

# Modulation of GABA-A receptors of astrocytes and STC-1 cells by taurine structural analogs

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**Abstract** Taurine activates and modulates GABA receptors in vivo as well as those expressed in heterologous systems. This study aimed to determine whether the structural analogs of taurine: homotaurine and hypotaurine, have the ability to activate GABA-A receptors that include GABA $\rho$  subunits. The expression of GABA-A receptors containing GABA $\rho$  has been reported in the STC-1 cells and astrocytes. In both cell types, taurine, homo-, and hypotaurine gated with low efficiency a picrotoxin-sensitive GABA-A receptor. The known bimodal modulatory effect of taurine on GABA $\rho$  receptors was not observed; however, differences between the activation and deactivation rates were detected when they were perfused together with GABA. In silico docking simulations suggested that taurine, hypo-, and homotaurine do not form a cation– $\pi$  interaction such as that generated by GABA in the agonist-binding site of GABA $\rho$ . This observation complements the electrophysiological data suggesting that taurine and its analogs act as partial agonists of GABA-A receptors. All the observations above suggest that the structural analogs of taurine are partial agonists of GABA-A receptors that occupy the agonist-binding site, but their structures do not allow the proper interaction with the receptor to fully gate its Cl $^-$  channel.

**Keywords** Astrocytes · Electrophysiology · GABA $\rho$  · Homotaurine · Hypotaurine

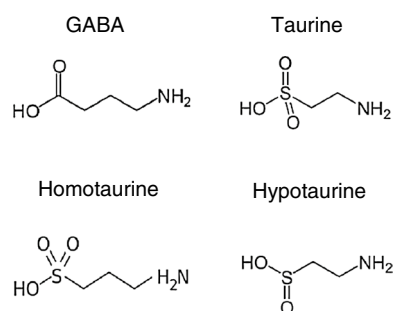
## Abbreviations

DMEM	Dulbecco's modified eagle's medium
FBS	Fetal bovine serum
GABA	$\gamma$ -aminobutyric acid
GLIC	<i>Gloeobacter</i> ion channel
EGTA	Ethylene glycol tetraacetic acid
GluCl	Glutamate-gated ion channel
HEPES	2-[4-(2-hydroxyethyl)piperazin-1-yl]ethanesulfonic acid
ICM	Internal coordinate mechanics
RsT	Rise time constant
STC-1	Secretin tumor cell line
TPMPA	(1,2,5,6-Tetrahydropyridin-4-yl)methylphosphinic acid

## Introduction

Taurine is one of the most abundant amino acids in the brain and spinal cord, leukocytes, heart and muscle cells, the retina, and indeed almost every tissue throughout the body; furthermore, taurine is known to have preventive effects against stroke and cardiovascular diseases in experimental models as well as in humans (Yamori et al. 2010; Ripps and Shen 2012). This amino acid lacks the carboxyl group and contains a sulfonate group instead; it does not form part of proteins. Taurine exerts multiple functions in the central nervous system including roles in cell volume regulation and neurotransmission. In neurons and astrocytes, taurine activates with low efficiency ionotropic GABA and glycine receptors (Pasantes-Morales et al. 1990, Moran et al. 1997; ElIdrissi and Trenkner, 2004; LeCorronc et al. 2011). Electrophysiological and competitive binding assays have indicated that taurine is a weak agonist of the GABA-A receptor complex, interacting with

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**Fig. 1** Structure of agonists of the GABA-A receptor used in this study

the muscimol-displaceable, bicuculline-sensitive GABA-binding sites thereby allosterically modulating the binding of benzodiazepines (Bureau and Olsen 1991; Frosini et al. 2003).

