

REVIEW

Current approaches to enhance glutamate transporter function and expression

Andréia C. K. Fontana

*Department of Pharmacology and Physiology, Drexel University College of Medicine, Philadelphia, Pennsylvania, USA***Abstract**

L-glutamate is the predominant excitatory neurotransmitter in the CNS and has a central role in a variety of brain functions. The termination of glutamate neurotransmission by excitatory amino acid transporters (EAATs) is essential to maintain glutamate concentration low in extracellular space and avoid excitotoxicity. EAAT2/GLT-1, being the most abundant subtype of glutamate transporter in the CNS, plays a key role in regulation of glutamate transmission. Dysfunction of EAAT2 has been correlated with various pathologies such as traumatic brain injury, stroke, amyotrophic lateral sclerosis, Alzheimer's disease, among others. Therefore, activators of the function or enhancers of the expression of EAAT2/GLT-1 could serve as a potential therapy for these conditions.

Translational activators of EAAT2/GLT-1, such as ceftriaxone and LDN/OSU-0212320, have been described to have significant protective effects in animal models of amyotrophic lateral sclerosis and epilepsy. In addition, pharmacological activators of the activity of EAAT2/GLT-1 have been explored for decades and are currently emerging as promising tools for neuroprotection, having potential advantages over expression activators. This review describes the current status of the search for EAAT2/GLT-1 activators and addresses challenges and limitations that this approach might encounter.

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Glutamate is the predominant excitatory amino acid neurotransmitter in the mammalian CNS and is essential for many aspects of normal brain function including cognition, memory, learning, developmental plasticity, and long-term potentiation (McEntee & Crook 1993; Weiler *et al.* 1995; Lopez-Bayghen & Ortega 2011; Peng *et al.* 2011).

The termination of glutamate neurotransmission is achieved by rapid uptake of the released glutamate by presynaptic and astrocytic high-affinity sodium-dependent transporters (Danbolt 2001; Beart & O'Shea 2007). Five structurally distinct subtypes of glutamate transporters have been cloned from animal and human tissue. Glutamate and aspartate transporter (GLAST) or EAAT1 (Storck *et al.* 1992) is localized in astroglial cells (Lehre *et al.* 1995).

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Address correspondence and reprint requests to Dr Andréia C. K. Fontana, Drexel University College of Medicine, Department of Pharmacology and Physiology, NCB, MS 488, 245 N. 15th Street, Philadelphia, PA 19012, USA. E-mail: andreia.mortensen@drexelmed.edu

Abbreviations used: 6-OHDA, 6-hydroxydopamine; AAV, adeno-associated viral; AD, Alzheimer's disease; Akt, protein kinase B; ALS, amyotrophic lateral sclerosis; A β , amyloid β ; A β PP, amyloid- β precursor protein; CREB, cAMP response element-binding protein; EAAC1, excitatory amino acid carrier 1; EAAT1-5, human excitatory amino acid transporter subtypes 1-5; EAAT2, human glutamate transporter 2; EAATs, excitatory amino acid transporters; GABA, gamma-aminobutyric acid; GLAST, glutamate and aspartate transporter; GLT-1, rat glutamate transporter 1; GltPh, glutamate transporter homolog from *Pyrococcus horikoshii*; GSK3 β , glycogen synthase kinase 3 beta; HAND, HIV-associated neurocognitive disorder; HD, Huntington's disease; HEK, human embryonic kidney; HP, hairpin loop; HSB, hybrid structure-based; JAK2, Janus-activated kinase-2; L-DOPA, L-3,4-dihydroxyphenylalanine; MCAO, middle cerebral artery occlusion; MS-153, [R]-[-]-5-methyl-1-nicotinoyl-2-pyrazoline; mTLE, mesial temporal lobe epilepsy; NF- κ B, nuclear factor kappa-light-chain-enhancer of activated B cells; NMDA, N-methyl-D-aspartate; NMDAR, N-methyl-D-aspartate receptor; N-myc, N-myc proto-oncogene protein; OGD, oxygen and glucose deprivation; PACAP, pituitary adenylate cyclase-activating peptide; PDC, L-trans-pyrrolidine-2,4-dicarboxylate; PD, human idiopathic Parkinson's disease; PI3K, phosphatidylinositol 3-kinase; PKC, protein kinase C; PS1, presenilin-1; SOD1, superoxide dismutase 1; TBI, traumatic brain injury; TBOA, threo-beta-benzoyloxyaspartate; TM, transmembrane; tPA, tissue plasminogen activator; WT, wild type.

GLT-1 or EAAT2 (Pines *et al.* 1992) is also primarily expressed in astroglial cells (Lehre *et al.* 1995), but at lower levels also in several neuronal populations (Berger & Hediger 1998; Chen *et al.* 2004; Furness *et al.* 2008; Petr *et al.* 2015). Excitatory amino acid carrier 1 or EAAT3 (Kanai & Hediger 1992), was initially localized to neuronal cell bodies and dendrites avoiding both axon-terminals and astrocytes (Rothstein *et al.* 1994). This finding has been somewhat controversial, but has recently been shown to be correct (Holmseth *et al.* 2012) and in agreement with a cell culture study suggesting that the C-terminus contains a sorting motif directing it to dendrites rather than terminals (Cheng *et al.* 2002). EAAT4 (Fairman *et al.* 1995) is a glutamate transporter subtype that was demonstrated to be predominantly expressed in cerebellar Purkinje cells by traditional antibody-based experiments (Dehnes *et al.* 1998) and by the use of bacterial artificial chromosome-transgenic mice (Gincel *et al.* 2007). EAAT5 is a transporter subtype that is expressed in vertebrate retina (Arriza *et al.* 1997). Interestingly, a report has shown that EAAT5 transporter-associated anion current hyperpolarizes the presynaptic terminal and thereby inhibits synaptic transmission by suppressing transmitter release. This demonstrates that EAAT5 does not seem to be important for removing glutamate, but rather acts like an inhibitory receptor, while EAAT1 in Müller glial cells mediates the bulk of glutamate uptake (Veruki *et al.* 2006).

Other transporters of glutamate include the glutamate–cysteine exchangers, present in neurons and glia (Sato *et al.* 2002; Sato *et al.* 2005; De Bundel *et al.* 2011) and the vesicular glutamate transporters (vGLUTs), a class of intracellular transporters that is only present in neurons (Li *et al.* 2013).

The glutamate transporter subtype EAAT2/GLT-1 (human/rat homolog, nomenclature that will be used interchangeably in this manuscript) is expressed throughout the brain, in the spinal cord, primarily in astrocytes, and also in neurons and oligodendrocytes. GLT-1 contributes 95% of the total glutamate transport activity and 1% of total brain protein in the CNS (Haugeto *et al.* 1996; Tanaka *et al.* 1997; Lehre & Danbolt 1998; Suchak *et al.* 2003; Scofield & Kalivas 2014). This important observation has been confirmed by several electrophysiology studies (Bergles & Jahr 1997; Diamond & Jahr 1997; Otis & Kavanaugh 2000; Herman & Jahr 2007). Hence, EAAT2 plays a central role in maintenance of extracellular glutamate homeostasis (Maragakis *et al.* 2004; Lauriat & McInnes 2007; Sheldon & Robinson 2007; Kim *et al.* 2011).

Molecular functioning and mechanisms of glutamate transporters

The transport of glutamate across the plasma membrane is coupled to the movement of cations (Na^+ , K^+ , and H^+)

(Kanner 1983; Kanner & Schuldiner 1987; Kanner 1989), that are necessary for glutamate uptake and transporter cycling as well as anions that are uncoupled from the flux of glutamate (Szatkowski & Attwell 1994; Zerangue & Kavanaugh 1996). Electrophysiological studies in *Xenopus* oocytes have suggested that glutamate diffusion and binding to transporters, rather than uptake, are likely to dominate the synaptic concentration decay kinetics (Wadiche *et al.* 1995a; Wadiche *et al.* 1995b). Later studies aimed to define the kinetic relationship between the two major access states that alternatively expose glutamate-binding sites to the extracellular or to the intracellular solution have proposed that transporter-associated anion currents were approximately twice as slow to rise and decay as stoichiometric transport currents, but the presence of permeant anions did not slow transporter cycling (Bergles *et al.* 2002). It is important to state that EAATs are not only secondary-active glutamate transporters, but also anion-selective channels and thus represent prototypical dual function membrane transport proteins. Recent studies combining molecular dynamics simulations and fluorescence spectroscopy of GltPh and patch-clamp recordings of mammalian EAATs have been able to determine that the transporters conduct anions through a lateral movement of the glutamate transport domain during intermediate transporter conformations that result in the formation of an anion-selective conduction pathway (Jensen *et al.* 2015; Machtens *et al.* 2015).

Additionally, voltage-clamped fluorometry assays measuring conformational changes in EAAT3 have shown that there are significant Na^+ -dependent conformational changes preceding glutamate binding, and that Na^+ and H^+ are cotransported with glutamate in the forward part of the transport cycle. The data further suggest that an increase in proton concentrations slows the reverse transport of glutamate, which may play a neuroprotective role during ischemia (Larsson *et al.* 2004). Recently, a reconstituted system and simulations suggest that the relative rates of net uptake and heteroexchange are comparable in EAAT2 (Zhou *et al.* 2014b).

Importantly, the identification of several crystal structures of a bacterial homolog of glutamate transporters GltPh has greatly facilitated our understanding of mechanistic aspects of the transport cycle (Yernool *et al.* 2004; Boudker *et al.* 2007; Reyes *et al.* 2009; Verdon & Boudker 2012; Reyes *et al.* 2013), setting the stage for structure-based drug discovery. GltPh consists of three identical protomers that associate within the membrane to form a bowl-like structure (Gether *et al.* 2006). Each protomer has eight transmembranes (TMs) and two helix-turn-helix motifs termed hairpin 1 (HP1) and HP2. Specifically, HP2 serves as the extracellular gate of the transporter (Qu & Kanner 2008). The individual protomers in the trimeric complex are held together by a central core; this core is made of TMs 1, 2, 4, and 5 from each protomer that will associate together,

while TMs 3, 6, 7, and 8, along with HP1 and HP2, form a transport domain (Vandenberg & Ryan 2013). Fluorescence spectroscopy studies have demonstrated that substrate binding to the transporter is coupled to the binding of all three Na^+ in a highly cooperative manner at both outward-facing and inward-facing conformations (Reyes *et al.* 2013). It has been suggested that transport domains translocate substrates by moving across the membrane within a central trimerization scaffold in an 'elevator-like' transport domain motion. Recently it was shown that two 'humanizing' mutations introduced in GltPh result in markedly increased transport domain dynamics, suggesting that these mutations favor structurally 'unlocked' intermediate states in the transport cycle at the interface between the transport domain and the trimeric scaffold (Akyuz *et al.* 2015).

