



Anion transport and GABA signaling

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Whereas activation of GABA_A receptors by GABA usually results in a hyperpolarizing influx of chloride into the neuron, the reversed chloride driving force in the immature nervous system results in a depolarizing efflux of chloride. This GABAergic depolarization is deemed to be important for the maturation of the neuronal network. The concept of a developmental GABA switch has mainly been derived from *in vitro* experiments and reliable *in vivo* evidence is still missing. As GABA_A receptors are permeable for both chloride and bicarbonate, the net effect of GABA also critically depends on the distribution of bicarbonate. Whereas chloride can either mediate depolarizing or hyperpolarizing currents, bicarbonate invariably mediates a depolarizing current under physiological conditions. Intracellular bicarbonate is quickly replenished by cytosolic carbonic anhydrases. Intracellular bicarbonate levels also depend on different bicarbonate transporters expressed by neurons. The expression of these proteins is not only developmentally regulated but also differs between cell types and even subcellular regions. In this review we will summarize current knowledge about the role of some of these transporters for brain development and brain function.

Keywords: GABA, pH, chloride, bicarbonate, ion transporter

INTRODUCTION

GABA signaling has a wide spectrum of functions in individual neurons and neuronal networks in the brain. It is well known that in the mature brain, GABA acts as the main inhibitory transmitter due to activation of hyperpolarizing chloride currents through GABA_A receptors (Farrant and Kaila, 2007). In contrast, during early brain development, GABAergic transmission is assumed to provide the main excitatory drive in neuronal networks, at a time when glutamatergic synaptic contacts are less frequent than GABAergic synapses (Ben-Ari et al., 1989). Although this functional switch from excitatory to inhibitory GABA action during brain development has been observed in a wide range of preparations and different animal species, most of the experimental evidence relies on *in vitro* studies (comprehensively reviewed by Ben-Ari et al., 2007). They were performed using several electrophysiological means like intracellular recordings (Mueller et al., 1984; Luhmann and Prince, 1991) and less invasive techniques including perforated patch (Owens et al., 1996; Yamada et al., 2004) and cell-attached measurements (Wang et al., 2003; Rheims et al., 2008; Kirmse et al., 2010). Consistently, several groups reported intracellular calcium increases in immature neurons upon GABA application, most likely due to depolarization mediated activation of voltage-gated calcium channels (Yuste and Katz, 1991; Owens et al., 1996; Yamada et al., 2004; Kirmse and Kirischuk, 2006; Kirmse et al., 2010). However, *in vivo* evidence for depolarizing GABA action in immature neuronal networks is rare (Brustein et al., 2003) and often indirect (Sipila et al., 2006). Metabotropic GABA_B-receptors are coupled to calcium or

potassium channels, and cyclic AMP signaling. Although there is quite recent evidence that the non-hyperpolarizing activation of GABA_B-receptors during development promotes neuronal migration and morphological maturation (Bony et al., 2013), this review will focus on GABA_A-receptor signaling and how this relates to anion-transport.

The functional relevance of GABA_A-receptor activation for activity patterns in immature neuronal networks has been investigated in different model systems *in vitro* and *in vivo*. In the immature hippocampus, it is widely accepted that GABAergic excitation drives the typical spontaneous network activity known as giant depolarization potentials or GDPs (Ben-Ari et al., 1989; Bonifazi et al., 2009). On the other hand in neocortex, glutamatergic excitation was shown to be dominating in the early generation of network activity like early network oscillations or ENOs (Garaschuk et al., 2000) and spindle-bursts (Minlebaev et al., 2007). However, more recent results suggest that GABA_A receptor activation also supports the generation of early neocortical network activity (Allene et al., 2008). It is worth noting that depolarizing GABA_A-receptor activation not necessarily needs to be excitatory (Morita et al., 2005, 2006). The GABA-induced increase in membrane conductance can also cause a so-called shunting inhibition, because according to Ohm's law, the drop in membrane resistance would decrease the voltage change caused by a certain depolarizing current (e.g., a glutamatergic synaptic input). The inhibitory effect of shunting does not depend on the polarity of a GABA-induced membrane potential change and therefore on the chloride and bicarbonate reversal potentials, but

is solely due to the GABA-induced drop in input resistance. Nevertheless, the paradoxical situation could happen that a depolarizing GABA_A-receptor activation leads to an inhibitory restriction of network activity (Minlebaev et al., 2007).

Although many initial steps in early neuronal network development are genetically determined, there is a large body of evidence, that the proper functional maturation of cortical neuronal circuits is highly activity-dependent (Katz and Shatz, 1996). However, to what extent the spontaneously occurring network activity, partly driven by GABA_A-receptor activation, contributes to the functional maturation in the pre-sensory period of the brain is still unclear. Undoubtedly, a fine tuned balance between excitation and inhibition at any stage of development is essential for providing a proper function of neuronal networks. In this context, the potentially depolarizing mode of GABA_A-receptor activation during early development has been considered to contribute to the higher liability to pathological events like epileptic seizures during childhood (see Kirmse et al., 2011 for review). Later in development, the increasing GABAergic inhibition governs the on- and offset of the so-called critical period in the visual system, which is crucial for the activity-dependent functional refinement of the participating cortical circuits (Hensch et al., 1998; Fagiolini and Hensch, 2000). Therefore, GABA_A-receptor activation plays a pivotal role at various developmental stages for the maturation, refinement and proper function of neuronal networks.