GABA-A receptor-mediated inhibitory effects of exogenously added taurine have been reported in several brain regions such as the cerebellum, hippocampus, striatum, and cerebral cortex (Chan-Palay et al. 1982; Mori et al. 2002; Wu and Xu 2003). Experiments with recombinant GABA receptors have revealed the ability of taurine to interact with receptors composed of different combinations of subunits (Horikoshi et al. 1998; Hadley and Amin 2007). In cloned GABA $\rho$ 1 subunits, taurine induces a bimodal effect, by modulating the GABA currents negatively when it is present at high concentrations but positively at low concentrations (Ochoa-de la Paz et al. 2008). This paradoxical effect could be explained by the transition of the taurine molecule from the *cis* to the *trans* conformation upon concentration changes of the molecule in solution. GABA $\rho$ 1 receptors are expressed mainly in retina, although many other areas of the central nervous system express this GABA-A subunit to a lesser extent (Martínez-Delgado et al. 2008). GABA $\rho$ 1, as well as GABA $\rho$ 2 and GABA $\rho$ 3, forms homo-oligomeric receptors when expressed in heterologous systems; upon activation by the agonist the receptor generates Cl<sup>−</sup> currents and exhibits virtually no desensitization. Thus, to explore if the dual effect of taurine on GABA $\rho$ 1 extends to other sulfonated amino acids, in this work we tested two structural analogues of taurine: hypotaurine and homotaurine (Fig. 1).

Hypotaurine is a strong antioxidant, scavenging OH<sup>−</sup> radicals with its sulfur group; this amino acid accumulates in the mammalian reproductive system where it acts as an osmolyte (Fontana et al. 2004). Homotaurine has been extensively studied because it binds A $\beta$ -amyloid in its soluble form; thus, its use was proposed to prevent Alzheimer's disease. Unfortunately, clinical trials of this molecule have not been conclusive (Santa-María et al. 2007).

In this study, we tested the action of the taurine analogs on the cell line STC-1, cells known to express a variety of

GABA-A subunits (Glassmeier et al. 1998; Jansen et al. 2000); in addition, we tested them in a primary culture of astrocytes from cerebellum, in which GABA-A, GABA $\rho$ 1 and GABA $\rho$ 2 are known to be expressed in vitro and in situ (Bormann and Kettenmann 1988; Martínez-Delgado et al. 2011). Finally, we generated in silico a structural model that explains the binding of homotaurine and hypotaurine to the GABA-binding site.

## Materials and methods

### Electrophysiology and pharmacology

STC-1 cells were maintained in Dulbecco's modified eagle's medium (DMEM) supplemented with 15 % horse serum, 2.5 % fetal bovine serum (FBS), 100 U/mL penicillin, and 100  $\mu$ g/mL streptomycin and maintained at 37 °C and 5 % CO<sub>2</sub> (Machuca-Parra et al. 2013). For the astrocyte primary culture, one newborn CD1 mouse was decapitated for each cellular culture; the cerebellum was removed, and dissociated cells were cultured in DMEM supplemented with 10 % FBS, 2 mM glutamine, 100 U/mL penicillin, and 100  $\mu$ g/mL streptomycin (Reyes-Haro et al. 2005; Martínez-Delgado et al. 2011). All experiments were conducted according to the guidelines of the National Institutes of Health Guide for Care and use of Laboratory Animals and approved by the Institutional Animal Care and Use Committee of the Universidad Nacional Autónoma de México.