Glutamate excitotoxicity and pathologies

Glutamate excitotoxicity is the pathological process of failure of the proper removal of glutamate by astrocytes from the synapses. This results in sustained elevation of extracellular glutamate levels and excessive activation of post-synaptic glutamate receptors resulting in Ca^{2+} influx (Nilsson *et al.* 1990a) and activation of a cascade of phospholipases, endonucleases, and proteases such as calpain that can lead to apoptotic or necrotic cell death (Raghupathi 2004). In fact, ischemic events in humans and in animals lead to an acute and sustained increase in extracellular glutamate concentrations (Benveniste *et al.* 1984; Faden *et al.* 1989; Brown *et al.* 1998; Vespa *et al.* 1998), indicating a lack of timely clearance by glutamate transporters. In excitotoxic states, the extracellular concentrations of glutamate may reach a millimolar range, causing degeneration of neurons through excessive stimulation of glutamate receptors (Meldrum & Garthwaite 1990; Clements *et al.* 1992; Rosenberg *et al.* 1992; Zhou & Danbolt 2014).

The speed of clearance of transmitter from the cleft influences many aspects of synaptic function, including the time course of the post-synaptic response and the peak post-synaptic receptor occupancy. Estimates of the time course of transmitter clearance, either by detailed theoretical modeling, or from the attenuation of synaptic transmission produced by a low-affinity competitive antagonist, have produced results that are in agreement, suggesting an average concentration of transmitter peaks in the range 1–5 mM, and a biphasic clearance, with time constants of approximately 100 μs and 2 ms (Clements 1996).

A study investigated the synaptic glutamate concentration using NMDARs (NMDA receptors) expressed by CA1 pyramidal cells in acute hippocampal slices, and found that its baseline concentration is near 25 nM (Herman & Jahr 2007). Such low resting concentrations will only be possible if the expression levels of GLT-1 are extremely high, as discussed in (Zhou & Danbolt 2013), therefore supporting

the notion that expression levels of GLT-1 are high, and ambient glutamate is, in physiological conditions, very low.

Furthermore, it is important to discuss how EAAT2 modulation could possibly affect the process of transmitter diffusion. Studies have demonstrated that rapid communication in the brain between glia and neurons relies on the release and diffusion of transmitter molecules across the synaptic cleft (Rusakov *et al.* 2011). By focusing on the interplay between extracellular and intramembrane diffusion processes, a recent study illustrates the remarkable versatility of signal formation in synapses. An integrative measure of three-dimensional glial coverage confirms that thin spine post-synaptic densities are more tightly surrounded by glia. This distinction suggests that diffusion-dependent synapse–glia communication and astrocytic glutamate uptake near 'learning' synapses (associated with thin spines) could be stronger than that near 'memory' synapses (associated with larger spines) (Medvedev *et al.* 2014).

Recent evidence shows that EAAT2 on rat astrocytes appears to be concentrated near synapses, where it acts acutely in a very dynamic manner to shape synaptic transmission by binding released glutamate and, through a process that involved movements of the transporters in the cell membrane. This movement was dependent on neuronal and glial activities and transporter concentration was strongly reduced in the vicinity of active glutamatergic synapses. Notably, glutamate uncaging at synaptic sites increased GLT-1 diffusion, displacing transporters away from this compartment and impairment of GLT-1 membrane diffusion through cross-linking slowed the kinetics of excitatory post-synaptic currents, resulting in a prolonged time course of synaptic glutamate. These data provide evidence for a physiological and highly dynamic role of GLT-1 in shaping synaptic transmission through surface diffusion (Edwards 2015; Murphy-Royal *et al.* 2015).

Therefore, one might suggest that the tight regulation of the glutamate signal by GLT-1 is of great importance for normal glutamate neurotransmission and when diminished in injured states activators of glutamate transport can prevent aberrant glutamate signaling.

Mechanisms involved in glutamate-mediated excitotoxicity include down regulation of glutamate transporters or glutamate efflux via transporter reversal, as further discussed below. See Fig. 1 for a simplified depiction of the role of EAAT2 at glutamatergic synapses at physiological (Fig. 1a) and pathological (excitotoxicity, Fig. 1b) states.

Glutamate transporter down-regulation

Reduced expression and function of EAAT2 has been reported in numerous neurological disorders, as will be further discussed below. The first report was on cerebral ischemia (Levy *et al.* 1995), with many other examples including amyotrophic lateral sclerosis (ALS) (Rothstein *et al.* 1995; Trotti *et al.* 2001; Wilson *et al.* 2003) and

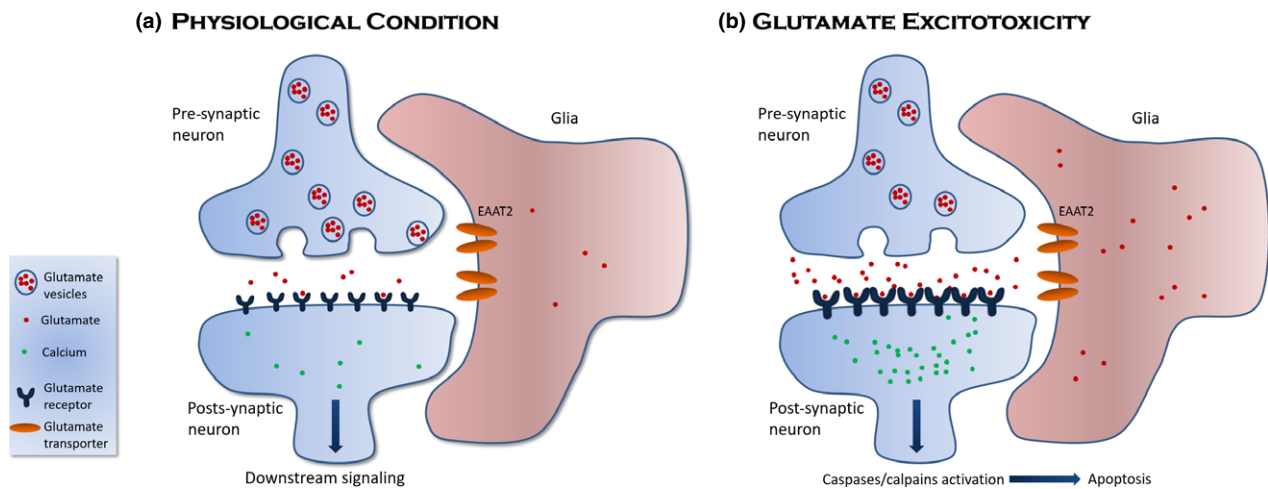


Fig. 1 Glutamatergic signaling under physiological (a) and excitotoxic (b) states. L-glutamate is stored in synaptic vesicles at presynaptic terminals. Glutamate (red dots) is released into the synaptic cleft, activating post-synaptic glutamate receptors. EAAT2 takes up released glutamate, removing it from the synaptic cleft to maintain the extracellular glutamate concentration below neurotoxic levels. In

excitotoxicity glutamate is released in higher concentration and glutamate clearance is decreased, causing excessive activation of glutamate receptors resulting in Ca^{2+} influx (green dots) and activation of a cascade of phospholipases, endonucleases, and proteases such as calpain that can lead to apoptotic or necrotic cell death.

Alzheimer's disease (AD) (Li *et al.* 1997) among others. Additionally, decreased GLT-1 expression was shown in an animal model of brain trauma (Rao *et al.* 1998) and hippocampal neuronal damage was exacerbated after brain injury in an antisense knockdown model of GLT-1 (Rao *et al.* 2001). These results are consistent with the decreased EAAT2 activity observed in patients of traumatic brain injury (TBI) and stroke (Ikematsu *et al.* 2002; van Landeghem *et al.* 2006; Yi & Hazell 2006).

Moreover, a study on antisense knockdown revealed that loss of glutamate transporters GLAST or GLT-1 produced elevated extracellular glutamate levels, neurodegeneration characteristic of excitotoxicity, and a progressive paralysis, while loss of the neuronal glutamate transporter excitatory amino acid carrier 1 did not elevate extracellular glutamate in the striatum but did produce mild neurotoxicity and resulted in epilepsy (Rothstein *et al.* 1996).

Furthermore, a very important work by (Tanaka *et al.* 1997) has shown that knockout of the EAAT2 gene resulted in exacerbated damage, with lethal spontaneous seizures and increased susceptibility to acute cortical injury, compared to their wild-type counterparts, following cerebral injury in mice. This was the first time a GLT-1 knockout mice model was described. Later studies confirmed severe disturbances in mice lacking GLT-1 glutamate transporter (Hakuba *et al.* 2000; Katagiri *et al.* 2001; Mitani & Tanaka 2003). Another study showed that the brains of mice lacking GLAST/GLT-1 developed normally but that GLAST/GLT-1 double-knockout mice died around embryonic days 17–18 and exhibited cortical, hippocampal, olfactory bulb disorganization, and impairment of several essential aspects of neuronal

development (Matsugami *et al.* 2006). Recently, a study with conditional knockout mice confirmed that GLT-1 plays main roles in the brain (Zhou *et al.* 2014a).

On the other hand, a study on GLT-1-deficient mice, GLT-1 homozygous ($-/-$), and heterozygous ($+/-$), has shown that a 50% reduction in GLT-1 levels had minor effects (Kiryk *et al.* 2008). The GLT-1 ($-/-$) mice had lower body and brain weight, mild neuronal loss in CA1 hippocampal region, focal gliosis, severe focal neuronal paucity in layer II of the neocortex, and short life-span. In contrast, GLT-1 ($+/-$) mice exhibited a 59% decrease in GLT-1 immunoreactivity in their brain tissue, with surprisingly no apparent morphological brain abnormalities, and their life-span and behavior not markedly different from controls, suggesting that GLT-1 ($+/-$) mice may serve as a potentially useful model to study neurodegenerative disease conditions with mild hyperglutamatergic activity.

Although the mechanism of down-regulation of EAAT2 in these conditions/diseases has yet to be fully established, it is clear that it results in impairment of their functions, which plays an important role in the etiology of neurological diseases. Consequently, there has been an extensive effort to identify molecular targets for enhancement of EAAT2 expression as a potential therapeutic approach.

