti-mouse tetramethylrhodamine isothiocyanate (TRITC)-conjugated secondary antibody (1 h; 1:150; Molecular Probes). Sections were finally washed with PB, mounted, air dried and coverslipped with Vectashield mounting medium (H-1000; Vector). In control experiments primary antibodies were omitted.

DATA COLLECTION

Data were collected from CA1, CA3 and first somatic sensory cortex (SI) using a Leica (TCS SP2) confocal laser microscope. To improve signal/noise ratio, 10 frames of each image were averaged. For low magnification confocal microscopy, images were collected with an air dry 10× lens and aligned to obtain composite images of hippocampus and SI using Adobe Photoshop CS2 (Adobe System, San Jose, CA, USA). For high resolution confocal microscopy (applied to eGFP co-localization and intensity studies; see below) images were acquired with a HCX PL APO 63× oil-immersion lens (1.4 numerical aperture), a pinhole of 1 Airy Unit, an image size of 512 × 512 pixels and a pixel size of 0.423 μm, with the exception of the astrocyte intensity studies in which the pixel size was 0.230 μm. Acquisition of astrocytic and neuronal eGFP signals was optimized through the 'QLUT' button, which permitted direct visualization of pixel saturation; photomultiplier gain was set so that the brightest pixels were just slightly below saturation, and the offset such that the darkest pixels were just above zero. To avoid bleed-through between green and red fluorescence, images were acquired sequentially.

eGFP co-localization studies

For astrocytes, eGFP/GFAP fields were randomly selected from stratum oriens (so), pyramidale (sp), and radiatum (sr) of CA1 and CA3 (6 fields/layer/animal for each region). In SI, microscopic fields (10 fields/layer/animal) were randomly selected from layer I and VI, where GFAP immunoreactivity is robust (Hajós, 2008). For neurons, sampling of eGFP/NeuN fields of CA1 and CA3 was carried out as for eGFP/GFAP studies, whereas in SI microscopic fields were sampled in all cortical layers (10 fields/layer/animal).

eGFP intensity studies

For the analysis of eGFP intensity in both astrocytes and neurons we used eGFP/NeuN labelled sections. Sampling was carried out in all hippocampal and cortical layers (10 fields/layer/animal for each hippocampal region and 10 fields/layer/animal for SI). We used eGFP positive (+)-NeuN+ cells for the analysis of eGFP intensity in neurons, and eGFP+-NeuN- cells for that of astrocytes. The rationale for considering eGFP+-NeuN- cells as astrocytes is that in pilot studies we observed that in layers I-VI of SI and in strata oriens, pyramidale, and radiatum of CA1 and CA3 CNPase+ cells (i.e., oligodendrocytes) and CD11b+ cells (i.e., microglial cells) were never eGFP+. Details on immunocytochemical procedure are given in the legend to **Figure 2**.

DATA ANALYSIS

eGFP co-localization studies

For each merged image, a threshold level of fluorescence for green and red channel was determined by setting a background intensity value. Background intensity values for green and red channel corresponded to 0.5 of the median of single channel intensity (estimated with Adobe Photoshop CS2); then, background values were subtracted from each channel, yielding a thresholded image. Background intensity was chosen within a range of values (from 0.25 to 0.75 of the median) which did not affect measurement accuracy (Melone et al., 2005; Ronneberger et al., 2008); background subtraction resulted in null intensity pixel value in eGFP- structures, e.g. blood vessels (Regan et al., 2007).

Thresholded images were split into green and red channel and then analyzed with NIH ImageJ Software (<http://rsb.info.nih.gov/ij>). A region of interest (ROI) of 5 × 5 μm was used to assess pixel positivity within neuron somata on the green channel, while a ROI of 3 × 3 μm was used for astrocytes. Cells with an average pixel intensity ≥ 1 arbitrary units were considered positive for eGFP expression and were therefore included in the analysis. Co-localization of eGFP+ cells with NeuN and GFAP was calculated as the percentage of eGFP+/NeuN+ and eGFP+/GFAP+ cells on the total of NeuN+ and GFAP+ cells.

eGFP cellular intensity studies

In these studies, as in eGFP co-localization studies, background value was set at 0.5 of the median level of the entire image. In a series of pilot studies, we studied the relationship between eGFP intensity values of cells and mean intensity of the whole microscopic field; we found that astrocytic and neuronal intensity levels measured were positively correlated to the mean intensity levels of acquired images ($r = 0.965$ for astrocytes, $P = 0.008$; $r = 0.860$ for neurons, $P = 0.028$; $n = 10$ images). Therefore, in order to obtain intensity values not affected by the mean intensity of images, we transformed intensity measures by matching the cellular with the background intensity ($I_{\text{cell}}/I_{\text{background}}$), as described by Marrs et al. (2001). After this transformation, correlation between intensity values was not significant ($P = 0.226$ for astrocytes; $P = 0.103$ for neurons), thus supporting the validity of the procedure. Then we normalized the data using the formula:

$$NI_v = \frac{v - v_{\min}}{v_{\max} - v_{\min}} (NI_{\max} - NI_{\min}) + NI_{\min}$$

where v stood for the value of $I_{\text{cell}}/I_{\text{background}}$ to be normalized, NI_v for Normalized Intensity of v , v_{\min} and v_{\max} for minimum and maximum values of v , NI_{\min} and NI_{\max} are the new minimum and maximum normalized values of v , i.e. 0 and 1 (Han and Kamber, 2006).