The ionotropic GABA_A receptors are pentamers of 19 different subunits, which are grouped into eight different families according to sequence homology (α 1–6, β 1–3, γ 1–3, δ , ϵ , θ , π , ρ 1–3; Farrant and Kaila, 2007). Although the different receptor assemblies have different properties and different distributions, only chloride and bicarbonate are conducted under physiological conditions (Bormann et al., 1987). It is assumed that the relative bicarbonate/chloride permeability of ionotropic GABA_A receptors ranges between 0.18 and 0.6 (Fatima-Shad and Barry, 1993). Because of a variety of different ion transporters within the plasma membrane, which mediate acid extrusion either by extrusion of H⁺ or by accumulation of bicarbonate, the bicarbonate equilibrium potential is much more depolarized (around –10 mV) than the resting membrane potential (Roos and Boron, 1981). Hence bicarbonate can only mediate a depolarizing current under normal conditions. Quite in contrast, the equilibrium potential for chloride is close to the resting membrane potential. Hence chloride can mediate both hyperpolarizing and depolarizing currents depending on the existing gradient, which is regulated during development. However, a depolarizing action of GABA does not exclude an inhibitory action but may result in shunting inhibition as outlined above.

ELECTROPHYSIOLOGICAL METHODS TO QUANTIFY GABA FUNCTION

Before fluorescent indicators have been available, ion-selective micro-electrodes were the gold standard for the measurement of chloride activity and pH in the intracellular compartment (Walker, 1971; Thomas, 1974; Ammann et al., 1981). Because of the invasive nature of this method – the cellular membrane needs to be impaled by a double-barreled sharp micro-electrode – it was impossible to determine the ion activity of interest quantitatively without changing it at the same time due to the measurement. Leakage currents at the site of impalement and intracellular perfusion by the solution

of the reference channel are only two possible sources of measurement errors. Nevertheless, this method provided first important insights into the ionic mechanisms of pH regulation or chloride homeostasis (Thomas, 1977; Vaughan-Jones, 1979).

In order to keep the ionic composition of the intracellular compartment unchanged, most other electrophysiological means are based on cell-attached patch clamp recordings. In the following, we describe several methods which have been developed to determine membrane potentials and membrane currents without disrupting the plasma membrane. To provide electrical access to the intracellular compartment without interference with the intracellular milieu, a technique called perforated patch clamp was developed, originally using ATP in the pipette solution as a membrane permeabilizing agent (Lindau and Fernandez, 1986). In subsequent modifications of this method, different ionophores were added to the pipette solution, which, during the experiment, incorporate into the membrane patch under the pipette tip. Ionophores are lipid-soluble molecules which form hydrophilic pores in the cell membrane and mediate electrical access to the intracellular compartment without destroying the barrier function of the membrane patch for the ion of interest. In early studies, nystatin and amphotericin B were used to achieve low resistance electrical access (Horn and Marty, 1988; Rae et al., 1991). However, as a major drawback these substances also lead to chloride redistribution. Subsequently, gramicidin D, a mixture of different antibiotics, was added because it is only permeable for monovalent cations and uncharged low molecular substances but impermeable for chloride, leaving its concentration gradient over the cell membrane intact (Ebihara et al., 1995; Kyrozis and Reichling, 1995). However, indirect changes in intracellular chloride concentration are conceivable because it is mainly regulated by cation/chloride co-transporters. Several groups have applied this method successfully to determine e.g., the chloride equilibrium potential in hippocampal cells during brain development (Mohajerani and Cherubini, 2005; Sipila et al., 2006; Tyzio et al., 2007; Pfeffer et al., 2009).

As mentioned previously, the chloride concentration gradient is not the only determinant governing GABA-induced membrane potential changes, because GABA_A-receptor channels are also permeable for other anions like bicarbonate. In order to quantify the GABA-reversal potential directly, Tyzio et al. (2003, 2006) developed a non-invasive method to measure the resting membrane potential and the GABA reversal potential at the same cell, using single *N*-methyl-D-aspartate and GABA channel recordings. The combination of both experimental approaches at the same cells provides the driving force for GABA-induced currents and the resting membrane potential in absolute numbers and, thereby, the GABA equilibrium potential. Although this is up to now the most reliable method to quantify these parameters, it is hardly applicable to complex preparations like *in vivo* recordings. If only the non-invasive quantification of the membrane potential change due to GABA-receptor activation is of interest, an alternative method described by Verheugen et al. (1999) can be applied (Kirmse et al., 2010). Assuming symmetrical potassium concentrations, this method uses the fact, that the reversal potential of voltage-dependent potassium currents in the cell-attached configuration represents a good estimate of the cell membrane potential

(Verheugen et al., 1999). However, potential changes in intracellular potassium concentration would flaw the correct membrane potential quantification. Because a voltage-ramp protocol has to be applied for every time point of interest, the time resolution of this method is rather slow and phasic membrane potential changes are difficult to catch. In these cases, applying a current-clamp recording protocol in the cell-attached configuration might be beneficial, because it can provide a good estimate of the polarity of an induced membrane potential change at high time resolution (Perkins and Wong, 1996; Mason et al., 2005).

In summary, various non-invasive electrophysiological methods provide valuable estimates of GABA equilibrium potentials under different conditions *in vitro* and *in vivo* and enable the measurement of relative or absolute cell membrane potential changes without disturbing the intracellular milieu. In concert with complementary optical methods for quantification of intracellular chloride concentration and pH, they draw a detailed image of GABA-mediated physiological processes.