Glutamate transporter reversal

As explained above, glutamate translocation is a multi-step process that utilizes energy stored in the Na^+/K^+ electrochemical gradient to move glutamate against its concentration. Under physiological conditions, the conventional transport direction is inward, however, in excitotoxic

conditions when the extracellular $[Na^+]/intracellular [K^+]$ ratio decrease and/or intracellular $[Na^+]/extracellular [K^+]$ ratio increase, glutamate is transported in the outward direction (Szatkowski *et al.* 1990; Billups & Attwell 1996; Jabaudon *et al.* 2000). This has been demonstrated in a study where raising extracellular potassium concentration evoked a reversed uptake current (Levy *et al.* 1998). Furthermore, massive increases in extracellular potassium and indiscriminate release of glutamate have been reported following concussive brain injury (Katayama *et al.* 1990) in addition to studies demonstrating a decreased sodium concentration after lateral fluid percussion (Soares *et al.* 1992). Also, GLT-1 reversal was shown in ischemic glutamate release (Phillis & O'Regan 1996; Grewer *et al.* 2008). Furthermore, the Na^+/K^+ -ATPase that maintains these ion gradients are compromised in the injured brain (Werner & Engelhard 2007; Lima *et al.* 2008). Additionally, GLT-1 can be rendered inactive due to oxidative stress, as demonstrated by reduction in GLT-1/EAAT2 function in an ALS-causing superoxide dismutase 1 (SOD1) mutation in mice and rats, in both *in vitro* and *in vivo* studies (Volterra *et al.* 1994; Trotti *et al.* 1999a).

In this context, it remains to be clarified whether EAAT2 activators will facilitate glutamate clearance under excitotoxic conditions, or will rather intensify reverse transport. A study showed that *in vitro* ischemia led to neuronal damage, however, the ischemia-induced reversal of GLT-1 contributed to the survival of astrocytes themselves, demonstrating that reversal of glutamate transporter activity, while damaging to neurons, is also important for the survival of astrocytes under conditions of ischemia and might contribute to later survival of neurons (Kosugi & Kawahara 2006). Also, several strategies involving the application of pharmacological agents such as ceftriaxone, riluzole, and others, as reviewed in (Krzyzanowska *et al.* 2014) have shown significant protection of nervous tissues against ischemia.

Collectively, these are indications that modulation of glutamate transporters as a therapeutic concept is relevant. However, this concept faces the same inherent obstacle as other drugs acting through glutamate-mediated mechanisms, such as a significant risk of inducing adverse effects, as glutamatergic signaling is involved in several functions including brain development, cell survival, and synapsis differentiation.

Next, we will briefly discuss the current understanding of the glutamate-mediated excitotoxicity and the role of GLT-1 in several acute and chronic CNS pathologies. For extensive reviews on glutamate excitotoxicity and pathologies, see (Rothstein 1995b; Mark *et al.* 2001; Hazell 2007; Kim *et al.* 2011; Mehta *et al.* 2013).

Stroke

The third leading cause of death in industrialized countries and the most frequent cause of permanent disability in adults

worldwide (Beresford *et al.* 2003), is primarily either hemorrhagic or ischemic, with almost 80% of the stroke being ischemic (Banerjee *et al.* 2011). Glutamate released during ischemia was shown to be responsible for cell death in models of retinal ischemia (Sisk & Kuwabara 1985; Louzada-Junior *et al.* 1992) and of cerebral ischemia. A pivotal study on transient cerebral ischemia monitored by intracerebral microdialysis revealed elevation of the extracellular concentrations of glutamate and aspartate in rat hippocampus, respectively, eight- and threefold during the ischemic period. These results suggest that the large increase in the content of extracellular glutamate and aspartate in the hippocampus induced by the ischemia may be one of the causal factors in the damage to certain neurons observed after ischemia (Benveniste *et al.* 1984), a result later confirmed in other studies (Taoufik & Probert 2008; Chao *et al.* 2010). A study showed that release of glutamate occurs within minutes of ischemic onset, with a tenfold higher concentration of glutamate within the cells resulting in failure of the energy and ion gradient (Nishizawa 2001). These studies indicate that abnormal function of glutamate transport plays an essential role in the excitotoxic neurodegeneration that occurs in the models.

Interestingly, a study supports the notion that glutamate transporters work in reverse during ischemic events, thus contributing to the increased extracellular glutamate by releasing glutamate that triggers neuronal death. Hence, to use GLT-1 modulators may in such situations potentiate neuronal death and complexity of the disease (Rossi *et al.* 2000).

On the other hand, other studies suggest that modulation of EAAT2 to inhibit glutamate-induced excitotoxicity may be a potential therapeutic target for the treatment of stroke and this hypothesis has been explored in many ways. First, decreased glutamate transporter expression was observed in ischemic neuronal death, and this may have disrupted the normal clearance of the synaptically released glutamate that contributed to the death (Rao *et al.* 2001). Then, studies using pharmacological manipulation of EAAT2 levels (Chu *et al.* 2007; Lipski *et al.* 2007) and increased expression of GLT-1 by using an adeno-associated viral vector expressing GLT-1 cDNA were shown to reduce the damage caused by brain ischemia, with decrease in the size of lesion and improvement of behavioral recovery (Harvey *et al.* 2011). Additionally, ceftriaxone, an up-regulator of EAAT2 protein expression and activity, confers neuroprotection in cerebral ischemia/reperfusion injury (Verma *et al.* 2010).

Since treatments for stroke are, to date, restricted to the administration of plasminogen activator (tPA) to break up blood clots in the arteries of the brain, and no treatments available to halt the progression of secondary damages, we expect that future research may further explore the potential role of EAAT2 as target for neuroprotection after ischemic stroke.

Epilepsy

A group of disorders characterized by recurrent spontaneous seizures that apparently result from complex processes involving several neurotransmitter systems including glutamate (Jabs *et al.* 2008; Wang & Qin 2010; Werner & Covenas 2015), that produce an imbalance between neuronal excitatory–inhibitory activities that has been suspected to occur during seizures and to underlie epileptogenesis (Coutinho-Netto *et al.* 1981; Kaila *et al.* 2014).

Increased extracellular glutamate levels were found in the hippocampi of patients with mesial temporal lobe epilepsy (mTLE) (During & Spencer 1993; Cavus *et al.* 2005; van der Hel *et al.* 2005). Evidence that GLT-1 is involved in epilepsy come from several studies, however, GLT-1 levels reported in animal models and human patients are not consistent, as discussed below.

Studies with mice lacking GLT-1 revealed that they are prone to exhibiting seizures (Tanaka *et al.* 1997). On the other hand, another study with GLT-1 knockdown in rats resulted in increased extracellular glutamate, but not seizures (Rothstein *et al.* 1996). These findings raise the question of whether GLT-1 is really involved in epilepsy.

Patients with mTLE have been reported with decreased levels of EAAT1 and EAAT2 as well as increased extracellular glutamate levels in the epileptogenic hippocampus both during and after clinical seizures (Mathern *et al.* 1999; Proper *et al.* 2002). On the other hand, (Tessler *et al.* 1999) proposed that major changes in the level of expression of the glutamate transporters do not play an important role in the development of human mTLE, but may be implicated in the etiology of other types of epilepsy. In the same line, another study found no significant change in the amount of EAAT2 in the human epileptogenic hippocampus, which led them to suggest a deficiency in glutamine synthetase in astrocytes as a molecular basis for extracellular glutamate accumulation and seizure generation (Eid *et al.* 2004). Later, a study with human epilepsy patients also claimed that EAAT2 is not altered (Bjornsen *et al.* 2007).

Regarding animal models of epilepsy, several studies claim that the levels of GLT-1 are unchanged, as reported in kindled rats (Akbar *et al.* 1997; Miller *et al.* 1997; Simantov *et al.* 1999), in rats fed with an anticonvulsant ketogenic diet (Bough *et al.* 2007), and in spontaneously epileptic rats (Guo *et al.* 2010).

On the other hand, decreased levels of GLT-1 were observed in pilocarpine (Lopes *et al.* 2013) and albumin models (David *et al.* 2009), as well as in tuberous sclerosis (Wong *et al.* 2003), in the FeCl₃-induced limbic epilepsy model (Ueda *et al.* 2007), and in the chest compression-induced audiogenic epilepsy model (Lu *et al.* 2008).

Interestingly, increasing GLT-1 expression protected mice against status epilepticus-induced death, neuropathological changes, and chronic seizure development (Kong *et al.* 2012).

Recently, it was demonstrated that conditional deletion of GLT-1 protected against fatal epilepsy (Petr *et al.* 2015). This study demonstrated that astrocytic GLT-1 is indeed involved in critical functions required for normal weight gain, resistance to epilepsy, and survival.

Together, these studies indicate that astrocytic glutamate uptake plays a crucial role in protecting neurons from hyperexcitability. However, how exactly this mechanism is disturbed in epilepsy is still under discussion. Furthermore, it is still unclear whether changes in transporters represent causative or compensatory changes during epileptogenesis. Finally, on the basis of some of the previous reports, the notion that enhancing GLT-1/EAAT2 protein expression is a potential therapeutic approach to treat epilepsy cannot be ruled out.

For more information on astrocytic targets (including GLT-1) and new avenues for therapeutic treatment of epilepsy, please see the reviews (Wetherington *et al.* 2008; Coulter & Eid 2012; Crunelli *et al.* 2015).

Autism

Autism is a disorder of unknown etiology, and co-occurrence with epilepsy being common. It is thought that reduced density of cerebellar Purkinje neurons is a key neuropathological component in autism (Courchesne 1997; Palmen *et al.* 2004), which results in excitotoxicity, mitochondrial dysfunction, and degeneration (Khodorov 2000). A recent report by the Autism Genome Project Consortium identified a new linkage peak for autism in the region of chromosome 11 where the gene for EAAT2 is located (Szatmari *et al.* 2007). Recent considerations revealed signs of astroglial, oligodendroglial, and microglial dysfunction in the autistic brain, suggesting that all these cellular processes may represent presumptive targets for novel therapeutic strategies (Zeidan-Chulia *et al.* 2014). Recently, a study investigating the effects of ceftriaxone and cefixime, activators of GLT-1, demonstrated that these drugs improved some symptoms of autism and decreased epilepsy seizures (Ghanizadeh & Berk 2015).

Traumatic brain injury

TBI is a complex pathology of many etiologies (Matute *et al.* 2006; Guerriero *et al.* 2015). TBI has been studied with multiple different animal models, including non-penetrating and penetrating impact, lesion studies, fluid percussion, shock waves (explosion), among others. For reviews on animal models of TBI, see (Finnie 2001; de Lanerolle *et al.* 2015).