Membrane currents were recorded using the whole-cell voltage-clamp technique (Hamill et al. 1981). A fragment of coverslip, on which astrocytes or STC-1 cells were cultured for 4–8 days, was placed in a recording chamber (300  $\mu$ l for STC-1 cells or 500  $\mu$ l for astrocytes) on the stage of an inverted microscope (Olympus IX-70, Japan). The recording chamber was continuously superfused (5 ml/min for STC-1 cells and 2 ml/min for astrocytes) with standard bathing solution containing (in mM): 150 NaCl, 2.5 KCl, 2 CaCl<sub>2</sub>, 1 MgCl<sub>2</sub>, 10 HEPES, and 10 glucose (pH 7.3). Recording pipettes were made with borosilicate capillaries (Sutter Instruments) and filled with a solution containing (in mM): 130 KCl, 0.5 CaCl<sub>2</sub>, 5 EGTA, 2 MgCl<sub>2</sub>, 10 HEPES, 3 ATP-Mg, and 0.4 GTP-Mg (pH 7.3), having a resistance of 3–5 M $\Omega$ . All experiments were carried at room temperature (20–23 °C). Currents were recorded with an Axopatch 200B amplifier (Axon Instruments, CA, USA) or EPC 10 amplifier (HEKA), filtered at 3 kHz, and acquired at 10 kHz. The data were analyzed with pClamp 8.2 (Axon Instruments, CA, USA) or TIDA software (5.19) and Microcal Origin 7 software (OriginLab, MA, USA). The data are given as mean  $\pm$  standard error. Comparison of the mean values among groups was performed by the Student's *t* test, where *p* < 0.05 was taken to indicate

a statistically significant difference. The rise time ( $R_s$ ) and decay time constants ( $\tau$ ) were calculated with Clampfit or TIDA software.

### Homology modeling and docking

We used the program MODELLER v 9.10 to develop a homology model of the human GABA $\rho$ 1 subunit, employing the high-resolution crystal structures of the *C. elegans* glutamate-gated chloride channel (GluCl), the pentameric ligand-gated ion channel of *Gloeobacter violaceus* (GLIC), and the refined structure of the nicotinic acetylcholine receptor, downloaded from the Brookhaven Protein Data Bank (PDB) using the accession numbers 3RIF, 3EHZ, and 2BG9, respectively (Brejc et al. 2001; Bocquet et al. 2009; Hibbs and Gouaux 2011). The model of GABA $\rho$ 1 was then projected fivefold on the pentameric structure of the GluCl channel to build a homopentamer of the GABA $\rho$ 1 receptor. The amino acid sequence was obtained from the Uniprot Server (protein accession number: P24046). The few observed clashes were corrected using GROMOS96 (part of DeepView/Swiss-Pdb Viewer v4.0.1 software) and refined by energy minimization using Molsoft ICM v3.5. All the ligand molecules were downloaded and edited from the ChemSpider database (IDs: Taurine, 1091; Hypotaurine, 96959; Homotaurine, 1584). For Docking, we used Pocket Finder in Internal Coordinate Mechanics (ICM) for automatic detection of small molecule-binding sites defined on an interface of the cytoplasmic domain, selecting the best fitting site for each superposed molecule in the electron density, and optimized on the model (ICM score). The images were generated using the PyMOL Molecular Graphics System, Version 1.3, Schrödinger, LLC.

## Results

### Taurine and analogs act as agonists of GABA-A receptors in STC-1 cells and astrocytes

To determine whether taurine and its structural homologs activate GABA-A receptors containing GABA $\rho$  subunits we tested astrocyte primary cultures from cerebellum and STC-1 cells, because they endogenously express functional GABA $_A$  receptors that include GABA $\rho$  subunits.

GABA generated larger currents in astrocytes (GABA 50  $\mu$ M,  $506 \pm 68$  pA;  $n = 9$ ) (Fig. 2a) than in STC-1 cells (GABA 100  $\mu$ M,  $150 \pm 29$  pA;  $n = 23$ , Fig. 2c), despite the fact that we doubled the concentrations of the agonist. A similar pattern for taurine (1 mM) was observed in astrocytes ( $193 \pm 98$  pA;  $n = 6$ ) and STC-1 cells ( $36 \pm 6$  pA;  $n = 6$ ) (Fig. 2b, d). Homotaurine

evoked currents with an amplitude of  $891 \pm 154$  pA ( $n = 4$ ) in astrocytes and  $93 \pm 13$  pA ( $n = 6$ ) in STC-1 cells, while hypotaurine (1 mM vs 3 mM) evoked currents with an amplitude of  $165 \pm 68$  pA ( $n = 9$ ) in astrocytes and  $196 \pm 43$  pA ( $n = 6$ ) in STC-1 cells (Fig. 2b, d). All the currents were almost completely blocked (>85 %) by picrotoxin (100  $\mu$ M), a selective antagonist of GABA-A receptors (Table 1) (Fig. 2a, c). Altogether, these results indicate that taurine, hypotaurine, and homotaurine activate GABA-A receptors in cerebellar astrocytes and STC-1 cells, with higher affinity for those expressed by astrocytes.