OPTICAL METHODS TO QUANTIFY GABA FUNCTION

A big step forward was the development of fluorescent indicator dyes which enabled the optical measurement of intracellular pH and chloride concentrations (Rink et al., 1982; Illsley and Verkman, 1987). The initially used small molecular fluorescent chloride indicator dyes were quinoline derivatives which change their fluorescent intensity upon changes in chloride concentration by a mechanism called collision quenching (Chen et al., 1988). With increasing chloride concentration the probability of a collision between a chloride ion and an indicator molecule increases and therefore, its fluorescence intensity decreases by quenching. A notable feature of this mechanism is, that different from the popular calcium indicator dyes, which change their fluorescence intensity upon binding to calcium, these chloride indicator dyes do not introduce any exogenous buffer capacity to the intracellular milieu, because no binding to the ion of interest takes place. Nevertheless, their excitation spectra in the ultraviolet range give rise to strong bleaching and photodynamic damage (Inglefield and Schwartz-Bloom, 1997). However, the combination of these dyes with two-photon imaging is able to reduce both side effects significantly (Marandi et al., 2002). Several years earlier, the measurement of intracellular pH has been revolutionized by the invention of BCECF, a fluorescence indicator derived from fluorescein, by Roger Tsien and coworkers (Rink et al., 1982). The absorption spectrum of BCECF is shifted depending on changes in pH and by applying ratiometric excitation the indicator can be calibrated to absolute pH units (Graber et al., 1986; Bright et al., 1987).

The unspecific loading of the exogenously applied fluorescent indicator dyes prevents a cell-specific labeling. Therefore, much effort was invested to develop genetically determined chloride indicator dyes (see Bregestovski et al., 2009 for review). Starting point of this development was the chloride binding property of the yellow fluorescent protein (YFP) a derivative of the green fluorescent protein (GFP). Because the sensitivity of wild-type YFP to chloride is low, many random chloride binding site mutations of YFP were tested and analyzed for improved sensitivity (Galiotta et al., 2001). Besides the possibility of cell-specific

expression of the chloride indicator, YFP-based indicator dyes have additional advantages. Different from quinolone-derived dyes, the optimal excitation wavelength is located in the visible range, providing less bleaching and photodynamic damage during the experiments. In addition, leakage during the measurements is less pronounced due to their relative large molecular weight of about 27 kDa (Bregestovski et al., 2009). Finally and different from calcium measurements, intracellular indicator concentrations are orders of magnitude smaller than that of the ion of interest, therefore exogenous buffering of chloride is negligible. On the other hand, there are also some disadvantages of YFP-based indicator dyes. Keeping in mind that changes in intracellular chloride concentration are often accompanied by changes in pH, the significant pH-sensitivity of many YFP derivatives is the most serious one. The only way to circumvent this restriction is the independent monitoring of pH changes and subsequent data correction. Another problem of YFP-based chloride indicator dyes are their rather slow kinetics or poor sensitivity, which either limits the detection of fast chloride concentration changes or leads to poor resolution at physiological chloride levels (Galiotta et al., 2001). Originally, YFP-based indicator dyes were not able to report absolute levels of chloride concentration, because they lack an isosbestic point, at which they are insensitive to chloride concentration changes. Because the absolute measurement of intracellular chloride concentrations is imperative to determine the chloride equilibrium potential or the driving force for chloride, Kuner and Augustine (2000) developed a ratiometric chloride indicator named Clomeleon. Clomeleon uses the chloride-dependent interaction of two fluorophores (cyan fluorescent protein (CFP) as donor and a variant of YFP called topas fluorescent protein (TFP) as acceptor) by Förster energy transfer (FRET). Upon chloride binding to TFP, the efficiency of FRET between CFP and TFP declines. As a consequence, the ratio of TFP and CFP fluorescence emission drops with increasing chloride concentrations. Because the emission spectrum comprises an isosbestic point, calibration to absolute chloride levels is possible (Kuner and Augustine, 2000). Unfortunately, the sensitivity of Clomeleon with an IC_{50} of more than 160 mM is rather low and, at physiological levels, makes reliable measurements of absolute intracellular chloride concentration very difficult (Kuner and Augustine, 2000). Following genetic engineering of the YFP chloride binding site yielded a higher sensitivity of the resulting indicator called Cl-sensor with an IC_{50} around 30 mM, much closer to physiological intracellular chloride concentrations (Markova et al., 2008). However, Clomeleon and Cl-sensor have slow response kinetics and share the pH sensitivity of all YFP-based chloride indicators (Bregestovski et al., 2009).

To overcome the main drawbacks of Clomeleon and Cl-sensor, a new ratiometric but non-FRET-based sensor was developed (Arosio et al., 2010). This new indicator, called ClopHensor, is suitable for the simultaneous quantification of intracellular pH and chloride concentration. Therefore, a variant of enhanced GFP (E^2GFP) with pH sensitivity and sensitivity to chloride comparable to Cl-sensor was fused with a pH- and chloride-insensitive monomeric DsRed. The E^2GFP part of the fused protein allows chloride-independent ratiometric quantification of pH by exciting it subsequently at 458 and 488 nm. The ratiometric measurement

of chloride concentration by alternative exciting E²GFP at 458 nm and the chloride-insensitive DsRed at 543 nm requires in addition the calibration at different pH values. However a simultaneous quantification of intracellular pH is now possible, ClopHensor still suffers from rather low sensitivity to chloride with an IC₅₀ around 40 mM (Arosio et al., 2010; Mukhtarov et al., 2013). A more recent variant of ClopHensor exhibits a higher sensitivity with an IC₅₀ of 20 mM, but at the expense of a significant lower dynamic range (Mukhtarov et al., 2013). In summary, the development of the ratiometric indicator ClopHensor provides a most valuable means for the simultaneous quantification of pH and chloride concentration, but variants with higher sensitivity to chloride are desired to increase the quantification accuracy at physiological concentration levels.