Studies on TBI in humans and in animals have demonstrated an acute increase in tissue glutamate concentrations that remain high for up to 5 days in humans, suggesting a lack of timely clearance by glutamate transporters (Nilsson *et al.* 1990b; Baker *et al.* 1993; Palmer *et al.* 1993; Brown *et al.* 1998; Vespa *et al.* 1998; Yamamoto *et al.* 1999). Clinical and experimental studies have determined that the resulting glutamate-mediated excitotoxicity is a significant

contributor to acute post-injury neurodegenerative events (Faden *et al.* 1989; Yi & Hazell 2006). The failure of proper glutamate removal results in sustained elevation of extracellular glutamate levels and overstimulation of the NMDA glutamate receptor that results in overloading of neurons with Ca^{2+} and Na^+ (Rothman & Olney 1995), leading to the activation of phospholipases, endonucleases, and proteases such as calpain resulting in the apoptotic and/or necrotic cell death (reviewed in (Raghupathi 2004).

Previous studies in animal models of lateral fluid percussion injury reported decreased GLT-1 expression between 6 and 72 h in the rat (Rao *et al.* 1998), other studies have shown differences in EAAT2 expression in the cerebral cortex relative to the survival time and severity of cerebral contusion after TBI in humans (Ikematsu *et al.* 2002; van Landeghem *et al.* 2006). In addition, antisense knockdown of GLT-1 in rat exacerbates neuronal damage following controlled cortical impact (Rao *et al.* 2001). Collectively, these studies suggest that targeting GLT-1 could be a strategy to find a therapy for the treatment of TBI.

Importantly, the pathophysiology of TBI differs between lesional and perilesional tissue and the degree of injury severity; cell injury proceeds more slowly in the brain tissue that is immediately proximal to the most severe injury and may be reversible in this penumbra. Similarly, following concussion the majority of brain injury is reversible, given the typical course of symptom resolution (Collins *et al.* 2006; Zlotnik *et al.* 2012; Eisenberg *et al.* 2013; Eisenberg *et al.* 2014; Burda *et al.* 2015). Hence, in brain areas with reversible damage, the opportunity to decrease acute glutamate excitotoxicity holds promise for neuroprotection, while other pharmacological approaches might be needed for other types of TBI.

Amyotrophic lateral sclerosis

ALS is a debilitating disease characterized by progressive loss of voluntary motor neurons leading to muscle atrophy, weight loss, and respiratory failure. The pathogenesis of ALS involves inflammation, oxidant stress, apoptosis, mitochondrial dysfunction, SOD1 protein aggregation, and astroglial dysfunction, including dramatic loss of EAAT2/GLT-1 in both in the motor cortex and in the spinal cord (Rothstein *et al.* 1992; Rothstein *et al.* 1995; Rothstein 1995a; Wilson *et al.* 2003). Additionally, mouse models of ALS including lines G85R, A4V, and I113T SOD1-mutants all have a marked loss or inactivation of glutamate transporters (Bruijn *et al.* 1997; Trotti *et al.* 1999b; Bendotti *et al.* 2001). Moreover, selective loss of EAAT2 has also been demonstrated in both sporadic and familial cases of ALS (Van Den Bosch *et al.* 2006).

Although several lines of evidence point to the possibility that glutamate transport defects and the related excitotoxicity may play a role in the late stage of ALS disease progression, there are no consistent data to support defects in EAAT2 as

having a primary role in the induction or the selectivity of motor neuron degeneration taking place in ALS. In this sense, a study showed that GLT-1 over-expression in SOD1G93A mouse cervical spinal cord did not protect motor neurons, preserve diaphragm function, or prolong animal survival, suggesting that focal restoration of GLT-1 expression in cervical spinal cord astrocytes was not an effective therapy for ALS (Li *et al.* 2015). Moreover, it is possible that multiple therapies aimed at simultaneously diminishing the overwhelming Ca^{2+} overload, endoplasmic reticulum stress, and mitochondrial dysfunctional pathways would be effective on halting ALS progression, as proposed by (Verche *et al.* 2011).

Extensive studies using ceftriaxone and other GLT-1 expression activators have been performed with hopes that it would restore GLT-1 expression to normal levels and slow, or even possibly, stop ALS progression in patients. These studies will be further discussed below.

Huntington's disease

Huntington's disease (HD) is a devastating progressive autosomal dominant neurodegenerative disorder in which multiple areas of the brain degenerate, mainly involving the dopamine, glutamate, and GABA. It is caused by a mutated form of the huntingtin gene, where excessive CAG repeats result in accumulation of the mutant protein as cytoplasmic and nuclear aggregate inclusions (Bates 2003; Walker 2007), resulting in oxidative stress and mitochondrial dysfunction (Browne & Beal 2006).

Moreover, excitotoxicity is thought to be important in the pathogenesis of HD (Massieu & Garcia 1998; Shin *et al.* 2005; Estrada-Sanchez & Rebec 2012). In this regard, it has also been shown that the EAAT2 mRNAs of post-mortem brains of HD patients are decreased in correlation to disease severity (Arzberger *et al.* 1997; Faideau *et al.* 2010).

Additionally, studies have shown that increase in the functional expression of GLT-1 can improve the behavioral phenotype of the mouse model of HD (Estrada-Sanchez *et al.* 2009; Miller *et al.* 2012a; Petr *et al.* 2013). Furthermore, ascorbate release and GLT-1 function are impaired in the striatum of transgenic mouse models of HD (Rebec 2013). These studies suggest that changes in GLT-1/EAAT-2 expression or function can potentiate or ameliorate the progression of HD, however, effective therapies are yet to be discovered and developed. However, in a R6/2 transgenic mouse model that is heterozygous for the null allele of GLT-1/EAAT-2 and carries the double mutation, weight loss, accelerating rotarod, climbing, and paw-clasping were not exacerbated, suggesting that decreased expression of GLT-1 in this model does not worsen disease progression.

Neuropathic pain

Neurons responsible for sensing noxious stimuli and conducting pain signals from periphery to the spinal cord are

predominantly glutamatergic (Tao *et al.* 2005). Decreases in glutamate uptake activity and expression of spinal glutamate transporters have been reported in animal models of pathological pain, with pharmacological inhibition or antisense down-regulation of spinal transporters being reported to induce/aggravate pain behaviors (Gegelashvili & Bjerrum 2014).

On the other hand, over-expression of GLT-1 in astrocytes in the spinal cord attenuated the induction, but not maintenance, of inflammatory and neuropathic pain, suggesting that up-regulation of spinal GLT-1 could be a novel strategy for the prevention of pathological pain (Maeda *et al.* 2008).

Additionally, emerging data point at key roles of glutamate transporters in molecular mechanisms of chronic pain and analgesia, including the development of opioid tolerance. However, precise pharmacological targeting of the glutamate transport system requires detailed elucidation of molecular factors and signaling pathways underlying expression and activity of individual glutamate transporters subtypes, including their splice variants, and molecular behavior on the pathological states of neuropathic pain (reviewed in (Gegelashvili & Bjerrum 2014).

HIV-associated neurocognitive disorder

A common neuropathology found in the brains of HIV infected individuals is astrocytes apoptosis. Excess glutamate is released upon HIV infection of macrophage/microglial cells, which has been associated with neurotoxicity mediated by gp120, transactivator of transcription and other HIV proteins (Sabri *et al.* 2003).

To prevent glutamate excitotoxicity in HIV-associated neurocognitive disorder (HAND), antagonists of NMDA receptors have been explored, however, this approach has failed as a result of side effects, and therefore alternatives are being sought. Another approach has been to regulate the glutamate transporters, and there is preliminary experimental evidence that these approaches have potential therapeutic utility for the treatment of HAND. These efforts, however, are at an early stage where the next steps are dependent on the identification of drug-like inhibitors as well as the development of predictive neuroAIDS animal models (Potter *et al.* 2013). Recently, a study showed that methamphetamine and HIV treatment activated trace amine associated receptor 1, leading to intracellular cAMP in human astrocytes and decreased EAAT-2 mRNA, significantly decreasing glutamate clearance. Furthermore, molecular alterations in astrocyte TAAR1 levels correspond to changes in astrocyte EAAT2 levels and function, confirming the involvement of EAAT2 in the context of HAND (Cisneros & Ghorpade 2014).

However, our precise understanding of host factors that mediate the neurotoxic insult launched by exposure of CNS to HIV-1 is still developing. We need to better understand the regional genetic variations in viral genes and further identify

opportunities to specifically block the neurotoxicity of viral and host proteins as well as identify potential neuroprotective factors that could be augmented to help alleviate the negative impact of HIV-1 on the brain (Rao *et al.* 2014).

Alzheimer's disease

AD is a progressive age-related neurodegenerative disorder. The pathophysiological characteristic of AD is abnormal deposition of fibrillar amyloid β ($A\beta$) protein, intracellular neurofibrillary tangles, oxidative damage, and neuronal death (Kumar *et al.* 2015). Aberrant glutamate stimulation and resulting synaptic dysfunction have been proposed as one of several mechanisms by which synapses are damaged in AD (Mattson & Chan 2003; Hynd *et al.* 2004; Talantova *et al.* 2013).

Moreover, a number of studies have found that GLT-1/EAAT-2 is significantly reduced or damaged in AD (Hardy *et al.* 1987; Li *et al.* 1997; Lauderback *et al.* 2001; Woltjer *et al.* 2010; Scott *et al.* 2011). Additionally, a report showed that deficits in GLT-1 function compound the effects of familial AD amyloid- β precursor protein/presenilin-1 mutant transgenes in younger animals, suggesting that this dysfunction may contribute to early occurring pathogenic processes associated with AD (Mookherjee *et al.* 2011), suggesting that GLT-1/EAAT-2 loss is capable of driving cognitive impairment in the context of $A\beta$ -related neuropathology (Schallier *et al.* 2011). Also, a study showed that $A\beta_{1-42}$ induces rapid GLT-1/EAAT-2 mislocalization and internalization in astrocytes, which leads to a marked reduction in the rate of glutamate clearance from the extracellular space (Scimemi *et al.* 2013).

Recently, a study revealed that GLT-1 loss also caused an apparent compensatory increase in insulin-degrading enzyme activity in the liver, an organ that has been shown to regulate peripheral $a\beta$ levels and expresses GLT-1, suggesting that partial GLT-1 loss can cause insulin/Akt (protein kinase B) signaling abnormalities observed in AD (Meeker *et al.* 2015).

