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Blood-Brain Barrier Dysfunction, TGF β Signaling, and Astrocyte Dysfunction in Epilepsy

UWE HEINEMANN¹, DANIELA KAUFER², and ALON FRIEDMAN^{3,*}

¹Institute of Neurophysiology, Charité Universitätsmedizin, Berlin

²Department of Integrative Biology, Helen Wills Neuroscience Institute, UC Berkeley, Berkeley, California

³Department of Physiology and Neurobiology, Zlotowski Center for Neuroscience, Ben-Gurion University of the Negev, Beer-Sheva, Israel

Abstract

Brain insults, including traumatic and ischemic injuries, are frequently followed by acute seizures and delayed development of epilepsy. Dysfunction of the blood-brain barrier (BBB) is a hallmark of brain insults and is usually surrounding the core lesion. Recent studies from several laboratories confirmed that vascular pathology is involved in the development of epilepsy and demonstrate a key role for astroglia in this process. In this review, we focus on glia-related mechanisms linking vascular pathology, and specifically BBB dysfunction, to seizures and epilepsy. We summarize molecular and physiological experimental data demonstrating that the function of astrocytes is altered due to direct exposure to serum albumin, mediated by transforming growth factor beta signaling. We discuss the reported changes and their potential role in the observed hyperexcitability as well as potential implications of these findings for the future development of new diagnostic modalities and treatments to allow a full implementation of the gained knowledge for the benefit of patients with epilepsy.

Keywords

buffering; connexins; astroglia; potassium channels; glutamate

BLOOD-BRAIN BARRIER DYSFUNCTION, EPILEPTOGENESIS, AND SEIZURES

Focal epilepsy typically arises either within or adjacent to a cortical lesion (Willoughby, 2000). While the characteristic hypersynchronous activity within the focus has been described extensively, epileptogenesis—the process modifying a functioning neuronal network into an epileptic one is less understood. In light of the high prevalence of acquired epilepsy, the high rate of drug resistance and the consequent neurological impairments, it is crucial to understand the epileptogenesis process, identify patients at risk, and develop antiepileptogenic strategies.

Long-lasting, persistent focal epilepsy can be elicited by inducing developmental cortical malformations (Jacobs et al., 1999; Wong, 2009), repeated electrical stimulation such as in

the kindling model (McNamara, 1988) or repeated application of ictogenic agents, such as penicillin (Opdam et al., 2002; Prince and Wilder, 1967) or pentetrazole (Barkai et al., 1990), or by chronic injury (Halpern, 1972; Pitkänen and McIntosh, 2006; Prince and Tseng, 1993). The most common animal models for temporal lobe epilepsy (TLE) involves induction of status-epilepticus (SE) by pilocarpine or kainic acid (Turski et al., 1987) reviewed by (Curia et al., 2008), or by repetitive electrical stimulation (Lothman et al., 1987; for review see Löscher, 2002). In these animal models (similar to patients), a latent period of days to weeks precedes the development of epileptic seizures (Hoffman et al., 1994; Prince and Tseng, 1993). In animal models, the reorganization in nervous tissue during this period ultimately leads to appearance of spontaneous and recurrent seizures. Hence, this period is referred to as epileptogenesis. In clinical practice, this is often not the case and reorganization following a lesion or SE maintains a level of excitability within affected networks which prevent appearance of seizures. This requires studies of mechanisms which shift the balance in reorganizational processes from a condition in which no seizures occur into a condition where every so often seizures emerge. We postulate that dysfunctional blood-brain barrier (BBB) might underlie such a shift in reorganizational balance towards epileptogenesis and/or the occurrence of seizures.