ION TRANSPORTERS INVOLVED IN THE REGULATION OF NEURONAL CHLORIDE AND BICARBONATE LEVELS

The role of cation-chloride co-transporters (**Figure 1**) in the regulation of the intraneuronal chloride concentration has been extensively studied and follows a well-defined developmental sequence with a high chloride concentration in immature neurons due to neuronal chloride accumulation. Chloride accumulation largely depends on the action of the Na⁺/K⁺/2Cl⁻ co-transporter NKCC1 (Yamada et al., 2004; Sipila et al., 2006; Achilles et al., 2007; Blaesse et al., 2009; Pfeffer et al., 2009). But other mechanisms to accumulate chloride exist and maintain GABA depolarizing even in the absence of NKCC1 (Pfeffer et al., 2009). One candidate is the anion-exchanger AE3, which normally accumulates chloride in exchange for intracellular bicarbonate and thereby raises intracellular chloride levels. The so-called GABA switch from excitatory to inhibitory is brought about by the incipient expression of the cation-chloride co-transporter KCC2 (Rivera et al., 1999; Hübner et al., 2001; Stein et al., 2004), which extrudes chloride out of the cell. From our knockout studies other KCl co-transporters like KCC1 (Rust et al., 2007), KCC3 (Boettger et al., 2003; Seja et al., 2012), or KCC4 (Boettger et al., 2002) appear to be less important in the control of neuronal chloride levels.

Several transporters either exchange bicarbonate and chloride or couple the transport of bicarbonate to sodium (**Figure 1**). For

bicarbonate transporters that also transport chloride, the net effect for GABA_A receptor signaling is difficult to predict. It is evident that changes of bicarbonate levels do not only directly affect the currents mediated by GABA_A receptors, but are also tightly linked to alterations of the pH both within the cells and in the extracellular space, which can have a multitude of different effects. In neurons with a high intracellular chloride concentration as in the immature nervous system, however, the effect of bicarbonate on the GABA_A reversal potential is quite low according to the Goldman equation (Farrant and Kaila, 2007).

ANION EXCHANGERS

Whereas there are numerous reviews addressing the role of cation-chloride co-transporters for GABA transmission (Blaesse et al., 2009), the role of bicarbonate and hence the role of neuronal mechanisms to control intracellular bicarbonate levels are less acknowledged. Bicarbonate transport is mediated by members of the SLC4A or the SLC26A family of proteins. As members of the SLC26A family appear to play a minor role for neurons (Dorwart et al., 2008; Majumdar and Bevensee, 2010), we restrict our review to some selected members of the SLC4A family with known relevance for neuronal function and refer to some other more complete reviews (Romero et al., 2004; Dorwart et al., 2008; Majumdar and Bevensee, 2010). The SLC4 family can be subdivided into four main branches: the sodium independent anion-exchangers (AE1, AE2, and AE3) recently reviewed in Alper (2009), and sodium-coupled bicarbonate transporters recently reviewed in Majumdar and Bevensee (2010; **Figure 2**). The role of AE4 is still unclear: although originally cloned as a sodium-independent anion-exchanger, there is evidence that it rather serves as a Na⁺/HCO₃⁻ co-transporter (Parker et al., 2002). It localizes highly specific to the basolateral membrane of mouse type B intercalated cells and is involved in chloride recovery by these cells (Chambrey et al., 2013). Na⁺/HCO₃⁻ co-transporters can be either electroneutral (NBCn1 and NBCn2) or electrogenic (NBCe1 and NBCe2), whereas the sodium-dependent anion-exchangers (NDAE or NDCBE and NCBE) are electroneutral. SLC4A11 differs from the other family members because it rather mediates borate transport and is hence termed as BTR1 (Park et al., 2004). In the following we will focus on

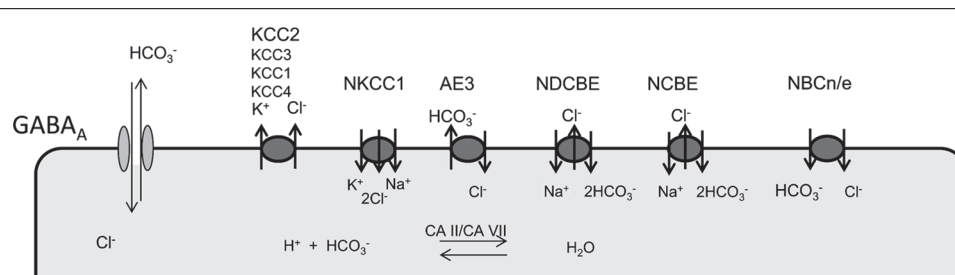
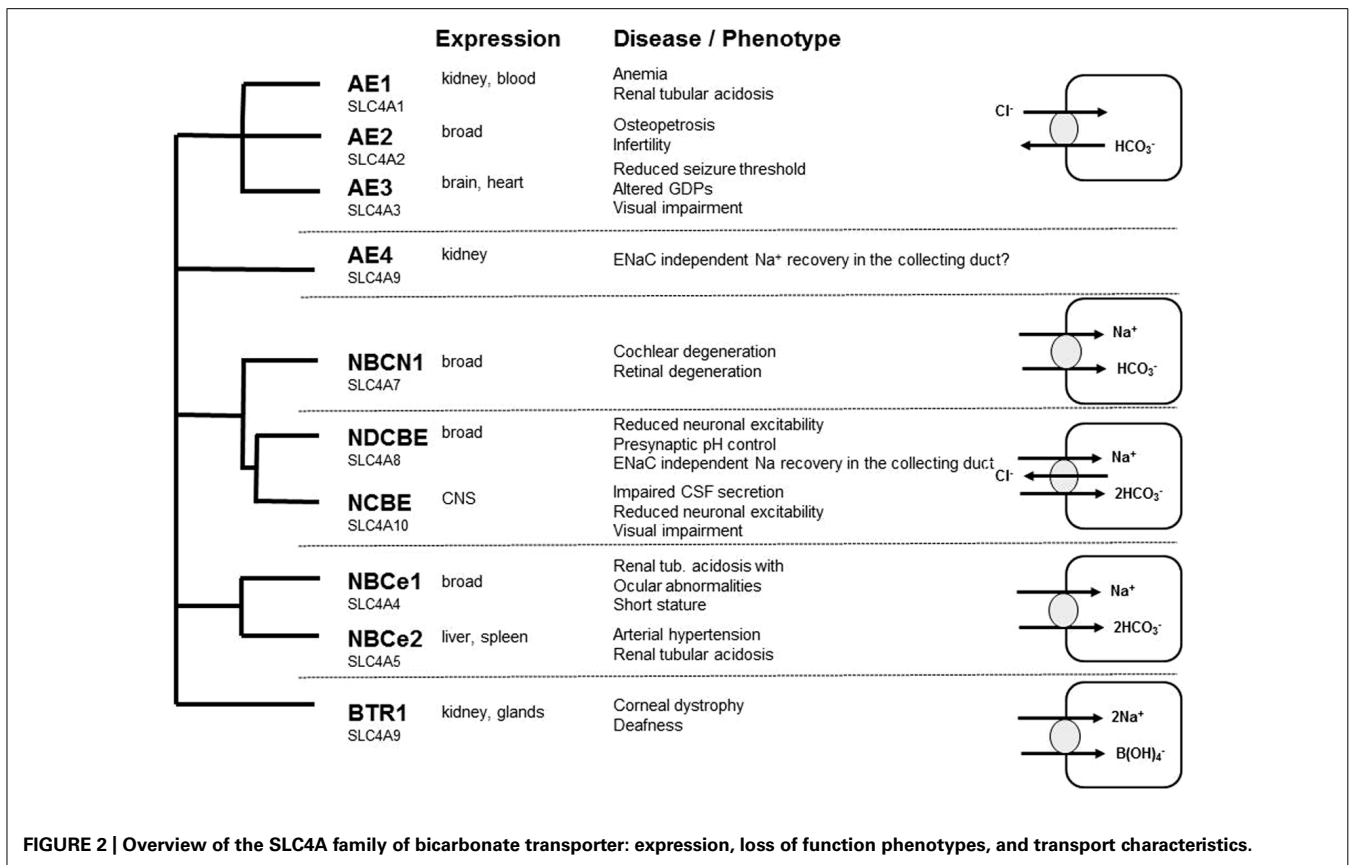