Nonetheless, it is not yet entirely clear whether GLT-1/EAAT-2 dysfunction plays a pathogenic role in AD, and whether $A\beta$ -related pathogenic processes have a functional impact on the rate at which astrocytes remove endogenous, synaptically released glutamate is unknown (Scimemi *et al.* 2013). Furthermore, a study suggested that GLT-1 is not involved in AD, as the authors reported decreased expression of glutamine synthetase but no changes in GLT-1 expression in the prefrontal cortex in a mouse model of AD (Kulijewicz-Nawrot *et al.* 2013). Therefore, the question whether GLT-1/EAAT-2 may be used as target for neuroprotective functions in AD pathogenesis remains to be further explored.

Currently available treatments for AD include acetylcholinesterase inhibitors (Hogan 2014) and NMDA receptor antagonist memantine (Zimmer *et al.* 2012; Kumar *et al.* 2015), however, they provide symptomatic relief and target

late aspects of the disease, and therefore a definite treatment for this disease is yet to be identified.

Human idiopathic Parkinson's disease

Human idiopathic Parkinson's disease (PD) is a progressive neurodegenerative movement disorder that is primarily characterized by degeneration of the dopaminergic neurons at the nigrostriatal pathway (Deumens *et al.* 2002). This degeneration is believed to lead to an overactivation of the subthalamic nucleus (Rodriguez *et al.* 1998), increasing the firing rate of the glutamatergic excitatory projections to the substantia nigra (Bamford *et al.* 2004; Ambrosi *et al.* 2014) causing a sustained exposure to glutamate that could accelerate the degeneration of dopaminergic neurons (Oster *et al.* 2014). Therefore, glutamate-mediated excitotoxicity may be involved in a lethal vicious cycle, which critically contributes to the exacerbation of nigrostriatal degeneration in PD.

The role of glutamate transporters in the pathogenesis of PD is not entirely clear, however, some studies clearly demonstrate a link between disturbed glutamatergic neurotransmission and glutamate transporter functioning in the striatum of an unilateral 6-hydroxydopamine animal model for PD (Massie *et al.* 2010). Another study using the same animal model showed a time-dependent bilateral effect of lesioning on the expression and the activity of GLT-1 (Carbone *et al.* 2012b). Additionally, acutely induced dysfunction of EAATs in the rat by single unilateral injection of their substrate inhibitor L-trans-pyrrolidine-2,4-dicarboxylate (PDC), triggers a neurodegenerative process mimicking several PD features (Assous *et al.* 2014). Moreover, it was reported that ceftriaxone, a drug that increases expression of GLT-1, ameliorated locomotor impairments in this animal model of PD (Chotibut *et al.* 2014; Kelsey & Neville 2014).

The major treatment for PD is L-3,4-dihydroxyphenylalanine (L-DOPA), to replace the progressive loss of dopamine; however repeated L-DOPA treatment leads to motor complications, such as dyskinesias associated with increased extracellular glutamate levels in the basal ganglia (Robelet *et al.* 2004), therefore alternative treatments are needed. Based on previous studies and data, GLT-1/EAAT-2 may be a potential target for PD (Sheldon & Robinson 2007).

EAAT2 as target for neuroprotection

Several classes of compounds that target glutamate-mediated excitotoxicity, such as NMDAR antagonists (McIntosh *et al.* 1990; Baker *et al.* 1993; Okiyama *et al.* 1997; Okiyama *et al.* 1998; Ikonomidou & Turski 2002; Ginsberg 2008) and compounds that target calcium influx (Okiyama *et al.* 1992; Cheney *et al.* 2000) have been shown to alleviate cellular damage and neurologic deficits. However, they were shown to exhibit substantial side effects, making them less attractive for translation to the clinic.

Interestingly, a substantial proportion of extrasynaptic NMDARs are located adjacent to glia, which express EAAT2 (Hardingham & Bading 2010). EAAT2 might be a more attractive therapeutic target since it is positioned to take up glutamate near neurotoxic extrasynaptic NMDARs.

Historically, modulation of that GLT-1/EAAT2 expression and/or activity has received surprisingly little attention as putative drug targets, considering the immense therapeutic potential in pharmacological intervention into glutamatergic neurotransmission and the fact that other neurotransmitter transporters are targeted by clinically administered drugs against epilepsy, depression, and attention-deficit hyperactivity disorder (Kristensen *et al.* 2011).

Nonetheless, finding alternative strategies to treat excitotoxic conditions without producing considerable side effects has become a very significant endeavor in the last decade. In this way, pharmacological manipulation of transporter expression or function is of high interest from a therapeutic point of view, and specially EAAT2, accounting for most of the glutamate clearance in the CNS, has become a high priority target for the development of therapeutic agents to prevent excitotoxicity (Bunch *et al.* 2009).

Albeit this review is focused on positive modulators of GLT-1 activity/expression, it is important to reflect that the drug development using EAAT2 as target started with the discovery of pharmacological agents that were developed to study the intrinsic properties and function of the EAATs, as reviewed in (Bridges *et al.* 1999). In this regard, a very important and pivotal series of medicinal chemistry work is illustrated by the development of the potent EAAT2 inhibitor TBOA (threo-beta-benzyloxyaspartate) and analogs (Lebrun *et al.* 1997; Shimamoto *et al.* 1998; Shigeri *et al.* 2004; Shimamoto 2008). These series of work revolutionized the field, as TBOA was the first non-transportable blocker for all subtypes of EAATs ever identified. Indeed, blockade of EAAT2 by TBOA results in marked neurotoxicity through the activation of NMDARs because of an increased extracellular glutamate level (Izumi *et al.* 2002; O'Shea *et al.* 2002; Bonde *et al.* 2005). Additionally, this discovery helped elucidate several functions of the EAATs; and it was also an encouraging step in the search for other modulators of the function of EAATs, as seen subsequently with the identification of positive ones, as will be discussed below.

Furthermore, it is important to consider developmental changes of expression when designing therapies based on GLT-1 activators. It has been demonstrated that the development of the total glutamate uptake activity in the forebrain corresponds to that of GLT-1, in agreement with the observation that GLT-1 is the predominant transporter in the adult brain (Ullensvang *et al.* 1997). Moreover, it has been shown that GLT-1 was present in fetal brain and spinal cord, with expression progressively increasing to adult levels throughout the neuraxis by postnatal day 26 (Furuta *et al.*

1997). Another study suggests a postnatal development- and age-dependent differential interaction of transcription factors NF- κ B (nuclear factor kappa-light-chain-enhancer of activated B cells) and N-myc (N-myc proto-oncogene protein) to their respective sequences and they act as positive and negative regulator, respectively, of GLT-1 gene expression in the brain during early developmental period in both cerebral and cerebellar cortices, which might be different in aging mice (Gupta & Prasad 2014).

In this regard, a very well-studied pathology that illustrates striking differences according to the stage of development is TBI, as observed by the differences between adult and pediatric TBI (Giza *et al.* 2007). Important distinctions of the younger brain after TBI include an increased propensity for apoptosis, age-dependent parameters for cerebral blood flow and metabolism, development-specific biomarkers, increased likelihood of early post-traumatic seizures, differential sensitivity to commonly used neuroactive medications, and anesthesia and altered neuroplasticity during recovery from injury (Huh & Raghupathi 2009). Questions that remain to be answered, among others, include whether, following pediatric injury, there is persistent decrease in GLT-1 and other proteins, leading to neuronal hyperexcitability, whether loss of these proteins is a direct result of epigenetic modulation of gene transcription, whether manipulation of DNA methylation reverses these losses, and, finally, whether therapies based on GLT-1 enhancement would be beneficial.

Many studies suggest that targeting GLT-1/EAAT2 provides a novel approach for the treatment of conditions that involve glutamate excitotoxicity. This review attempts to review the current state of research on the development of compounds with the ability to increase levels of EAAT2 protein expression (summarized in Table 1) or compounds that interact directly or indirectly with EAAT2 enhancing its catalytic activity (summarized in Table 2).

Transcription/translational modulators

GLT-1 expression can be regulated in several ways, at both the transcriptional and translational level (Anderson & Swanson 2000; Sheldon & Robinson 2007). Even though GLT-1 protein is already highly expressed, several studies show evidence that GLT-1 can be up-regulated. In this manuscript, we will focus on approaches demonstrated to increase glutamate transporter expression with a therapeutic goal, and we will present the development of these regulators in a chronological sequence below and in Table 1 [with regulators classified as proteins (a), endogenous molecules (b) or synthetic molecules (c)]. For a review on the mechanisms of regulation of glutamate transporter function discussing transcriptional regulation and post-transcriptional regulation and modifications, see (Grewer *et al.* 2014).

Early studies showed increased expression of GLT-1 protein in astroglial cultures by the epidermal growth factor

and this was thought to be mediated through intracellular signaling involving cAMP as dibutyryl-cAMP incubation had a similar effect (Zelenaia *et al.* 2000). Pituitary adenylate cyclase-activating peptide was also found to stimulate GLT-1 expression (Figiel & Engele 2000).

Dexamethasone, a glucocorticoid, increases EAAT2 mRNA levels and subsequently up-regulates EAAT2 protein expression and activity (Wen *et al.* 2005; Zschocke *et al.* 2005). A recent study shows that dexamethasone induces neuroprotection in hypoxic-ischemic brain injury in newborn rats, partly mediated via Akt activation (Feng *et al.* 2014b).

Importantly, studies concluded that Akt induces the expression of GLT-1 through increased transcription and that Akt can regulate GLT-1 expression without increasing GLAST expression in astrocytes (Li *et al.* 2006).

Corticosterone and retinol are both able to increase the translation of EAAT2 transcripts and, moreover, many disease-associated insults affected the efficiency of the induced translation of the EAAT2 transcript by these compounds (Tian *et al.* 2007).

GPI-1046, a synthetic neuroimmunophilin ligand that selectively increases GLT-1 expression and activity, was shown to protect motor neurons against excitotoxicity *in vitro* and increases the survival rate of transgenic ALS mice (Ganel *et al.* 2006) and more recently was shown to have a role in reduction of ethanol consumption (Sari & Sreemantula 2012).