### Modulation of GABA $_A$ receptors by taurine and its analogs

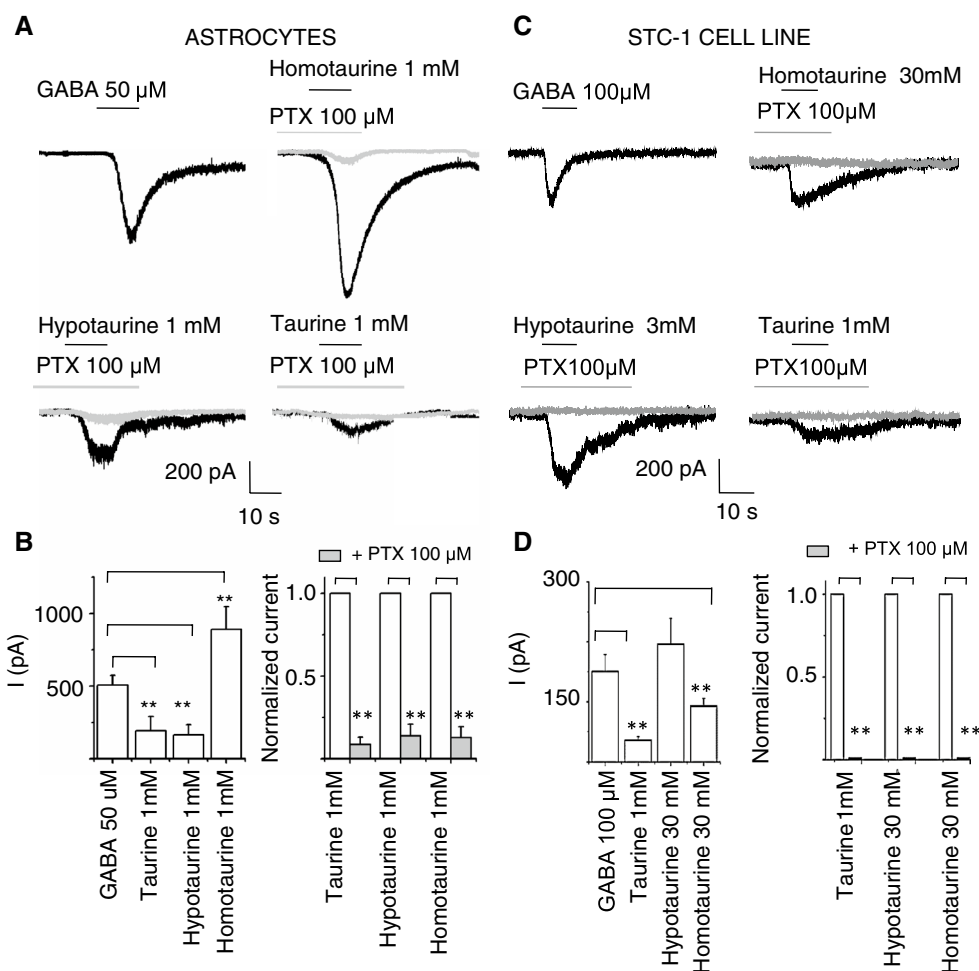
We tested the modulatory effect of taurine and its analogs on GABA-evoked currents in astrocytes and STC-1 cells (Fig. 3a, e). In astrocytes, taurine or one of the analogs was co-applied at 1 mM with 50  $\mu$ M GABA; the current amplitudes were not different from the control (50  $\mu$ M GABA), and the deactivation constant ( $\tau$ ) remained unaffected (Fig. 3b, c); however, the rise time constant ( $R_sT$ ) estimated for GABA ( $5.39 \pm 0.68$ ) increased when taurine was present in the bath (+57 %;  $n = 5$ ;  $p = 0.0204$ ) and decreased with homotaurine (−40 %;  $n = 7$ ;  $p = 0.034$ ), while hypotaurine did not affect the  $R_sT$  (Fig. 3d). On the other hand, the amplitude of GABA currents (100  $\mu$ M) was not modified significantly for STC-1 cells with the co-application of taurine or any of the analogs (Fig. 3f). Nevertheless, the estimated  $\tau$  for GABA ( $4.5 \pm 0.8$  s) increased when taurine ( $35 \pm 3.7$  s;  $n = 9$ ;  $p < 0.001$ ), homotaurine ( $20 \pm 3.5$  s;  $n = 8$ ;  $p < 0.018$ ), or hypotaurine ( $11 \pm 2.5$  s;  $n = 8$ ;  $p < 0.001$ ) was co-applied (Fig. 3g). Furthermore, the estimated  $R_sT$  for GABA ( $0.86 \pm 0.18$  s;  $n = 23$ ) increased significantly when taurine ( $2.23 \pm 0.35$  s;  $n = 9$ ;  $p = 0.002$ ), homotaurine ( $2.26 \pm 0.36$  s;  $n = 8$ ;  $p < 0.001$ ), or hypotaurine ( $3.08 \pm 0.87$  s;  $n = 8$ ;  $p = 0.001$ ) was co-applied (Fig. 3h).