FIGURE 1 | Ion transporters involved in the regulation of neuronal chloride and bicarbonate levels. GABA receptors are permeable for both chloride and bicarbonate. Several anion transporters are expressed in neurons and may thus also affect GABA signaling. Whereas NKCC1 is the main chloride accumulating transporter in most neurons, KCC2 is the most important chloride extruder. Anion-transporters of the SLC4A family of

bicarbonate transporters can be sub-classified in Na⁺ dependent (NDCBE, NCBE) and Na⁺ independent anion exchangers (AE1,2,3). According to the stoichiometry Na⁺-coupled bicarbonate co-transporters can be either electroneutral (NBCn) or electrogenic (NBCe). Although carbonic anhydrases cannot change the existing bicarbonate gradients, they promote anion-transport by members of the SLC4A family and replenish bicarbonate levels.



AE3, NCBE, and NDCBE which are strongly expressed in the brain.

AE3

In nervous tissue, the AE3 transporter has been localized to neurons (Kopito et al., 1989; Raley-Susman et al., 1993), and to Muller cells and horizontal neurons within the retina (Kobayashi et al., 1994). In many neurons, anion-exchange is thought to be mainly mediated by AE3 (Kopito et al., 1989; Hentschke et al., 2006; Romero et al., 2013). In contrast, AE1 plays an important role for bicarbonate recovery of intercalated cells in the kidney and for red blood cells, where it is one of the most abundant proteins of the plasma membrane (band3). Accordingly, mutations in AE1 can cause renal tubular acidosis and/or hemolytic anemia (Alper, 2009). AE2 is the most widely expressed anion-exchanger, which localizes to the basolateral side in most epithelial cells (Romero et al., 2013). Like AE1 it appears to play a minor role for the control of the neuronal pH. The *SLC4A3* gene encoding AE3 employs two different promoters to generate the brain and the cardiac variant, the latter having a shorter amino-terminal amino acid sequence. Because of its broad neuronal expression, the brain variant of AE3 is also often referred to as the neuronal anion-exchanger. Transcripts were already detected at early developmental stages of murine brain development starting around E11 (Hentschke et al., 2006). Because of its early expression and its predicted role to raise the intracellular chloride concentration above the electrochemical equilibrium, AE3 may also contribute to early

GABAergic excitation. In particular, it has been hypothesized that AE3 is responsible for chloride accumulation in lateral superior olivary neurons at P0–P3 (Becker et al., 2003). During this time window, these neurons express AE3 but not NKCC1 and depolarize in response to glycine (Balakrishnan et al., 2003). Surprisingly, however, anion-exchange was nearly absent from cultured fetal neurons, although AE3 mRNA was found in both fetal and adult hippocampal neurons (Raley-Susman et al., 1993).

Interestingly, AE3 expression levels in cultured hippocampal neurons from rat increased during long-term exposure to ammonia and caused an ammonia induced increase of the intracellular chloride concentration (Irie et al., 1998), thus supporting a role of AE3 for the regulation of the intraneuronal chloride concentration. At the protein level, a clear band corresponding to AE3, which was absent from knockout tissues, was detected in murine P1 brain lysates with increasing signal intensities at P5 and P15 (Pfeffer et al., 2009). Unfortunately, the subcellular localization of AE3 in the brain is still unclear, because no antibody has been reported that reliably detects endogenous AE3 in brain sections. The GABA reversal potential and GABA-evoked Ca²⁺ responses of CA1 neurons of AE3 knockout mice did not differ between AE3 knockout and WT mice at P1 (Pfeffer et al., 2009), suggesting that in this type of neuron at this time point chloride accumulation by AE3 may be marginal compared to NKCC1. Nevertheless, this may change with increasing expression levels of AE3 during brain maturation. Supporting that AE3 modulates GABAergic transmission, similar to NKCC1 knockout mice GDPs, which largely depend on