The antiepileptic drug sodium valproate (valproic acid) was shown to up-regulate hippocampal glutamate transport following chronic treatment (Hassel *et al.* 2001). Because the mechanism of action of valproate involves modulation of multiple mechanisms its beneficial effects cannot be contributed to only EAAT2 regulations. These other mechanisms include the regulation of GABA and glutamate neurotransmissions, activation of pro-survival protein kinases, and inhibition of histone deacetylase. However, evidence for its neuroprotective properties is emerging. Recently, it was suggested that valproate reduces the development of chronic pain after nerve injury, in part by preventing down-regulation of GLT-1 (Yoshizumi *et al.* 2013) and to prevent retinal degeneration in a murine model of normal tension glaucoma (Kimura *et al.* 2015).

Minocycline, a broad-spectrum tetracycline antibiotic, was shown in neuropathic rats to maintain the expression of GLT-1 in the spinal dorsal horn and this was suggested to be involved in the attenuation of behavioral hypersensitivity (Nie *et al.* 2010). This study supports the idea that GLT-1 may be a potential target for the development of analgesics.

The tricyclic antidepressant amitriptyline was shown to reverse the down-regulation of EAAT1 and EAAT2 and to induce up-regulation of EAATs in an animal model of spared nerve injured (Mao & Yang 2010), suggesting that this up-regulation may be one of the therapeutic mechanisms of amitriptyline in the treatment of neuropathic pain.

Table 1 Neuroprotective activators of EAAT2 expression

Compound/protein	Mechanisms	Key findings	References
EGF ^b , dibutyryl-cAMP ^b , PACAP ^b	Endogenous hormones or second messengers, inducers of GLT-1 expression	Early studies showed increased expression of GLT-1 protein by EGF, cAMP, and PACAP, setting the stage for studies aimed to further understand the regulation of GLT-1	Figiel & Engele (2000) and Zelenia <i>et al.</i> (2000)
Dexamethasone ^c	Synthetic glucocorticoid, inducer of EAAT2 protein expression	Neuroprotection in hypoxic-ischemic newborn rat brain injury	Wen <i>et al.</i> (2005), Zschocke <i>et al.</i> (2005), and Feng <i>et al.</i> (2014b)
Akt ^a	Protein kinase B, inducer of GLT-1 expression	Basic research that identified Akt as a control signal for GLT-1 expression <i>in vivo</i>	Li <i>et al.</i> (2006)
Corticosterone ^b and retinol ^b	Steroid hormone and metabolite of vitamin A, modulators of EAAT2 mRNA translation	Identification of glucocorticoids as EAAT2 protein expression inducers	Tian <i>et al.</i> (2007)
GPI-1046 ^c	Neuroimmunophilin, inducer of GLT-1 expression	Attenuates ethanol intake in part through GLT-1 up-regulation	Ganel <i>et al.</i> (2006) and Sari & Sreemantula (2012)
Valproate ^c	Multiple mechanisms through regulation of GABA and glutamate neurotransmissions, activation of prosurvival protein kinases, inhibition of histone deacetylase	Reduces chronic pain after nerve injury and prevents retinal degeneration in glaucoma model	Hassel <i>et al.</i> (2001), Yoshizumi <i>et al.</i> (2013), and Kimura <i>et al.</i> (2015)
Minocycline ^c	Broad-spectrum tetracycline antibiotic, ameliorates down-regulation of GLT-1 expression in neuropathy model	Attenuates behavioral hypersensitivity in neuropathic rats	Nie <i>et al.</i> (2010)
Amitriptyline ^c	Tricyclic antidepressant, inducer of EAAT2 expression	Attenuates mechanical allodynia, used in the treatment of neuropathic pain	Mao & Yang (2010)
mTOR ^a	Serine-threonine protein kinase mammalian target of rapamycin, induces EAAT2 over-expression	Up-regulation of GLT-1 via mTOR- NF- κ B signaling cascade in OGD may promote glutamate uptake in brain ischemia and neurodegenerative diseases	Wu <i>et al.</i> (2010), Pignataro <i>et al.</i> (2011), and Ji <i>et al.</i> (2013)
Pyridazine derivatives ^c	Small-molecule activators of EAAT2 expression	A cell-based assay identified 61 compounds as modulators of EAAT2 translation, thiopyridazine derivatives further explored.	Colton <i>et al.</i> (2010) and Xing <i>et al.</i> (2011)
JAK2 ^a	Non-receptor tyrosine kinase, inducer of EAAT2 expression	JAK2 is proposed to participate in protection against excitotoxicity	Hosseinzadeh <i>et al.</i> (2011)
Tamoxifen ^c	Antagonist of the estrogen receptor, increases TGF- α and GLT-1 expression	Regulates GLT-1 via the CREB and NF- κ B pathways	Karki <i>et al.</i> (2013) and Lee <i>et al.</i> (2013)
Estradiol ^b	Estrogen hormone, enhancer of GLT-1 function	Neuroprotective in clinical and experimental models of neuronal injury.	Karki <i>et al.</i> (2014)
GSK3 β ^a	Kinase, stimulates the expression of GLT-1	Constitutively active GSK3 β is important in controlling the expression of functional glutamate transporters on the plasma membrane.	Jimenez <i>et al.</i> (2014)
Histamine ^b	Neuromodulator, selectively increases GLT-1 expression	Neuroprotective after OGD and MCAO	Fang <i>et al.</i> (2014)

(continued)

Table 1. (continued)

Compound/protein	Mechanisms	Key findings	References
Harmine ^c	Beta-carboline alkaloid, increases GLT-1 expression	Neuroprotective in animal of ALS and cerebral ischemia	Li <i>et al.</i> (2011) and Sun <i>et al.</i> (2014)
Sulbactam ^c	β -lactamase inhibitor, GLT-1 expression up-regulator	Protects pyramidal neurons against brain ischemia.	Cui <i>et al.</i> (2015)
Ceftriaxone ^c	Transcriptional activator of GLT-1 expression	Neuroprotective in several animal models of brain trauma and neurodegeneration, failed clinical trials on ALS patients	Rothstein <i>et al.</i> (2005), Beghi <i>et al.</i> (2006), Chu <i>et al.</i> (2007), Lipski <i>et al.</i> (2007), Hota <i>et al.</i> (2008), Lee <i>et al.</i> (2008), Melzer <i>et al.</i> (2008), Miller <i>et al.</i> (2008), Thone-Reineke <i>et al.</i> (2008), Verma <i>et al.</i> (2010), Pan <i>et al.</i> (2011), Leung <i>et al.</i> (2012), Wei <i>et al.</i> (2012), Berry <i>et al.</i> (2013), Cudkowicz <i>et al.</i> (2013), Rebec (2013), Cui <i>et al.</i> (2014), Feng <i>et al.</i> (2014a), Hu <i>et al.</i> (2015), and Kelsey & Neville (2014)
LDN/OSU-0212320 ^c	Pyridazine derivative, translational activator of GLT-1 expression	Protects cultured neurons from excitotoxicity and delays motor function decline and extend life span in animal model of ALS	Kong <i>et al.</i> (2014)

This class of regulators can be divided into three major subcategories: ^aproteins, ^bendogenous molecules, and ^csynthetic molecules.

Table 2 Neuroprotective activators of EAAT2 activity

Compound	Mechanisms	Key findings	References
Riluzole	Sodium channel blocker, activator of glutamate uptake	Neuroprotective <i>in vitro</i> , approved by FDA to ALS patients, efficacious in preclinical models of spinal cord injury	McIntosh <i>et al.</i> (1996), Zhang <i>et al.</i> (1998), Miller <i>et al.</i> (2000), Frizzo <i>et al.</i> (2004), Traynor <i>et al.</i> (2006), Dagci <i>et al.</i> (2007), Fumagalli <i>et al.</i> (2008), Banas <i>et al.</i> (2010), Carbone <i>et al.</i> (2012a), Miller <i>et al.</i> (2012b), Mu <i>et al.</i> (2000), and Wilson & Fehlings (2014)
Guanosine	Guanine-based purine, activator of glutamate uptake	Currently examined for treatment of ischemic brain injury	Frizzo <i>et al.</i> (2001), Frizzo <i>et al.</i> (2005), Moretto <i>et al.</i> (2005), and Hansel <i>et al.</i> (2014)
Nicergoline	Ergot derivative activator of glutamate uptake	Currently examined for cerebrovascular disease and dementia	(Fioravanti & Flicker (2001), Nishida <i>et al.</i> (2004), and Fioravanti <i>et al.</i> (2014)
MS-153	R]-[-]-5-methyl-1-nicotinoyl-2-pyrazoline, activator of GLT-1 activity, inhibitor of glutamate release and calcium channels	Neuroprotective in MCAO, focal ischemia and lateral fluid percussion brain injury in rats	(Umemura <i>et al.</i> (1996), Kawazura <i>et al.</i> (1997), Shimada <i>et al.</i> (1999), Uenishi <i>et al.</i> (1999), Nakagawa <i>et al.</i> (2001), Abekawa <i>et al.</i> (2002b), Li <i>et al.</i> (2004), Nakagawa <i>et al.</i> 2005, Abekawa <i>et al.</i> (2002a), Alhaddad <i>et al.</i> (2014), and Fontana <i>et al.</i> (2015)
Parawixin1	Specific activator of EAAT2 activity	Neuroprotective in animal model of retinal ischemia	Fontana <i>et al.</i> (2003) and Fontana <i>et al.</i> (2007)
GT949 and GT951	Direct activators of EAAT2 activity, no effect on NMDA-mediated currents	Ongoing research	Unpublished

mTOR (a family of serine-threonine protein kinase mammalian target of rapamycin) is involved in the control of a wide variety of cellular processes, and there is evidence for the involvement of PI3K(phosphatidylinositol 3-kinase)/Akt/

mTOR signaling regulation of GLT-1 in astrocytes (Pignataro *et al.* 2011). A study showed that treatment with PI3K or Akt inhibitors suppresses the phosphorylation of Akt and mTOR and decreased GLT-1 up-regulation in astrocytes.

Moreover, treatment with the mTOR inhibitor rapamycin decreased GLT-1 protein and mRNA levels and did not affect Akt phosphorylation, suggesting that mTOR is a downstream target of the PI3K/Akt pathway regulating GLT-1 expression (Wu *et al.* 2010). Another study demonstrated up-regulation of GLT-1 via mTOR-Akt- NF- κ B (nuclear factor- κ B) signaling cascade in a model of oxygen and glucose deprivation, suggesting that this signaling cascade may work to promote glutamate uptake in brain ischemia and neurodegenerative diseases (Ji *et al.* 2013).

However, even though mTOR inhibitors, such as rapamycin and its analogs may represent novel, rational therapies for a variety of neurological disorders, they have not been tested yet in patients with epilepsy, Alzheimer's, Huntington's, and Parkinson's disease. With continuing progress in basic and clinical research, there are reasons to be optimistic that the clinical indications of mTOR inhibitors for neurological disease will continue to expand in future (Wong 2013).