STATISTICAL ANALYSIS

Comparison between CA1, CA3 and SI (obtained by collapsing data from all layers of each region) was performed by one way ANOVA ($\alpha = 0.05$) with Bonferroni-Dunn post test ($\alpha = 0.017$ due to Bonferroni correction for multiple comparisons). Comparisons among layers within CA1 and CA3 were performed with one way ANOVA ($\alpha = 0.05$) with Bonferroni-Dunn post test ($\alpha = 0.017$). Comparison between CA1 and CA3 data was assessed also with

two way ANOVA considering hippocampal regions as the between factor and hippocampal layers as the within factor. Post tests used to study significant differences among levels of the between and the within factors were paired t-test and one way ANOVA. Comparison among layers of SI was assessed by one way repeated measures ANOVA ($\alpha = 0.05$) and Bonferroni-Dunn post test ($\alpha = 0.003$) was applied. Statistical analysis was performed with SPSS (v.13.0; SPSS Inc, IL, USA).

RESULTS

In line with the description of Regan et al. (2007), we observed numerous cells displaying GLT-1 promoter activity (eGFP+ cells) in SI and hippocampus (Figures 1A,B). Double-labelling studies with GFAP and NeuN antibodies (see below for details) showed that eGFP+ cells were in most cases astrocytes and not rarely neurons (Figures 1C,D), whereas double-labelling studies with CNPase and CD11b antibodies revealed that in SI and hippocampal gray matter eGFP+ cells were never oligodendrocytes or microglial cells (Figure 2). In addition, we observed that the intensity of eGFP+ cells varied remarkably, with neurons exhibiting low eGFP fluorescence intensity (Figures 1E,F). We therefore studied quantitatively both the presence and the intensity of eGFP fluorescence in astrocytes and neurons of CA1, CA3 and SI of adult GLT-1-eGFP BAC reporter mice.

GLT-1 PROMOTER ACTIVITY IN ASTROCYTES AND NEURONS

We first examined the presence of eGFP reporting for GLT-1 in GFAP+ and NeuN+ cells of CA1, CA3 and SI (Table 1). GFAP/eGFP co-localization studies (Figure 1C) revealed that in CA1 and CA3 (stratum oriens, so; stratum pyramidale, sp; stratum radiatum, sr), and SI (layers I and VI) $96.5 \pm 1.4\%$, $95.5 \pm 1.3\%$, and $97.5 \pm 0.04\%$ of GFAP+ cells were eGFP+, in the order. Statistical analysis showed that GFAP/eGFP co-localization was not significantly different in these regions. NeuN/eGFP co-localization studies (Figure 1D) showed that in CA1, CA3 and SI $5.5 \pm 1.6\%$, $34.5 \pm 9.0\%$, and $37.0 \pm 3.7\%$ of all NeuN+ neurons were eGFP+, in the order (Figure 3). NeuN/eGFP co-localization was higher in CA3 and SI than in CA1 ($P = 0.0008$ and $P < 0.0001$, respectively), and in CA3 it was similar to SI ($P = 0.7$).

Next, we studied the laminar distribution of GFAP/eGFP+ and NeuN/eGFP+ cells in CA1, CA3 and SI. The distribution of GFAP/eGFP+ cells was similar in different layers of CA1 and CA3 (so, sp, and sr) and SI (layers I and VI), whereas that of NeuN/eGFP+ cells exhibited striking differences. In particular, in CA1 NeuN/eGFP+ cells were $6.5 \pm 2.3\%$ in so, $1.8 \pm 0.9\%$ in sp, and $8.2 \pm 4.1\%$ in sr (Figures 4A,D, white columns); in CA3 they were $10.9 \pm 4.5\%$ in so, $85.8 \pm 2.9\%$ in sp, and $6.7 \pm 2.8\%$ in sr (Figures 4B,D, black columns); and in SI they were $13.8 \pm 6.2\%$, $18.6 \pm 3.5\%$, $31 \pm 4.7\%$, $51.3 \pm 6.6\%$, $37.1 \pm 1.4\%$, and $70 \pm 3.0\%$ in layers I-VI (Figures 4C,E). NeuN/eGFP co-localization was different in CA3 layers ($P < 0.0001$), with sp exhibiting the highest degree of co-localization ($P < 0.0001$ with CA3 so and sr), but not in CA1 layers ($P = 0.3$), and it was higher in CA3 sp than in the same layer of CA1 ($P < 0.0001$) (Figure 4D). In SI, NeuN/eGFP co-localization was different between layers ($P < 0.0001$), with layer VI significantly higher than layers I-III and V ($P < 0.0001$) and layer IV ($P = 0.0007$) (Figure 4E).