a depolarizing action of GABA (Leinekugel et al., 1997; Ben-Ari et al., 2007), were reduced in terms of frequency and amplitudes at postnatal day 5 in AE3 knockout mice (Figure 3; Pfeffer et al., 2009), but these changes may also be related to changes in neuronal pH homeostasis. Although the intraneuronal pH at steady-state conditions in principal neurons of the adult mouse hippocampus did not differ between genotypes, the recovery from an alkaline load was drastically reduced in neurons devoid of AE3 (Hentschke et al., 2006). Hence, the role of AE3 for chloride accumulation in hippocampal neurons should be re-addressed at later developmental stages and in different types of neurons. Indeed, in spinal cord motoneurons chloride accumulation was in part bicarbonate-dependent and sensitive to anion-exchange blockers (Gonzalez-Islas et al., 2009). These findings are in accordance with a previous report on GABA currents in embryonic motoneurons, which were dampened by bumetanide and removal of extracellular bicarbonate (Kulik et al., 2000). It has been estimated that NKCC1 is responsible for approximately two-thirds of the steady-state chloride accumulation, whereas AE3 for the remaining third (Gonzalez-Islas et al., 2009). NKCC1 and AE3 may thus have distinct functions in the recovery of chloride levels following chloride depletion in embryonic motoneurons.

Overall, no obvious behavioral or morphological alterations of the brain of AE3 knockout mice have been reported (Hentschke et al., 2006; Alvarez et al., 2007). Notably, the seizure threshold in response to various proconvulsive agents was significantly reduced upon disruption of AE3 (Hentschke et al., 2006). This observation

supports a previous report that a susceptibility locus for common idiopathic generalized epilepsy maps to chromosomal region 2q36 (Sander et al., 2002), which also includes *SLC4A3*. Indeed, in a subsequent study a common polymorphism within the coding sequence of *SLC4A3*, which entails the amino acid exchange Ala867Asp, was associated with an increased risk to develop idiopathic generalized epilepsy (Sander et al., 2002). Moreover, the Ala867Asp variant had a significantly reduced anion-exchange activity compared to wild-type in a heterologous expression system, whereas differences in expression levels or protein trafficking to the plasma membrane were excluded (Vilas et al., 2009). Nevertheless, it is still unclear whether the above-mentioned polymorphism 867Asp itself confers the increased risk for epileptic seizures or another gene in close proximity of *SLC4A3* is involved.

Inner retina defects with late onset photoreceptor degeneration with optic nerve and retinal vessel anomalies, which resulted in reduction of the b-wave in electroretinograms, were noted in an independent AE3 knockout mouse (Alvarez et al., 2007). In the retina, the brain variant of AE3 localized to Müller cells, whereas the cardiac variant was detected in horizontal cells. Immuno-labeling of astrocytes showed that inner retina vessels were wrapped by dense astrocytic processes at 8 months of age in AE3 knockout mice. Moreover, inner retina blood vessels formed sporadic loops in the knockout, a finding which was not observed in wild-type mice. Immunoblotting analysis revealed that the $\text{Na}^+/\text{HCO}_3^-$ co-transporter (NBC1), and carbonic anhydrases

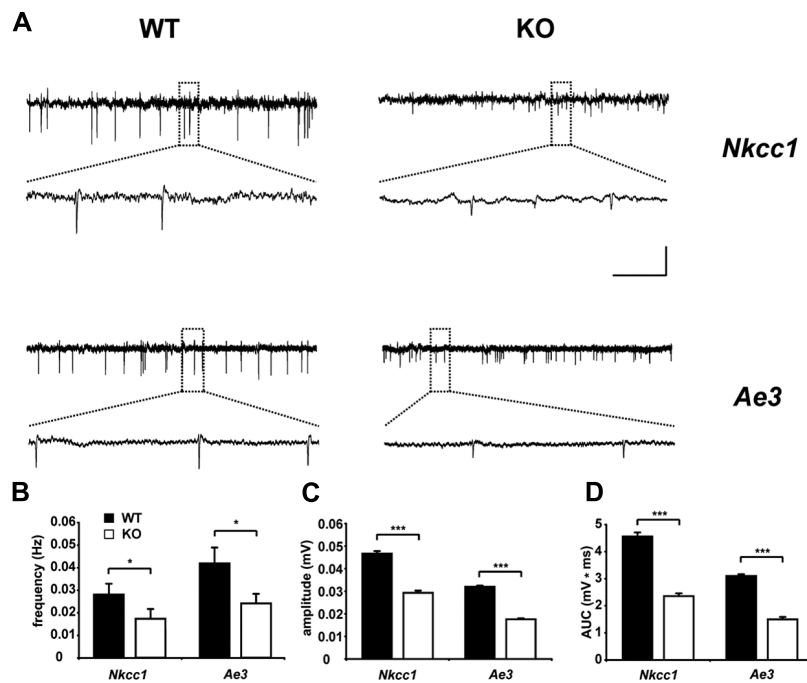


FIGURE 3 | Reduced spontaneous electrical activity in hippocampal slices of NKCC1 and AE3 knockout mice. (A) Representative extracellular recordings from the stratum pyramidale (CA3) of postnatal day 5 NKCC1 and AE3 WT and KO slices. The framed parts are shown as enlargements below the original trace. Calibration: horizontal, 2 min (original trace), 7.5 s

(enlargement); vertical, 0.04 mV. Quantification of frequency (B), amplitude (C), area under curve (AUC), and (D) of single spontaneous electrical events. The asterisks indicate significant difference (* $p < 0.05$, *** $p < 0.001$, t -test). Error bars indicate SEM. Reprinted from Pfeffer et al. (2009).