An approach to identify translational activators using a cell-based enzyme-linked immunosorbent assay identified 61 compounds that showed a dose-dependent increase in EAAT2 protein levels (Colton *et al.* 2010). In addition, the same group developed thiopyridazine and pyridazine derivatives that increase EAAT2 expression, although its neuroprotective properties remain to be evaluated (Xing *et al.* 2011).

Targeting the Janus-activated kinase-2 (JAK2) has also been demonstrated to be a potential therapeutic avenue for regulating glutamate transporter function as it was described to be a powerful regulator of glutamate transporters, by showing that co-expression of JAK2 with EAAT2 in *Xenopus* oocytes increased glutamate-induced current by 67%, and thus JAK2 was proposed to participate in the protection against excitotoxicity (Hosseinzadeh *et al.* 2011). However, thus far EAAT2 has not been validated as target for JAK2 and its activation has not been investigated in animal models of excitotoxicity.

Tamoxifen, a selective estrogen receptor modulator that is used to treat breast cancer, was shown to enhance the expression and function of GLT-1/EAAT2 in rat astrocytes (Lee *et al.* 2013). Tamoxifen regulates GLT-1/EAAT-2 via the cAMP response element-binding protein and NF- κ B pathways (Karki *et al.* 2013).

Estradiol (17 β -estradiol), one of the most active estrogen hormones possessing neuroprotective effects in both *in vivo* and *in vitro* models, has been shown to increase expression of both GLAST and GLT-1 mRNA and protein and glutamate uptake in astrocytes (Lee *et al.* 2013), an effect that seems to be mediated by growth factors such as transforming growth factor- α . However, the adverse effects associated with long-term use of estradiol have hampered its clinical utility (Karki *et al.* 2014).

Another promising therapeutic target for regulating GLT-1 expression is GSK3 β (glycogen synthase kinase 3 beta), a

kinase involved in multiple cellular processes including neuronal development and synaptic plasticity, that was found through both co-expression and pharmacological studies to stimulate the expression of GLT-1 (while, interestingly, reduced that of GLAST) in COS-7 cells and *Xenopus laevis* oocytes. The authors suggest that this differential modulation may be particularly relevant in pathological conditions such as AD or ischemia in which glutamate transporters and GSK3 β are involved (Jimenez *et al.* 2014). Recent evidence suggests that GSK3 β regulation of GLT-1 expression in the spinal dorsal horn plays a role with neuropathic pain (Weng *et al.* 2014).

A recent study showed that histamine increased GLT-1 expression in pure cultured astrocytes, decreased the extracellular glutamate content and alleviated neuronal cell death induced by exogenous glutamate. Importantly, histamine was neuroprotective in brain slices after oxygen and glucose deprivation and against ischemic injury in animal model of middle cerebral artery occlusion (Fang *et al.* 2014).

Harmine, a beta-carboline alkaloid, was identified through a screening of a library of 1040 FDA-approved compounds and natural products in fetal-derived human-immortalized astroglial cells stably expressing a firefly luciferase reporter under the control of EAAT2 promoter. Harmine effectively increased GLT-1 expression and transporter activity *in vivo* and presented neuroprotection effects in a rat model of ALS, with the beneficial effects specifically because of up-regulation of GLT-1 (Li *et al.* 2011). A recent study shows that harmine also provides neuroprotection in a global cerebral ischemia model (Sun *et al.* 2014).

Treatment with sulbactam, a β -lactamase inhibitor, was shown to protect pyramidal neurons against brain ischemia. A study showed that both antisense knockdown of GLT-1 expression or inhibition of the GLT-1 uptake activity with dihydrokainate, a selective inhibitor of GLT-1, significantly blocked the neuronal protective effect of sulbactam, indicating that sulbactam has a neuronal protective effect though up-regulation of GLT-1 (Cui *et al.* 2015).

Although β -lactams have been historically used as antimicrobials, a notable ancillary effect in the host was identified by (Rothstein *et al.* 2005), with a blind screen of over 1000 FDA-approved drugs that discovered ceftriaxone enhances the expression of EAAT2. This work was the foundation for the search of therapies using ceftriaxone as up-regulator of GLT-1 expression, and currently there are over 100 reports of the use of ceftriaxone as a neuroprotective drug that increases GLT-1 in many animal models of excitotoxic diseases.

Some of these studies include: ceftriaxone treatment delayed neuronal death and muscle strength loss and increases survival in a mouse model of ALS (Melzer *et al.* 2008). Ceftriaxone was also reported to reduce neurodegeneration and motor deficits in rodent models of stroke (Lipski *et al.* 2007; Thone-Reineke *et al.* 2008) and to attenuate the

damage observed in models of both acute and chronic neurodegenerative disorders (Hota *et al.* 2008; Lee *et al.* 2008, Verma *et al.* 2010). In addition, ceftriaxone treatment before transient focal ischemia up-regulates GLT-1 mRNA and protein and reduces infarct volume in rats (Chu *et al.* 2007). Ceftriaxone was shown to have antinociceptive role in chronic neuropathic pain in rats (Hu *et al.* 2010). Ceftriaxone was shown to reduce the level of brain glutamate, brain edema, and neuronal death following lateral cortical impact injury in rats (Pan *et al.* 2011; Wei *et al.* 2012; Cui *et al.* 2014). There is also evidence that ceftriaxone, by removing glutamate from the synapse more efficiently through up-regulation of GLT-1, resulted in a decrease in post-traumatic seizures after TBI (Goodrich *et al.* 2013). Additionally, in the 6-hydroxydopamine-lesioned model of PD, ceftriaxone has been shown to increase striatal GLT-1 expression and attenuate the normally observed motor symptoms (Leung *et al.* 2012). In the same model, the therapeutic effects of ceftriaxone were equal to L-DOPA, the current gold standard treatment for PD; however, unlike L-DOPA, ceftriaxone did not result in dyskinesia as a side effect (Kelsey & Neville 2014). Additionally, ceftriaxone was shown to have neuroprotective effects in HD (Miller *et al.* 2008; Rebec 2013). Ceftriaxone was also reported to alleviate early brain injury after subarachnoid hemorrhage by increasing EAAT2 expression via the PI3K/Akt/NF-kappaB (Feng *et al.* 2014a). Recently ceftriaxone was shown to protect neurons against global brain ischemia via up-regulation of GLT-1 expression and glutamate uptake (Hu *et al.* 2015) and to preserve glutamate transporters and prevent intermittent hypoxia-induced vulnerability to hypoxia in an *in vitro* model of chronic intermittent hypoxia (Jagadapillai *et al.* 2014).

Regarding translational studies, an open trial of 108 ALS patients showed that ceftriaxone did not substantially improve muscle strength and disability scores (Beghi *et al.* 2006). Ceftriaxone has failed to show significant efficacy in clinical trials for stroke (Nederkoorn *et al.* 2011). Nevertheless, a clinical trial of ceftriaxone treatment for ALS patients was conducted, which reported to be a safe and tolerable drug for humans, in stages 1 and 2 (Berry *et al.* 2013). Unfortunately, stage 3 was discontinued because of lack of efficacy in increasing length of patient survival or preventing a decline in function (Cudkowicz *et al.* 2013).

Recently, the effects of β -lactam antibiotics ampicillin, cefazolin, and cefoperazone were investigated in a model of ethanol intake in alcohol-preferring rats. Chronic ethanol consumption is known to down-regulate expression of GLT-1, which increases extracellular glutamate levels in subregions of the mesocorticolimbic reward pathway. The compounds significantly reduced ethanol intake and significantly up-regulated both GLT-1 and pAKT expressions in the nucleus accumbens and prefrontal cortex, demonstrating that these compounds appear to be potential therapeutic com-

pounds for treating alcohol abuse and/or dependence (Rao *et al.* 2015).

In contrast to studies demonstrating positive effects of beta-lactams on glutamate expression, other studies have shown inconsistent results. A study found that ceftriaxone did not increase GLT-1 promoter activity in rat hippocampal slices, although a neuroprotective effect was shown, suggesting that the neuroprotection was not associated to the level of GLT-1 protein expression (Lipski *et al.* 2007). Another study showed that chronic administration of ceftriaxone in rats failed to increase GLT-1 mRNA and protein levels in different brain regions, however, intriguingly, it increased GLT-1 activity and conferred neuroprotection in a stroke model (Thone-Reineke *et al.* 2008). Additionally, no effect of ceftriaxone has been detected in various CNS regions in WT C57BL/6 mice, including the spinal cord, however, the drug attenuated disease course and severity in a model of autoimmune CNS inflammation (Melzer *et al.* 2008). Moreover, ceftriaxone slightly reduced GLT-1 levels and significantly decreased glutamate uptake activity in mouse striatal astrocytes (Carbone *et al.* 2012a). Thus, it seems that some yet unidentified factors characteristic to different model systems can prevent the beta-lactam-dependent up-regulation of GLT-1. Therefore, ceftriaxone may not always exhibit antinociceptive or neuroprotective efficacies.

There are many possible explanations for these discrepant results on GLT-1 regulation by beta-lactam antibiotics. First, there seems to be indications that the protective effects of beta-lactam antibiotics are highly dependent on the experimental model, as suggested by the studies presented above. Second, the many ways that GLT-1 expression is measured can have different results. One way is indirect, by measuring survival or amelioration of behavior defects; however, this is not a reliable approach as the drugs might have indirect effects on other targets. Other approaches include to measure transporter expression using immunoblots or measuring mRNA levels, or to measure uptake activity in synaptosome assays. There is some skepticism that these measures are comparable, this is illustrated by one example and challenging idea that synaptosomal assays primarily detect the 5–10% of GLT-1 present in neurons rather than the 90–95% present in astrocytes (Furness *et al.* 2008). If this is correct, one might expect that the uptake activity in synaptosomes could go up dramatically with minor changes to the total brain activity of GLT-1. This could have an impact on the interpretation of a number of studies, and this could also explain why some studies on EAAT2 activators demonstrate lack of up-regulation of expression, despite the fact that neuroprotection is observed. Third, the preservation of tissue used in the assays should be considered, e.g., from human autopsy versus freshly obtained from animals, as these approaches can yield significant different levels of transport (Li *et al.* 2012).

Further studies on ceftriaxone, other beta-lactam antibiotics and derivatives are necessary to determine whether an

analog compound with good brain penetrance, pharmacokinetic properties, and with less risk of side effects and undesired toxicity can be developed into a therapeutic drug.