Indeed, clinical data and animal experiments support the hypothesis that primary vascular lesions and, specifically BBB dysfunction, trigger a chain of events leading to epilepsy (For reviews see Friedman et al., 2009; Shlosberg et al., 2010). Significant and long-lasting dysfunction of the BBB are a hallmark of cortical injuries, regardless of etiology (Cervos-Navarro and Lafuente, 1991; Tomkins et al., 2001, 2008). BBB dysfunction is often observed in both human and animal models of traumatic and ischemic brain injuries, within and surrounding brain tumors, infectious and inflammatory brain diseases and neurodegenerative diseases, including Alzheimer's and vascular dementia (Abbott et al., 2006; Neuwelt, 2004). In epilepsy, magnetic resonance imaging studies in patients with posttraumatic epilepsy demonstrated permeability of BBB to contrast agents, co-localized with the presumed epileptic focus (Tomkins et al., 2008, 2011). Ultrastructural studies on human-resected epileptic tissue show clear BBB abnormalities, including increased micropinocytosis, a thickening of the basal membrane, and the presence of abnormal tight junctions (Cornford, 1999; Cornford and Oldendorf, 1986; Kasantikul et al., 1983). However, no prospective clinical study was performed to test directly to what extent BBB impairment can predict the development of epilepsy in injured patients. In animal studies, a role for BBB opening was suggested in the progression of TLE based on the finding of serum albumin presence in brain parenchyma following SE, and a positive correlation between the extent of BBB opening and the number of seizures (van Vliet et al., 2007). Experimental focal opening of the BBB in the rat neocortex has been shown to result in epileptogenesis, evident by the delayed development of paroxysmal hypersynchronous activity recorded *ex vivo* (acute slice preparation) and *in vivo* (Ivens et al., 2007; Seiffert et al., 2004; Tomkins et al., 2007). This epileptogenesis was recapitulated by exposure of brain cortex to serum albumin. Extravasation of serum albumin into the cerebral cortex microenvironment activates a transforming growth factor beta (TGF β) receptor-mediated signaling cascade in astrocytes (Cacheaux et al., 2009; David et al., 2009; Ivens et al., 2007 and see below).

The possible involvement of serum albumin in astrocytic activation and proliferation is supported by previous studies showing serum albumin inducing proliferation of fibroblasts (Tigyi et al., 1995), calcium signaling as well as DNA synthesis in cultured astrocytes (Nadal et al., 1995). On the basis of their studies, Nadal et al. (1995) concluded that there albumin's effect is receptor-mediated. Although albumin is the most abundant protein in the serum, other blood-born proteins may also have a role in the epileptogenic process. For example, it has been recently shown that the serum protein, thrombin, through protease-

activated receptor 1 (PAR1), lowers epileptic seizures threshold in the hippocampus CA3 region and produces a long-lasting enhancement of CA1 neurons reactivity to afferent stimulation (Maggio et al., 2008). Thus, it seems plausible that damage to the microvasculature during brain insults and the extravasation of serum proteins, lead to the transformation of the neighboring astrocytes as a primary step in the epileptogenesis. Additionally, increase in vessels permeability to blood proteins leads to antibodies extravasation into brain tissue (van Vliet et al., 2007; Rigau et al., 2007), including autoantibodies to neuronal receptors and channels (Bien and Scheffer, 2011). It is not known to what extent such autoantibodies contribute to the pathogenesis of the disease; however, in some cases a clear respond to immunotherapy has been shown (Vincent et al., 2010).

THE TGF β PATHWAY AND EPILEPTOGENESIS

TGF β s are pleiotropic cytokines that play a pivotal role in intercellular communication (for review see Massague and Wotton, 2000; Shi and Massague, 2003), and are involved in cell growth, embryogenesis, differentiation, morphogenesis, wound healing, immune response, and apoptosis in a wide variety of cells (Blobe et al., 2000; Gold and Parekh, 1999). TGF β signaling is mediated mainly by two serine threonine kinase receptors, TGF β R1 and TGF β R2, which activate an intracellular signaling system, such as phosphorylation of the Smad protein complex and the p38 mitogen-activated protein kinase (MAPK) pathway. TGF β is upregulated in many disease conditions (for reviews see Szelenyi, 2001; Vitkovic et al., 2001): TGF β 1 expression is upregulated in the brains of individuals suffering from multiple sclerosis, AIDS, Alzheimer's disease, stroke, tumors, or trauma. TGF β has also been shown to be elevated in the cerebro-spinal fluid of some patients following brain injury (Phillips et al., 2006), to be produced in neurons after ischemia (Zhu et al., 2000), to be involved in pericyte-induced alterations in BBB function (Dohgu et al., 2005) and in microglial activation (Schilling and Eder, 2003). Smad3 null mice show reduced glial scarring after cortical stab wound injury (Wang et al., 2007), further supporting the central role of TGF β in injury. Indeed, while many researchers consider TGF β 1 to be a "protective" cytokine (Brionne et al., 2003; McNeill et al., 1994; Prehn et al., 1993; Zhu et al., 2002), it has also been found to exacerbate excitotoxicity (Mesples et al., 2005; Prehn et al., 1994). These apparent contradicting results in different experimental systems and models can be explained partly by data showing that TGF β 1 actions are dependent on cell type and condition, resulting even in opposing outcomes.