(CAs) II and CA XIV protein expression were elevated in AE3 knockout mice mouse retinas, suggesting a partial compensation for loss of AE3. Anion-exchange activity mediated by AE3 is promoted by the action of extracellular CAs (Svichar et al., 2009). AE3 associates with the CAs and forms a bicarbonate transport metabolon to maximize bicarbonate fluxes across the plasma membrane (Casey et al., 2009).

Sodium-coupled anion exchangers

Historically, sodium-dependent anion-exchange (NDAE) which extrudes chloride from cells was the first acid–base transport mechanism described to play a role in the control of intracellular pH (Boron and De Weer, 1976). A cDNA encoding a protein that mediates NDAE (also referred to as NDCBE) was initially cloned from *Drosophila* (Romero et al., 2000). The mRNA was expressed throughout *Drosophila* development with a prominent signal in the central nervous system and its disruption resulted in a lethal phenotype in *Drosophila*. A related cDNA coding for another protein mediating NCBE was cloned from a mouse insulinoma cell line (Wang et al., 2000). This initial transport characterization was subsequently confirmed for rat (Giffard et al., 2003; Damkier et al., 2010), whereas the human cDNA was rather characterized as an electroneutral $\text{Na}^+/\text{HCO}_3^-$ cotransporter (NBCn2) with chloride self-exchange activity (Parker et al., 2008b). Some of the controversy may be explained by the different expression systems used in the different studies like mammalian cells and *Xenopus* oocytes, temperature, and composition of solutions, the transfection/injection efficiency or molecular tagging of the transport proteins.

Sodium-coupled anion exchange is activated by intracellular acidification (Schwiening and Boron, 1994), suggesting that regulation of the chloride gradient by NDAEs may be closely linked to the regulation of cellular pH. As prolonged neuronal activity can cause neuronal acidification by efflux of bicarbonate through GABA_A receptors (Kaila and Voipio, 1987), sodium-coupled anion exchange may help to maintain a hyperpolarizing chloride reversal potential and thus promote the inhibitory action of GABA. Thus activation of sodium-coupled anion exchange by acidosis may also contribute to seizure termination by promoting a more negative chloride reversal potential and thus promoting the inhibitory effects of GABA.

NDCBE. Several transcript variants have been reported for human and murine NDCBE. The functional comparison of the NDCBE variants expressed in *Xenopus* oocytes demonstrated that the variants with a shorter C-terminus had a reduced functional expression, whereas the different amino termini did not affect the basal functional expression of NDCBE (Parker et al., 2008a). NDCBE is encoded by *SLC4A8* and is broadly expressed in different tissues including brain (Romero et al., 2004). A down-regulation of NDCBE protein expression was shown in different brain regions after chronic hypoxia with a different profile in neonates and adult mice (Chen et al., 2008a). Immunoreactivity for NDCBE was detected in different brain regions with no overlap to astrocyte markers (Chen et al., 2008b). This was also confirmed in a knockout controlled study with an independent polyclonal antibody against NDCBE. Moreover, this study reported that

NDCBE localization overlapped with markers of presynaptic glutamatergic but not GABAergic nerve terminals (Sinning et al., 2011). From Western analysis of different brain lysate subfractions and immunogold electron microscopy studies on isolated synaptosomes, it was further concluded that NDCBE is enriched in presynaptic nerve endings of excitatory neurons. The localization in presynaptic glutamatergic terminals was also shown in an independent study, but in contrast to the previous report the latter study also detected NDCBE in terminals of parvalbumin-positive GABAergic cells (Burette et al., 2012). Hence, the authors speculated that NDCBE may play a role as a regulator of GABAergic neurotransmission.

Confirming the important role of NDCBE for pH regulation in neurons, its disruption caused a sustained decrease of the steady-state pH of cultured hippocampal neurons (Sinning et al., 2011). In accordance with the observation that NDCBE co-localizes with presynaptic glutamatergic nerve terminals, the frequency of miniature excitatory postsynaptic currents (mEPSCs) was drastically reduced in a pH-dependent manner in hippocampal neurons of mice devoid of NDCBE, whereas miniature inhibitory postsynaptic currents (mIPSCs) were unchanged. Importantly, the effect on mEPSCs could be at least in part restored by shifting the pH, strongly arguing against a structural defect (Sinning et al., 2011). Its role during early brain maturation, however, has not been studied.

Whether NDCBE also significantly contributes to the control of the intraneuronal chloride concentration in some neurons is still unclear. It was observed that dopaminergic neurons in the rat substantia nigra do not express KCC2, but still exhibit inhibitory responses to GABA that are dependent upon the presence of extracellular bicarbonate (Gulacsi et al., 2003). As the GABA reversal potential was significantly less negative in bicarbonate-free buffer in dopaminergic neurons, a sodium-dependent anion might substitute KCC2 in this type of neuron. This assumption was also supported by the demonstration that complex-spike activity in some auditory interneurons results in a pH-dependent negative shift of the glycine reversal potential, and it was suggested that sodium-coupled anion exchange via *Slc4a8* may account for the reduction of intracellular chloride (Kim and Trussell, 2009).

In *Caenorhabditis elegans* sodium-coupled anion exchange is mediated by ABTS-1. While animals lacking ABTS-1 or KCC2 displayed only mild behavioral defects, disruption of both chloride extruders resulted in a paralytic phenotype (Bellemer et al., 2011). Although direct electrophysiological data were not provided, the authors speculated that the disruption of both transporters results in a reversal of chloride fluxes through GABA_A receptors thus rather exciting than inhibiting cells. Moreover, neuronal expression of both transporters was up-regulated during neuronal differentiation and ABTS-1 expression was increased in mutants devoid of KCC2, suggesting that both transporters are important to control the cellular chloride gradient.