Recently, a pyridazine derivative (LDN/OSU-0212320) that increases EAAT2 expression through translational activation was shown to protect cultured neurons from glutamate-mediated excitotoxic injury and to delay motor function decline and extend lifespan in an animal model of ALS (Kong *et al.* 2014).

Final remarks on EAAT2 expression enhancers

Reported transcriptional activators of GLT-1, such as ceftriaxone, have been tested in many disease models and are capable of providing neuronal protection. This is also illustrated the newly identified LDN-212320, that was developed to identify more GLT-1/EAAT-2 modulators with improved pharmacological and bioactivity properties.

Moreover, transgenic approaches for EAAT2 over-expression confirm its neuroprotective effects, as shown in a study with double transgenic mice created from crossing the ALS mouse model to a mouse model over-expressing EAAT2, that resulted in an animal that displays delayed grip strength decline and motor neuron loss and increased life expectancy (Guo *et al.* 2003). In addition, HEK (Human Embryonic Kidney) cells engineered to over-express EAAT2 protect cultured motor neurons exposed to glutamate toxicity (Wisman *et al.* 2003).

However, transition from preclinical animal studies to human clinical trials has been challenging, as successful preclinical studies often fail in subsequent clinical trials. Furthermore, increasing the *in vivo* expression levels of EAAT2 could have profound impact on glutamatergic neurotransmission throughout the brain with the intrinsic risk of inducing adverse effects (as exemplified in (Matos-Ocasio *et al.* 2014). Another potential issue is that many of the mechanisms targeted underlie the expression of numerous other genes, and thus transcriptional/translational modulators could potentially exert off-target effects outside the glutamatergic system. Interestingly, the kinetics and duration of the modulation exerted by at least some of these modulators are likely to be significantly different from the properties of those targeting the EAAT protein, which could constitute an advantage or a problem depending on the specific use. Additional issues for EAAT2 translation activators lay in the effect that, although having neuroprotective properties, this class of compounds must be administered prophylactically to be neuroprotective; therefore they have low clinical relevance for acute conditions.

Compounds that increase transporter activity

It has been known for many years that the activity of transporters can be regulated by many means, such as phosphorylation, sulfhydryl oxidation, arachidonic acid,

among others (Conradt & Stoffel 1997; Gegelashvili *et al.* 2000).

On the other hand, positive allosteric modulation of GLT-1/EAAT2 with pharmacological compounds represents a relatively novel approach to the treatment of conditions/diseases involving glutamate excitotoxicity. Table 2 summarizes known compounds that interact directly or indirectly with EAAT2 augmenting its catalytic activity.

Riluzole, a highly studied synthetic compound modulator of EAAT2, is a sodium channel-blocking benzothiazole anticonvulsant drug that was demonstrated to decrease excitotoxicity and neurodegeneration through many different mechanisms, including increased glutamate uptake (McIntosh *et al.* 1996; Zhang *et al.* 1998; Azbill *et al.* 2000; Mu *et al.* 2000; Brothers *et al.* 2013). Riluzole significantly increased GLT-1-mediated uptake in transfected HEK293 (Fumagalli *et al.* 2008), rat cortical astrocyte cultures (Frizzo *et al.* 2004), and within striatal astrocytes (Carbone *et al.* 2012a).

Moreover, riluzole was shown to have neuroprotective effects on rat glial cells subjected to glutamate excitotoxicity (Dagci *et al.* 2007), to attenuate the behavioral effects of chronic unpredictable stress, a rodent model of depression (Banasr *et al.* 2010) and to delay the onset of muscle weakness and extends life span by 4–6 weeks in animal models of ALS (Gurney *et al.* 1996; Waibel *et al.* 2004; Del Signore *et al.* 2009).

Currently riluzole is approved by the U.S. Food and Drug Administration for the treatment of ALS (Traynor *et al.* 2006), albeit it prolongs patient life by only a few months (Miller *et al.* 2012b). Additionally, riluzole showed efficacy in preclinical models of spinal cord injury in reducing the extent of sodium and glutamate-mediated secondary injury (Wilson & Fehlings 2014).

Guanosine is a guanine-based purine that activates glutamate uptake and exerts neurotrophic and neuroprotective effects in several models of CNS trauma (Frizzo *et al.* 2001; Frizzo *et al.* 2005), including a model of neonatal hypoxia-ischemia (Moretto *et al.* 2005), with recent evidence suggesting that guanosine is a promising therapeutic agent for the treatment of ischemic brain injury (Hansel *et al.* 2014).

Nicergoline, an ergot derivative marketed under the trade name Sermion for the treatment of cognitive, affective, and behavioral disorders of older people (Fioravanti & Flicker 2001) has been shown to have neuroprotective actions through increase of glutamate transport in rat cortical synaptosomes and cloned glutamate transporters (Nishida *et al.* 2004). This compound was recently subject of a systematic review and meta-analysis that suggested a good safety profile for cerebrovascular disease and dementia (Fioravanti *et al.* 2014).

Additionally, MS-153 ([R]-[S]-5-methyl-1-nicotinoyl-2-pyrazoline), characterized to be a glutamate uptake enhancer acting through GLT-1 (Shimada *et al.* 1999), was first

demonstrated to inhibit elevated brain glutamate levels induced by rat middle cerebral artery occlusion (Umemura *et al.* 1996), then later characterized as neuroprotective in a model of focal ischemia in rats (Kawazura *et al.* 1997). Further studies showed that MS-153 inhibited high voltage-gated calcium channels through interactions with protein kinase C, preventing massive release of glutamate from nerve terminals in ischemic conditions (Uenishi *et al.* 1999). MS-153 has also been shown to affect other conditions that are believed to involve excessive glutamate signaling, including the inhibition of morphine tolerance and dependence (Nakagawa *et al.* 2001), the attenuation of behavioral sensitization development to phencyclidine (Abekawa *et al.* 2002a; Abekawa *et al.* 2002b), an anxiolytic effect in fear conditioning (Li *et al.* 2004), and attenuation of the conditioned rewarding effects of morphine, methamphetamine and cocaine (Nakagawa *et al.* 2005). MS-153 was also recently suggested to have potential as a therapeutic drug for the treatment of alcohol dependence (Alhaddad *et al.* 2014).

Our laboratory has studied the actions of MS-153 in the lateral fluid percussion model of TBI. We found that administration of MS-153 in the acute post-traumatic period provides acute and long-term neuroprotection for TBI, and that the neuroprotective actions of MS-153 can be explained on the basis of inhibition of calcium channels and interactions with PKC (protein Kinase C), rather than activation of glutamate transport. We conclude that MS-153 displays promising potential for developing a much needed therapy for treatment of TBI (Fontana *et al.* 2015).

Our approach to identify compounds that modulate glutamate transport was through the characterization of spider venoms, a valuable source of compounds bioactive on synaptic transmission (Fontana *et al.* 2003; de O Beleboni *et al.* 2004; Estrada *et al.* 2007; Fontana *et al.* 2007). Our group has purified a compound from *Parawixia bistriata* spider venom, referred to as Parawixin1, that was shown to stimulate glutamate uptake in rat synaptosomes and to protect retinal tissue from ischemic damage (Fontana *et al.* 2003) and further characterized to act directly and selectively on the glutamate transporter EAAT2, specifically by facilitating conformational transitions involved in substrate translocation (Fontana *et al.* 2007).

This spider venom provides proof of principle for the development of drugs that act through a novel mechanism and also as a useful tool to further understand the mechanisms that may be involved in augmentation of the transporter function.

The exogenous EAAT2 activators compounds reported herein have, in their majority, non-specific effects that are associated with side effects that limit their utility or exhibit rather promiscuous pharmacological profiles (Table 2). With a goal to identify direct, selective, and potent EAAT2 activators, our laboratory has performed mutagenesis studies that identified a structural region within EAAT2 that is

important for the transporter-enhancing activity, comprised of TM domains 2, 5, and 8 (Mortensen *et al.* 2015). Subsequently, this unique structural information was employed in hybrid structure-based virtual screening of a large library to identify novel EAAT2 activating compounds. This approach successfully identified hit compounds (GT949 and GT951) that enhance the activity of EAAT2 in cultured cells and that have neuroprotective properties in glia-neuronal cultures subjected to glutamate excitotoxicity (Fontana, A.C.K. *et al.*, unpublished data). Further characterization of these novel allosteric modulators of EAAT2 using medicinal chemistry approaches and *in vivo* pharmacokinetic advanced profiling is currently being performed.

Concluding remarks and prospects

L-glutamate-mediated excitotoxicity is involved in a wide range of acute and chronic pathologies. Over the last decade the scientific community has concluded that drugs targeting glutamate receptors are limited and inefficient, as well as other drugs designed to target the secondary damages of glutamate-mediated excitotoxicity, due to unwanted side effects. Currently, there are no safe and effective drugs for the prevention or treatment of glutamate-mediated excitotoxicity diseases/conditions, thus, there is a need for better therapeutics.

The concept of pursuing GLT-1/EAAT-2 to prevent excitotoxicity has made great progress in recent years; it is now accepted that GLT-1/EAAT-2 transporters are a major target to combat neurotoxicity and provide novel potential therapeutic opportunities for the treatment of neurological diseases. However, the true potential of EAAT2 as target for therapeutics still remains to be understood, appreciated, and explored. Also, as reviewed in this article, the molecular behavior of glutamate transporters in excitotoxicity models is more complex than just general down-regulation or reversal of GLT-1/EAAT2. There is still much to further understand regarding when and how to manipulate the glutamate transport system as a target for excitotoxicity therapies, and also to further elucidate the intracellular signaling pathways that accompany the changes on disease states.

Nevertheless, small molecules designed to up-regulate the activity of GLT-1/EAAT2 are still being developed, and these efforts may reveal intriguing and promising therapeutic avenues. Moreover, efforts toward identification of direct EAAT2 allosteric activators seem to have gained more attention recently as an approach to target these proteins. Additionally, positive EAAT2 allosteric modulators can be administered acutely and be advantageous over the expression enhancers that generally require to be administered prophylactically. It remains to be determined whether small molecules allosteric activators of EAAT2, as for many biologics, lend themselves to conventional pharmacokinetic analysis and are efficacious in animal models of CNS injury.

Perhaps the future will reveal that such a class of compounds is effective in chronic conditions and in combination therapies.

We expect to see, in the upcoming years, the identification of an arsenal of selective pharmacological compounds that target these fascinating transporters and the investigation of their translational possibilities.

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