Is TGF β associated with epileptogenesis? The potential involvement of TGF β in epileptogenesis is supported by animal experiments showing TGF β upregulation in neurons from amygdale-kindled rats (Plata-Salaman et al., 2000). Aronica et al. (2000) showed TGF β expression in astrocytes from the hippocampus of SE-experienced rats. Recent studies in rats in BBB-disrupted animals demonstrated that serum albumin binds to TGF β R and activate TGF β signaling (Cacheaux et al., 2009; Ivens et al., 2007). Accordingly, transcriptome analysis revealed a strikingly similar transcription modulation patterns in animals exposed to BBB dysfunction, serum-derived albumin or following direct brain exposure to physiological levels of TGF β 1. However, the detailed mechanisms and cellular pathways bridging TGF β signaling to seizures in different cell types are still a matter of investigation.

THE ROLE OF ASTROCYTES IN EPILEPTOGENESIS FOLLOWING BBB DYSFUNCTION

The interactions observed between serum albumin and TGF β receptors and the uptake of serum albumin preferentially into glia cells within a few hours following BBB opening and prior to the development of seizures (Ivens et al., 2007) raised the hypothesis that glia

functions and dysfunctions play a key role in the generation of the epileptic network. Furthermore, direct neocortical application of TGF β 1 on the brain surface and activation of TGF β signaling, resulted in a prominent transcriptional-mediated change in expression levels of astrocytic genes (David et al., 2009).

Changes in glia morphology and function are a hallmark in the epileptic tissue in many of the patients (for reviews see Binder and Steinhauser, 2006; Heinemann et al., 2000; Jabs et al., 2008; Wetherington et al., 2008) and are summarized in detail in this special issue. Recent studies demonstrating a broad range of physiological effects of brain glia on neuronal activity, including the direct modulation of synaptic transmission and plasticity, point to a potential role of glia in epileptogenesis. It is thus hypothesized that the “transformation” or “activation” of glia in the presence of dysfunctional BBB leads to the reorganization of the neuronal network, which characterizes the epileptic brain. The term “activated” glia is often used, but this is unfortunately a very imprecise term and it is not clear what “activation” really means, to what extent it is a “single uniform” state, and how persistent is it. Is it just the increased expression of glial fibrillary acidic protein (GFAP), or is it associated with alterations in specific functions such as the expression of pumps and ion channels? How the connectivity between astrocytes is altered and what role activation of astrocytes has in the synthesis of cytokines and their release? Since cells which express GFAP and therefore termed “astrocytes” include radial glia such as Müller and Bergmann cells, but also astrocytes which have been termed glutamate transporter or passive cells as well as astrocytes which express AMPA receptors and voltage gated ion channels (Glu-R cells or complex astrocytes), it is unclear what activation of astrocytes really means and to what extent all types of glia (or even astrocytes) are affected in the same way. Different “types” of glia thus differ not only in morphology, but also in functions (e.g. in the extent of electrical coupling between them, spatial buffering of extracellular potassium, transport of glutamate and attachment to vessels). Here, we will summarize physiological changes observed in activated astrocytes following BBB opening, and from these observations we will deduce a working hypothesis on their role in remodeling the neuronal network towards lowering its threshold to large-scale synchronization, development and propagation of seizures.

Several functional changes in astrocytic properties, which may be relevant to the epileptogenesis process, have been found in the BBB-impaired cortex. These include: (1) Reduced expression of potassium inward rectifying channels (Kir4.1) and water channels (aquaporin 4, AQP4); (2) Reduced expression of gap junctions; and (3) impaired glutamate metabolism.