INTENSITY OF eGFP-GLT-1 FLUORESCENCE IN ASTROCYTES AND NEURONS

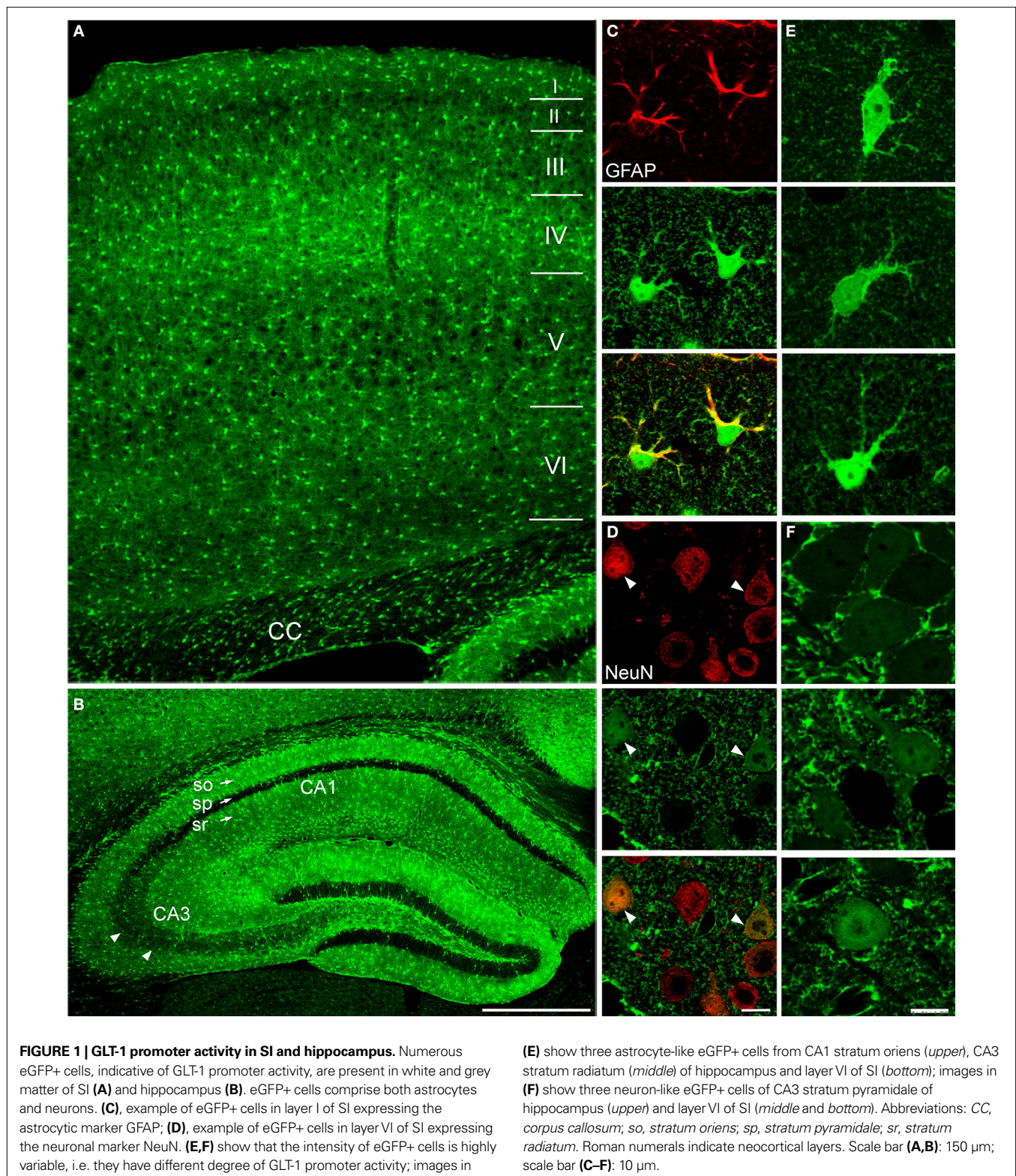
Since astrocytes and neurons displayed different levels of eGFP fluorescence intensity, i.e., of GLT-1 promoter activity (Figures 1E,F), we analyzed eGFP fluorescence intensity in different layers of CA1, CA3, and SI (Table 1). To do this, we measured fluorescence intensity within eGFP+ cell bodies and calculated its normalized intensity (NI), as described in Material and Methods. Mean NI of astrocytes was different between regions ($P < 0.0001$): it was higher in SI than in CA1 ($\sim 35\%$, $P < 0.0001$) and CA3 ($\sim 47\%$, $P < 0.0001$), but it was similar in CA1 and CA3 ($P = 0.13$) (Figure 5A). Mean NI of neurons was different between regions ($P = 0.0492$) and higher in SI than in CA1 (up to $\sim 50\%$, $P < 0.0146$) but not in CA3 (Figure 5B).