### Homology modeling of the interaction of taurine and its analogs with the agonist-binding site of GABA $\rho$ 1

Dose–response curves of taurine acting on the GABA $\rho$ 1 receptor, as well as TPMPA blocking of the generated currents suggested that these molecules bind to the GABA-binding site of the receptor. Thus, to understand whether the structural analogs interact with the receptor, docking simulations were performed using a model based on acetylcholine receptor, GLIC and GluCl. Taurine bound to the GABA-binding site is shown in Fig. 4a. The electrostatic representation illustrates that the aromatic box of the receptor that permits receptor binding of GABA through a cation– $\pi$  interaction does not seem to play this role for

**Fig. 2** Taurine and its analogs act as agonists of GABA-A receptors expressed in astrocytes and in the STC-1 cell line.

**a, c** Representative currents elicited by GABA (50  $\mu$ M for astrocytes,  $n = 9$ ; 100  $\mu$ M for STC-1 cells,  $n = 6$ ), homotaurine (1 mM for astrocytes,  $n = 4$ ; 30 mM for STC-1 cells,  $n = 6$ ), hypotaurine (1 mM for astrocytes,  $n = 9$ ; 3 mM for STC-1 cells,  $n = 6$ ) and taurine [1 mM for astrocytes ( $n = 6$ ) and STC-1 cells ( $n = 6$ )]. All the currents were antagonized by the GABA-A antagonist picrotoxin (PTX, 100  $\mu$ M) in astrocytes (**A**,  $n \geq 3$ ) and in the STC-1 cell line (**c**,  $n = 6$  for all). **b, d (left)**, Summary of the current amplitudes recorded with each molecule in astrocytes and in the STC-1 cell line. Notice that GABA-A receptors expressed in each cell type responded differently to homotaurine and hypotaurine **b, d (right)**. The currents evoked by taurine and its analogs were normalized, and the antagonism by picrotoxin in astrocytes and STC-1 cells is summarized. Data are given as the mean  $\pm$  S.E.M.  $**p < 0.01$



**Table 1** Comparison between the current amplitudes generated by GABA, taurine and structural analogs in astrocytes and STC-1 cells