Reduced Kir 4.1 and AQP4

BBB opening, brain exposure to albumin or to TGF β 1 resulted in downregulation of both Kir4.1 and AQP4 (David et al., 2009; Perillan et al., 2002). Both channels are co-localized most abundantly in astrocytic endfeet and considered crucial for the regulation of the brain's extracellular potassium ($[K^+]_o$) and water fluxes. Spatial buffering of $[K^+]_o$ occurs when its concentrations increase nonhomogeneously throughout the different cortical layers. This is usually the case during physiological activation and even more so during repetitive activation (Dietzel et al., 1989; Pumain and Heinemann, 1985). In the hippocampus, $[K^+]_o$ is usually maximally increased in pyramidal layer of Ammon's horn or in the granular layer of the dentate gyrus (Krnjevic et al., 1982). In the neocortex, $[K^+]_o$ accumulation is often maximal in the deep layers (Hablitz and Heinemann, 1987). Local increase in $[K^+]_o$ depolarizes astrocytes which carry large resting conductance to potassium. If these are spatially extended and/or electrically coupled (Wallraff et al., 2006) this depolarization will spread through the glial network; consequently, the local depolarization does not correspond to the local accumulation of $[K^+]_o$, leading to a driving force for potassium

influx at sites of maximal $[K^+]_o$ elevation and efflux at remote sites. This mechanism underlies the observed mismatch between transient and focal increase in $[K^+]_o$ and the associated glial depolarization (Futamachi and Pedley, 1976). This prevents excess increase in $[K^+]_o$ by dissipating it much faster than expected from pure diffusion (Gardner-Medwin and Nicholson, 1983). The influx of potassium into glia also contributes to the generation of slow negative potentials at the site of maximal K^+ uptake which may affect neurons by field effects mediating ephaptic interactions. Finally, due to differences in ion transport numbers, transglial influx of potassium is associated with a shrinkage of the extracellular space at sites of maximal neuronal activity: As K^+ ions enter the glial compartment, potential gradient emerge; The corresponding current in the extracellular space is predominantly carried by Na^+ and Cl^- , the predominant ions. Thus, Na^+ is transported to the site of K^+ accumulation while Cl^- moves away. The Cl^- and K^+ ions are only partially replaced by Na^+ , resulting in a decrease of extracellular osmolarity, leading to a water flux into the cells, and a shrinkage of the extracellular space. At remote sites, the opposite effect is expected due to K^+ outflow from glia and Cl^- transport to these sites (Lux et al., 1986). Which K^+ channels account for the potassium movements into astrocytes? It is likely that Kir 4.1 channels are very important in this regard. These ion channels are very sensitive to low concentrations of barium which significantly leads to depolarization of astrocytes. In addition, it is likely that two pore domain K^+ channels also contribute to the influx of K^+ (Pasler et al., 2007). These channels are also sensitive to barium but require higher concentrations. These differences in pharmacological sensitivity to barium permit testing changes in functional expression of Kir 4.1 channels during epileptogenesis in the dysfunctional BBB model. Indeed, while in control cortex low concentrations of barium increased $[K^+]_o$ levels and slowed the decay of $[K^+]_o$ in response to iontophoretic application, this effect was almost absent in the epileptogenic tissue (i.e. 24 h after BBB opening) and resembled the situation found in human epileptic cortex (Jauch et al., 2002). These results, together with mRNA and protein levels analysis confirmed that downregulation of Kir 4.1 channels characterizes transformed astrocytes in the epileptogenic tissue (David et al., 2009; Ivens et al., 2007). Loss of I_{kir} was also demonstrated in cortical astrocytes following exposure to fluid percussion model of traumatic brain injury with serum extravasation (Stewart et al., 2010). Several lines of evidence support a role for Kir 4.1 down regulation in epileptogenesis: (1) mice with glial specific Kir 4.1 deletions suffer from epilepsy and die relatively soon after birth (Djukic et al., 2007); (2) homozygous missense mutations in the KCNJ10 gene (encoding for the Kir 4.1 channel) has been recently described in five children with EAST (epilepsia, ataxia, moderate sensorineural deafness, and a renal salt losing tubulopathy) syndrome (Bockenbauer et al., 2009); (3) Impaired $[K^+]_o$ buffering due to downregulation of Kir4.1 channels has been reported in the hippocampus of pilocarpine-treated epileptic rats (Gabriel et al., 1998) and in the sclerotic hippocampus of TLE patients (Jauch et al., 2002; Kivi et al., 2000; Schroder et al., 2000).

Reduced expression of aquaporin channels may also contribute to the epileptogenic role of transformed astrocytes. AQP4 is expressed in astrocytes throughout the central nervous system, particularly at the BBB. AQP4-null mice indeed show reduced $[K^+]_o$ buffering and prolonged duration of induced-seizures (Binder et al., 2006; Strohschein et al., 2011).