In CA1 and CA3, analysis of astrocytic NI revealed laminar differences ($P = 0.0002$ for CA1; $P = 0.0015$ for CA3); NI was lower in astrocytes of so than in those of sp ($\sim 15\%$, $P < 0.0001$) and sr ($\sim 10\%$, $P = 0.0031$) of CA1, whereas it was lower in astrocytes of sr of CA3 than in so ($\sim 13\%$, $P = 0.0152$) and sp ($\sim 20\%$, $P = 0.0004$) (Figure 6A). Two way ANOVA revealed significant differences in astrocytic NI between CA1 and CA3 ($P < 0.0001$): each CA1 layer had an higher NI than the corresponding CA3 layer ($\sim 10\%$ for so, $P = 0.0075$; $\sim 16\%$ for sp, $P = 0.0114$; $\sim 14\%$ for sr, $P = 0.0001$) (Figure 6A). The discrepancy between the results of one and two way ANOVA (see above) is determined by the fact that in one way ANOVA data from so, sp, and sr were collapsed, whereas in two way analysis, mean values of each layer were compared. In SI, no differences of astrocytic NI were noted. Neuronal NI in so, sp, and sr of CA1 and CA3 showed no laminar differences; however, in CA3 sp NI was higher than in CA1 sp ($\sim 60\%$, $P = 0.0007$) (Figure 6B). In SI, intensity of eGFP+ neurons showed significant laminar differences ($P < 0.0001$): layer VI was higher than layers I ($\sim 72\%$, $P < 0.0001$), II ($\sim 74\%$, $P = 0.0001$), III ($\sim 70\%$, $P = 0.0001$), IV ($\sim 55\%$, $P = 0.0002$), and V ($\sim 46\%$, $P = 0.0017$) (Figure 6C).

Finally, to compare GLT-1 promoter activity in neurons and astrocytes, we calculated the ratio between neuronal and astrocytic NI ($NI_{neurons}/NI_{astrocytes}$; NI_n/NI_a). In CA1, CA3 and SI, mean NI_n/NI_a was 0.027 ± 0.004 , 0.055 ± 0.01 , and 0.033 ± 0.003 , in the order; these values indicated that GLT-1 promoter activity in neurons was 2.7, 5.5, and 3.3% of those measured in astrocytes. Statistical analysis showed a significant difference between regions ($P = 0.0275$), with CA3 NI_n/NI_a higher than CA1 ($P = 0.0150$) but not than SI NI_n/NI_a . Laminar evaluation of NI_n/NI_a revealed differences in SI ($P < 0.0001$), with NI_n/NI_a higher in layer VI (0.07 ± 0.01) than in layers I (0.01 ± 0.005 , $P < 0.0001$), II (0.017 ± 0.003 , $P < 0.0001$); III (0.023 ± 0.006 , $P < 0.0001$); IV (0.037 ± 0.008 , $P = 0.0003$), and V (0.036 ± 0.007 , $P = 0.0003$).