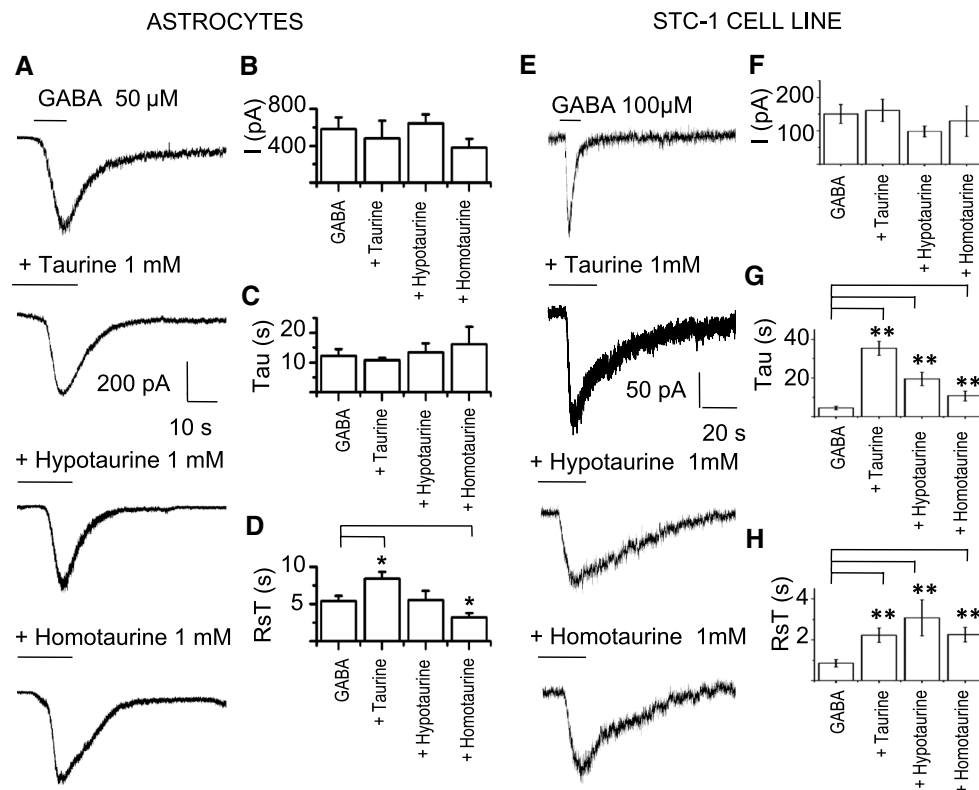
Amplitude (pA)					
Astrocyte			STC-1 cells		
Molecule		+Picrotoxin 100 $\mu$ M	Molecule		+Picrotoxin 100 $\mu$ M
GABA 50 $\mu$ M	506 $\pm$ 68 ( $n = 9$ )	70 % ( $n = 6$ )	GABA 100 $\mu$ M	150 $\pm$ 29 pA ( $n = 23$ )	100 % ( $n = 6$ )
Taurine 1 mM	193 $\pm$ 98 ( $n = 6$ )	92 % ( $n = 4$ )	Taurine 1 mM	36 $\pm$ 6 pA ( $n = 6$ )	100 % ( $n = 6$ )
Homotaurine 1 mM	891 $\pm$ 154 ( $n = 4$ )	87 % ( $n = 3$ )	Homotaurine 30 mM	93 $\pm$ 13 pA ( $n = 6$ )	100 % ( $n = 6$ )
Hypotaurine 1 mM	165 $\pm$ 68 ( $n = 9$ )	86 % ( $n = 5$ )	Hypotaurine 3 mM	196 $\pm$ 43 pA ( $n = 6$ )	100 % ( $n = 6$ )

Blockage effect of picrotoxin (100 M) is shown in the right column

taurine binding. Instead, we found the following residues to be fundamental: Q104, R125, Y219, and S263 (Fig. 4b). Hypotaurine fits the GABA-binding site; however, the docking analysis does not suggest a cation- $\pi$  interaction but rather, the generation of two polar contacts with R125 and K238 in the aromatic box B (Fig. 4b). Finally, homotaurine docking reveals only a weak interaction, a polar contact with R125 in the aromatic box C of the receptor (Fig. 4b).