Role of Gap Junctions

Gap junctions are functional channels between cells that allow for controlled transcellular movement of molecules between cells, and are comprised of connexin proteins. As described above, astrocytes are coupled via gap junctions to form large cellular networks which facilitate spatial buffering of small molecules (e.g., K^+). Interestingly, the expression of the astrocytic gap junction proteins connexin 30 and 43 is reduced in the BBB-induced epileptogenic cortex. This reduction may contribute to the activity-dependant accumulation

of $[K^+]_o$ (David et al., 2009). Notably, while block of Kir 4.1 and 2 Pore domain channels have dramatic effects on K^+ homeostasis, the effects of loss of connexins are less pronounced. Mice lacking connexin 30 and 43 in astrocytes show only mild disturbances in $[K^+]_o$ homeostasis. Nevertheless, afferent stimulation results in a larger rise in $[K^+]_o$ in these mice compared with controls, lower threshold for seizures, accelerated propagation of spreading depolarization, and enhanced locomotor activity (Theis et al., 2003; Wallraff et al., 2006).

Impaired Glutamate Metabolism

Within the neurovascular unit, glial cells have a key role in the uptake and metabolism of glutamate. Following BBB dysfunction activated astrocytes show reduced levels of mRNA encoding for the astrocytic glutamate transporters of the solute carrier family 1, subfamily A members SLC1A2 and SLC1A3 (Chaudhry et al., 1995; Su et al., 2003). The effect was specific to astrocytes since SLC1A1 (preferentially expressed in neurons; see (Rothstein et al., 1994) did not show significant changes in expression levels. In addition, the astrocytic glutaminase and glutamine synthetase (Derouiche and Frotscher, 1991) were also downregulated. Astrocytic buffering of glutamate (by glutamine synthetase) plays a lead role in neuronal hyperexcitability regulation (Eid et al., 2008). Data showing reduced glutamate buffering capacity during epileptogenesis further stress the potential role of altered glutamate metabolism in astrocytes in the development of epilepsy (David et al., 2009). In addition, astrocytic glutamate release has been described in several preparations and implicated to contribute to a slow, TTX resistant, NMDA sensitive neuronal inward current (Tian et al., 2005). It is not clear, however, to what extent such release contributes to epileptogenesis under BBB dysfunction. There is no direct evidence for increased glutamate release from transformed astrocytes, although one potential mechanism is the upregulation of TNF α which is a prominent modulator of glutamate release (Bezzi et al., 2004). Finally, reduced glutamate uptake in transformed astrocytes may also interfere with the production of glutathione: astrocytes use glutamate to uptake cystine, which is used to synthesize glutamyl-cysteine, which is released from astrocytes for synthesis of glutathione in neurons. Downregulation of neuronal and glial glutathione would weaken defense mechanisms against free radicals and would be expected to result in increased damage (Schuchmann and Heinemann, 2000). Interference with glutamate transport into astrocytes will also affect the detoxification of glutamate to glutamine and thus might interfere with detoxification of ammonium. Ammonium disturbs Cl^- transporters and thus might contribute to a reduced efficacy of GABA-mediated synaptic inhibition.

SUMMARY

Figure 1 summarizes the role played by vascular injury and transformed (activated) astrocytes in reducing seizure threshold. However, it is important to distinguish between seizure generation and epileptogenicity—which carries a more complex and long-lasting network modifications. The response of astrocytes to vascular injury is rapid, and seems to be directly related to the diffusion of the most common serum protein, albumin, via the dysfunctional barrier. Thus, it is proposed that transformation of astrocytes starts during the latent period of epileptogenesis and is regulated by TGF β signaling. Under these conditions, astrocytes show a less “rigid” control on potassium and glutamate levels, most significantly during repetitive activation. We postulate that transformed astrocytes may allow enhanced axonal sprouting and synaptogenesis as part of the brain response to injury. One view may thus be that epilepsy develops when the normal network response to injury has not been properly controlled due to lasting dysfunction of the local microvasculature. This hypothesis, while still requiring careful experimental control and appropriate experimental design, may change our view on the mechanisms underlying epileptogenesis and potential targets for its prevention. If there is a critical time window in which BBB dysfunction

predicts epileptogenesis following insult, it may become an early biomarker for the development of epilepsy and would facilitate studies on its prevention (Pitkänen, 2010). To address BBB dysfunction as a biomarker for the prediction of epilepsy, large prospective human studies will be required. Quantitative methods for evaluating focal BBB damage noninvasively are mandatory for such a study. While few imaging approaches have been proposed (Tomkins et al., 2001, 2008, 2011) there is still an urgent need for the development of such methods. In parallel, animal studies are required for better understanding the mechanisms underlying BBB damage and repair following insult, to allow a selective and efficient targeting and modification of these processes.