DISCUSSION

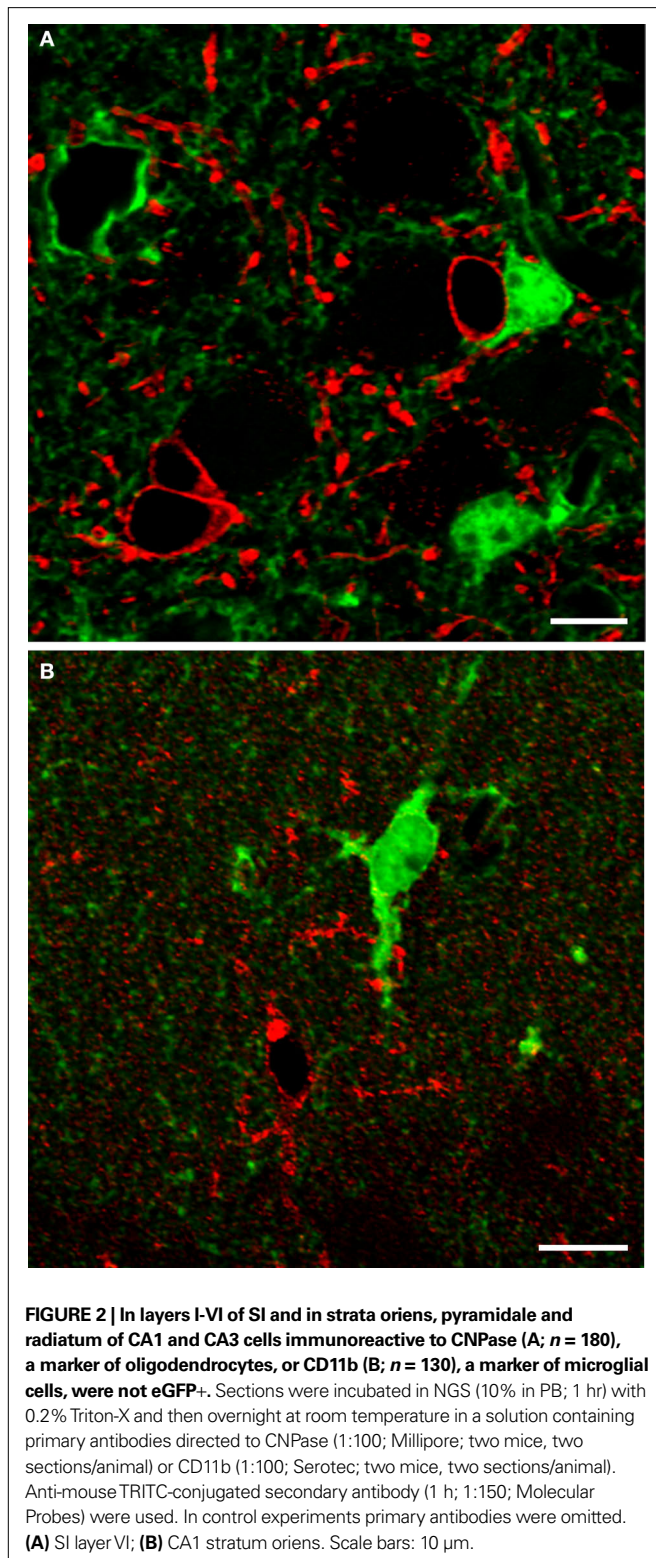
These studies in GLT-1 eGFP BAC reporter transgenic mice showed that virtually all GFAP+ astrocytes and numerous neurons exhibited GLT-1 promoter activity; that the degree of activation was much higher in astrocytes than in neurons; and that GLT-1 promoter activity was highly variable across regions and layers.



GLT-1 PROMOTER ACTIVITY IN ASTROCYTES AND NEURONS

There is a long history, albeit a controversial one, on the cellular localization of GLT-1, the major glutamate transporter in the mammalian brain, and particularly on its postulated neuronal expres-

sion, as briefly summarized in Introduction. Here, we have used GLT-1 eGFP BAC reporter transgenic mice to investigate GLT-1 promoter activation in astrocytes and neurons, as this preparation allows both the visualization of individual cells whose GLT-1

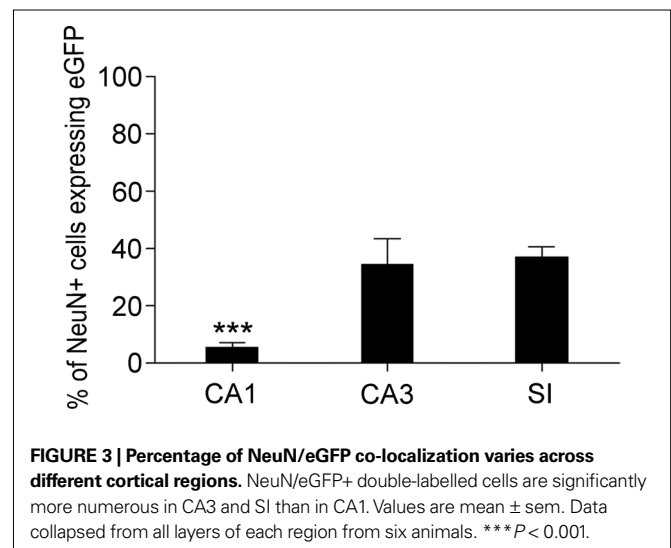


promoter is active and its quantitation. The latter feature appears particularly attractive on considering that GLT-1 eGFP fluorescence appears well correlated to GLT-1 mRNA (Yang et al., 2009), GLT-1 protein expression (Regan et al., 2007), and, most importantly, to GLT-1 function (Regan et al., 2007).

Table 1 | Number of cells studied in co-localization and eGFP intensity studies.