## Discussion

It is now well established that GABA as well as glycine receptors can be activated by paracrine release of taurine (Le-Corronc et al. 2011). In the brain and retina, taurine accumulates in glial cells, especially in astrocytes and the Bergmann glia of the cerebellum; this amino acid is released in a non-synaptic fashion independently of calcium (Zhang and Ottersen 1992; Rodríguez-Navarro et al.



**Fig. 3** Co-application of GABA with taurine and its analogs in astrocytes and STC-1 cells. **a, e** Representative currents elicited by GABA alone (50 μM for astrocytes,  $n = 10$ ; 100 μM for STC-1,  $n = 23$ ) or co-applied with taurine [1 mM,  $n = 6$  (**a**) and  $n = 9$  (**e**), hypotaurine (1 mM;  $n = 7$  (**a**) and  $n = 8$  (**e**)) or homotaurine (1 mM,  $n = 7$  (**a**) and  $n = 8$  (**e**)), in astrocytes (**a**) or STC-1 cells (**e**), respectively. **b, f** Summary of current amplitudes showed no significant differences between GABA alone or co-applied with taurine, hypotaurine, or homotaurine for both astrocytes (**b**) and STC-1 cells (**f**). **c, g** Summary of the inactivation constant ( $\tau$ ) with GABA and GABA

co-applied with taurine, hypotaurine, or homotaurine. No significant differences were observed in astrocytes, while in STC-1 cells all the molecules co-applied with GABA significantly increased  $\tau$ . **d, h** Summary of the rise time (RsT) calculated for GABA alone or co-applied with taurine, hypotaurine, or homotaurine. All the molecules significantly modified the RsT for both astrocytes (**d**) and STC-1 cells (**h**) except for hypotaurine in astrocytes, where no significant change was observed. Data are given as the mean  $\pm$  S.E.M. \* $p < 0.05$ ; \*\* $p < 0.01$

2009). GABA $\rho$ 1 subunits are both positively and negatively modulated by taurine, depending on its concentration; in addition, taurine acts as a weak agonist of the receptor when expressed in *X. laevis* oocytes (Ochoa-de la Paz et al. 2008). This study aimed to understand whether taurine executes the same actions on GABA $\rho$ 1 when this subunit is expressed endogenously in astrocytes and STC-1 cells and to establish whether structural analogs of taurine, hypo-, and homotaurine, activate GABA-A receptors. These cells exhibit GABA currents generated by three populations of ionotropic receptors: (1) GABA-A, made up of  $\alpha$ ,  $\beta$ ,  $\gamma$  and/or  $\delta$  subunits, (2) GABA $\rho$  subunits, and (3) heteromeric receptors composed of GABA $\rho$  and other GABA-A subunits. Thus, activation and modulation of taurine and structural analogs are expected to occur on all these GABA receptors.

Partial agonism of the taurine analogs is consistent with the docking simulation of their interaction with

the agonist-binding site of GABA $\rho$ 1. In both cases, the cation- $\pi$  interaction necessary for GABA-binding (Lummis et al. 2005; Padgett et al. 2007) is not predicted, and the interactions with the GABA-binding pocket are weak. It has been suggested that taurine can induce an allosteric modulation of GABA-A receptors by binding to the benzodiazepine site (Quinn and Miller 1992); the results obtained in our co-application assay are not yet conclusive and do not establish whether hypo- and homotaurine can also allosterically modulate GABA $\rho$ 1; however, it is clear that both molecules interact with the agonist-binding site. In any case, GABA $\rho$ 1 is not modulated by benzodiazepines (Polenzani et al. 1991; Shimada et al. 1992). On the other hand, isethionic acid, which did not induce any detectable current in oocytes that expressed GABA $\rho$ 1 (not shown), did not fit in the GABA-binding site, and docking simulations (not shown) corroborated the electrophysiological findings.





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