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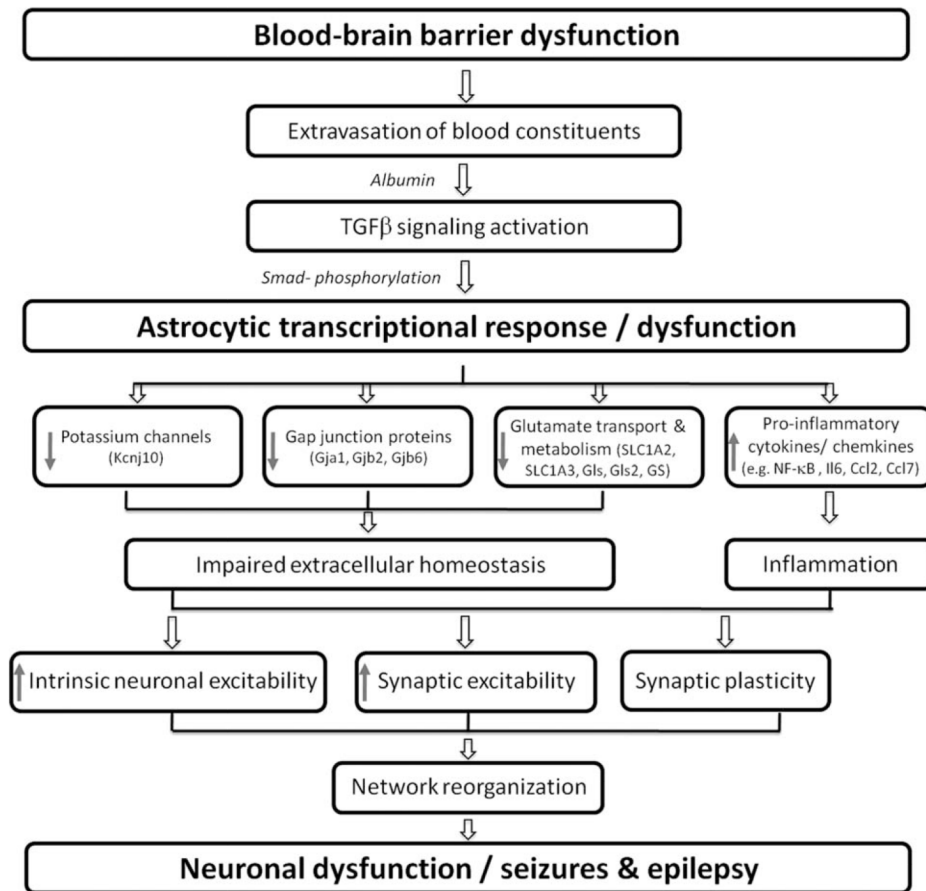
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**Fig. 1.**

Summary diagram on the role of BBB dysfunction and astroglia functions in epileptogenesis. The damage to endothelial cells leads to dysfunction of the BBB and the extravasation of serum proteins into the neuropil. Albumin activates TGF β signaling in astrocytes leading to transcriptional response associated with their transformation into “active” astrocytes. Transcriptional response includes the down regulation of Kcnj10 (inward rectifier 4.1 potassium channel) together with Gja1, Gjb2, and Gjb6 (gap junction proteins, connexins 43, 26, and 30, respectively). In addition, genes associated with glutamate metabolism are downregulated, including the mRNA coding for the astrocytic glutamate transporters of the solute carrier family 1, subfamily A members SLC1A2 and SLC1A3. Glutaminase (Gls, Gls2) and glutamine synthetase (GS) are also downregulated. Together, homeostasis of the extracellular brain environment is impaired, leading to enhanced neuronal excitability. In addition, upregulation of proinflammatory cytokines and chemokines, as part of the local inflammatory response, may also contribute to the increase in neuronal excitability, associated with reorganization of the local neuronal network, typical hallmark of the epileptic brain.