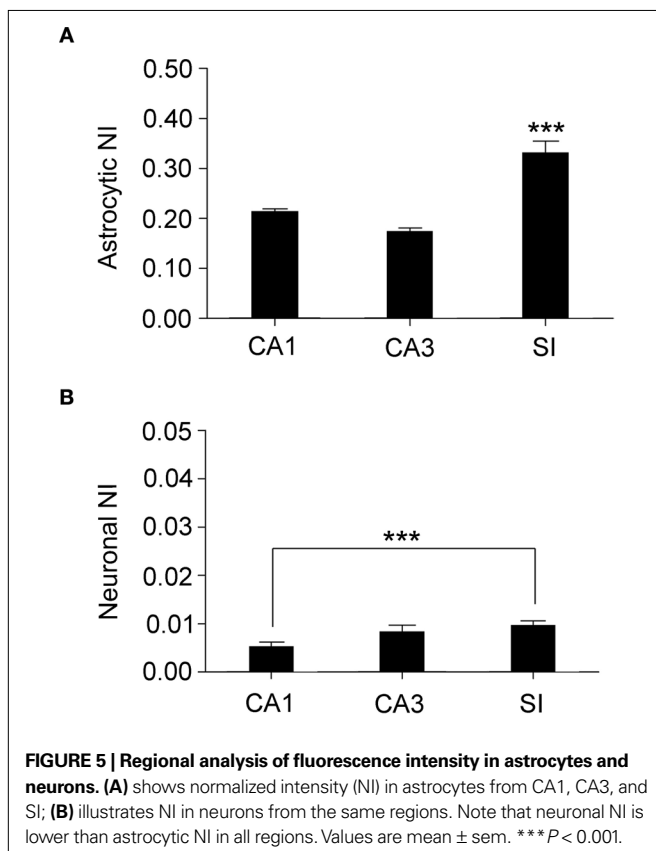
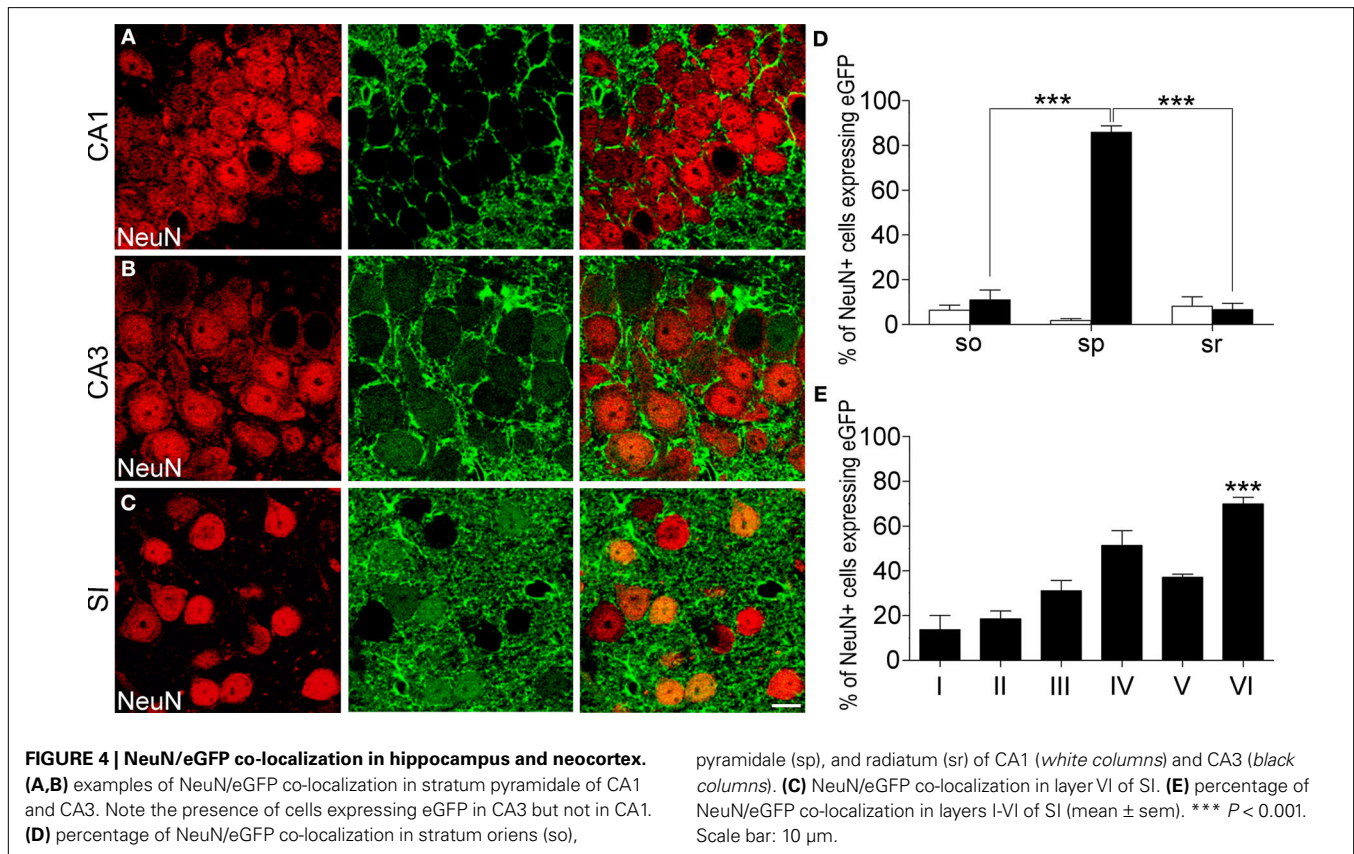
	Co-Localization		eGFP Intensity	
	Astrocytes	Neurons	Astrocytes	Neurons
CA1	392 (so 150; sp 70; sr 172)	2616 (so 116; sp 2420; sr 80)	261 (so 104; sp 76; sr 91)	33 (so 10; sp 15; sr 8)
CA3	404 (so 132; sp 67; sr 205)	2376 (so 69; sp 2191; sr 116)	274 (so 92; sp 75; sr 107)	1551 (so 7; sp 1536; sr 8)
SI	464 (I 167; VI 297)	8039 (II 113; II 1152; III 1595; IV 2073; V 1245; VI 1861)	311 (I 67; II 48; III 43; IV 51; V 52; VI 50)	3453 (I 18; II 207; III 463; IV 1046; V 432; VI 1287)

Data from 6 animals. Roman numbers indicate cortical layers. Abbreviations: so, stratum oriens; sp stratum pyramidale; sr, stratum radiatum.



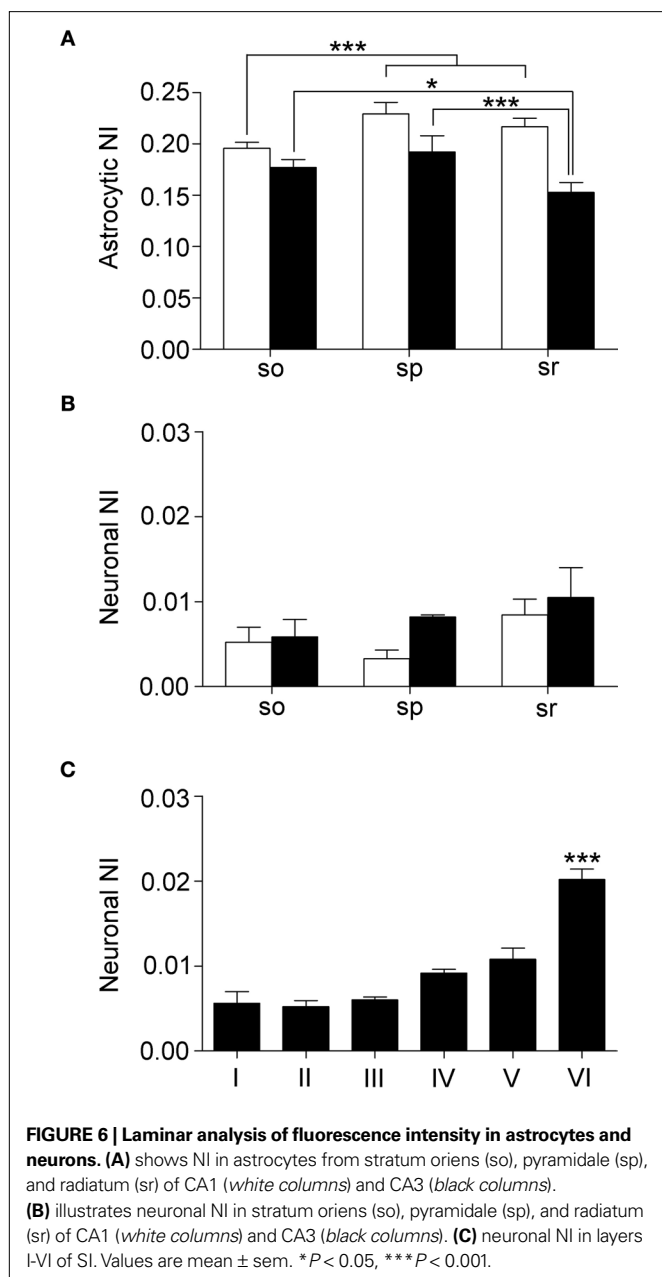
We showed that virtually all GFAP+ astrocytes expressed GLT-1 promoter activity, in line with existing data on GLT-1 localization in hippocampus and neocortex (Rothstein et al., 1994; Chaudhry et al., 1995; Lehre et al., 1995; Schmitt et al., 1996; Danbolt, 2001), and confirmed that astrocytes are the predominant cell type expressing GLT-1. We also showed that intensity of eGFP-GLT-1 fluorescence in astrocytes was much higher than in neurons; in CA3 sp and in layer VI of SI fluorescence intensity in neurons represented 4% and 7% of astrocyte values, respectively. Interestingly, Furness et al. (2008) have recently estimated that GLT-1 protein levels in axon terminals are about 10% of those measured in astrocytes. The similarity between our data and those of Furness et al. (2008) provides further support to the notion that the degree of GLT-1 promoter activity appears correlated to protein expression (Regan et al., 2007).

We have also observed that a considerable number of neurons in CA3 pyramidal layer (~80%) and in layers I–VI of SI (10–70%) exhibits GLT-1 promoter activity. These results extend those of previous studies that described qualitatively the presence of GLT-1 mRNA



in hippocampal and neocortical neurons (Torp et al., 1994, 1997; Schmitt et al., 1996; Berger and Hediger, 1998; Chen et al., 2004; Berger et al., 2005), and are compatible with the unexpected observation that $\sim 3/4$ of nerve terminals in CA1 stratum radiatum (Schaffer collaterals from CA3 pyramidal neurons) exhibit GLT-1-mediated D-aspartate uptake (Furness et al., 2008). Therefore, it is conceivable that the activation of eGFP GLT-1 promoter observed in more than 80% of pyramidal neurons of CA3 may be translated into GLT-1 protein in nerve terminals forming synapses in CA1, although this remains to be demonstrated. The discrepancy between the distribution of GLT-1 promoter activity revealed by eGFP positivity and that of GLT-1 protein reported in numerous immunocytochemical studies is conceivably related to the low levels of GLT-1 protein found in neurons, which makes it difficult to detect it immunocytochemically (Schmitt et al., 1996; Chen et al., 2004; Furness et al., 2008). This view is supported by the observation that eGFP-GLT-1 fluorescence in neurons was in general very low (*present results*) and that an high resolution confocal microscopy analysis of terminals expressing vesicular glutamate transporter 1 (VGLUT1) in CA1 stratum radiatum showed that only few of them ($<5\%$) were also GLT-1 positive (de Vivo, Melone, Rothstein, and Conti *unpublished observations*). Alternatively, it is possible that in neurons exhibiting low levels of eGFP-GLT-1 fluorescence the activation of the promoter is not followed by production of GLT-1 protein.

The percentage of neurons exhibiting eGFP-GLT-1 activity was significantly higher in CA3 pyramidal neurons and in layer VI neurons than in those of other regions or layers. Whereas CA3 pyramids were known to express GLT-1 at their axon terminals (Chen et al.,



2004; Furness et al., 2008), the unexpected presence of numerous layer VI neurons showing eGFP-GLT-1 fluorescence raises stimulating questions on their functional significance. Since most layer VI neurons project to the thalamus (Jones, 1984) and contribute

to the filtering of the thalamo-cortical flow of information, it is conceivable that GLT-1 may play a role in the gating of thalamically-mediated afferent information to the neocortex.

VARIATIONS OF eGFP-GLT-1 INTENSITY

Quantitative analysis of eGFP-GLT-1 fluorescence intensity in hippocampal and neocortical neurons and astrocytes revealed significant variations across brain regions and layers as well as between cells of the same region or layer, as previously noticed (Rothstein et al., 1994; Lehre et al., 1995; Milton et al., 1997; Regan et al., 2007; Holmseth et al., 2009). Mean normalized fluorescence values indicated that astrocytes in SI had a higher GLT-1 promoter activity than CA1 and CA3 astrocytes (SI > CA1 > CA3). Moreover, laminar analysis of astrocytes eGFP promoter activation in different SI layers showed a similar degree of activation, whereas in CA1 and CA3 significant differences were observed between strata oriens, pyramidale and radiatum. Neurons had the highest level of eGFP fluorescence in layer VI of SI and in CA3 stratum pyramidale, and showed laminar differences only in SI.

A large number of endogenous and exogenous molecules regulates GLT-1 expression in *in vitro* and *in vivo* conditions (Gegelashvili and Schousboe, 1997; Robinson, 1998; Anderson and Swanson, 2000; Beart and O'Shea, 2007; Lauriat and McInnes, 2007). In human astrocytes several GLT-1 regulators, including EGF, TGF- α , and dBcAMP, exert their effects by acting on the GLT-1 promoter (Su et al., 2003; Li et al., 2006). If the same transcriptional mechanism operates in neurons, the high variability of eGFP GLT-1 observed here in astrocytes and neurons reflects in all likelihood the differential regulation of GLT-1 promoter. It is conceivable that such a differential regulation, though largely constitutive, is at least in part induced dynamically by differential homeostatic needs in both physiological and pathophysiological conditions, in line with the crucial role GLT-1 plays in regulating extracellular glutamate and in shaping excitatory transmission (Tong and Jahr, 1994; Conti and Weinberg, 1999; Danbolt, 2001; Tzingounis and Wadiche, 2007). In this context, it is worth mentioning that a few dozen transporters distributed quasi-randomly inside the cleft attenuate AMPARs activation (Zheng et al., 2008), and that GLT-1 up-regulation affects synaptic plasticity (Omrani et al., 2009).

Thus, the variability of eGFP intensity described here raises the possibility that GLT-1 eGFP BAC reporter transgenic mice may reveal themselves useful tools to investigate quantitatively the regulation of GLT-1 promoter activation in *in vivo* physiological or pathophysiological conditions.

ACKNOWLEDGMENTS

Supported by MIUR and Università Politecnica delle Marche.

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- Conflict of Interest Statement:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.
- Received: 05 November 2009; paper pending published: 25 November 2009; accepted: 23 December 2009; published online: 11 January 2010.
- Citation: de Vivo L, Melone M, Rothstein JD and Conti F (2010) GLT-1 promoter activity in astrocytes and neurons of mouse hippocampus and somatic sensory cortex. *Front. Neuroanat.* 3:31. doi: 10.3389/neuro.05.031.2009
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