NCBE. For NCBE two different splice variants have been identified with different expression profiles (Giffard et al., 2003). The variant missing a 39-bp insert at the 3' end is predicted to result in a protein with a C-terminal PDZ motif (Giffard et al., 2003). How this might relate to function has not been studied. Our expression

analysis in the developing mouse brain with a probe detecting both transcript variants revealed a broad neuronal expression pattern and a particularly strong labeling of the choroid plexus (Hübner et al., 2004). At the protein level, NCBE localized to the basolateral membrane of choroid plexus epithelial cells (Jacobs et al., 2008). There, NCBE serves as a basolateral sodium entry pathway. According to this model, its disruption is predicted to impair cerebrospinal fluid secretion, which is supported by the finding that mice with a targeted disruption of NCBE display a collapse of their brain ventricles (Jacobs et al., 2008). Immunohistological studies revealed that the NCBE protein mainly localized to dendrites and somata of principal neurons, but not to axons or astrocytes (Chen et al., 2008b; Jacobs et al., 2008). There was also a considerable overlap between GABAergic interneurons as identified by the colocalization of GAD and NCBE (Jacobs et al., 2008). However, to which extent NCBE can be detected in different interneuron subtypes, still remains to be addressed. Notably, the ultrastructural analysis also localized NCBE preferentially to dendrites and spines both in the hippocampus as well as in the cerebellum (Jacobs et al., 2008).

Although there was no difference in the steady-state pH of principal neurons of the CA1 hippocampal region of NCBE knockout mice (Jacobs et al., 2008), the recovery to an acid load was delayed. How this affects network excitability was studied in the 4-aminopyridine model of interictal discharges in acute brain slices. The frequency of the interictal-like events at baseline levels did not differ between genotypes, however, the decreased frequency upon a propionate pulse was prolonged in the knockout. In accordance, knockout mice had an increased seizure threshold in response to different seizure inducing agents including pentylentetrazole or pilocarpine. Quite in contrast to the mouse findings, in some patients with epilepsy larger heterozygous genomic deletions involving *SLC4A10* were described (McMilin et al., 1998), however, the genetic evidence that the epilepsy phenotype is directly linked to the heterozygous loss of NCBE or rather to some other genes within this chromosomal region is obscure. Because of the different sites of NCBE expression both in excitatory and inhibitory neurons as well as in the choroid plexus different effects may add up in the total knockout. Thus, the exact role of NCBE for network excitability and synaptic transmission still needs to be addressed in more specific mouse models.

NCBE is also strongly expressed within the retina, where it localizes to ON and OFF bipolar cell axon terminals and to dendrites of OFF bipolar cells, where it co-localized with the main neuronal chloride extruder KCC2 (Hilgen et al., 2012). NCBE was also expressed in starburst amacrine cells, but was absent from neurons known to depolarize in response to GABA, like horizontal cells. These data suggest that NCBE may indeed contribute to the regulation of intracellular chloride and bicarbonate concentration in retinal neurons. Supporting this assumption, knockout mice displayed a decreased visual acuity and contrast sensitivity in behavioral experiments and smaller b-wave amplitudes and longer latencies in electroretinograms (Hilgen et al., 2012).

CARBONIC ANHYDRASES

In the mature rat hippocampus, intense GABA_A receptor activation causes neuronal excitation which is strictly dependent on the

presence of bicarbonate and suppressed by membrane-permeant inhibitors of CA activity (Staley et al., 1995; Kaila et al., 1997; Fujiwara-Tsukamoto et al., 2007). Fifteen members of the CA family have been identified which differ in tissue distribution and subcellular localization. At least 13 family members catalyze the reversible hydration of CO₂ to form bicarbonate and H⁺, accelerating this spontaneous reaction several thousand-fold. Thereby CAs influence the kinetics and amplitudes of pH transients in distinct intra- and extracellular compartments (Chesler, 2003; Casey et al., 2009) and can affect proton-sensitive membrane proteins involved in neuronal signaling such as GABA_A receptors, NMDA receptors, and many more. CA also associate with anion exchangers to form bicarbonate transport metabolons, which enhance bicarbonate fluxes across the plasma membrane (McMurtrie et al., 2004). By forming isoform-specific metabolons with distinct acid-base transporters intracellular neuronal CAs may contribute to developmentally and spatially distinct pH_i microdomains. In the brain, extracellular space CA activity is due mainly to isoforms CA IV and CA XIV, which both play important roles in the regulation of intracellular pH in hippocampal neurons by facilitating AE3-mediated Cl⁻/HCO₃⁻ exchange (Casey et al., 2009; Svichar et al., 2009).

CA II and CA VII are the only cytosolic isoforms present in both somata and dendrites of mature hippocampal CA1 pyramidal neurons. The functional expression of CA VII in mouse brain starts around postnatal day 10 (P10) and that of CA II around P20 and coincides with the appearance of bicarbonate-dependent high frequency stimulation (HFS)-induced tonic GABAergic excitation (Ruusuvuori et al., 2004, 2013). Synchronous neuronal activity in the form of GDPs, however, starts much earlier with an onset at approximately P0 (Ben-Ari et al., 1989) and is largely independent from the presence of bicarbonate (Ruusuvuori et al., 2004). These GDPs disappear with the on-going expression of the chloride extruding K⁺-Cl⁻ cotransporter KCC2 which is up-regulated from P0 to P12 thus rendering GABA_A responses hyperpolarizing (Rivera et al., 1999; Hübner et al., 2001; Stein et al., 2004). To study the role for cytoplasmic neuronal CAs for bicarbonate-dependent GABAergic depolarization, we recently established a CA VII knockout mouse model. Remarkably, CA VII knockout mice have a normal life span and show no gross behavioral abnormalities. At P13–14, when CA II is not yet expressed, CA VII KO mice show a complete absence of electrographic seizures (Ruusuvuori et al., 2013). These results point to a crucial role for the developmental expression of intrapyramidal CAs in shaping integrative functions, long-term plasticity and susceptibility to epileptogenesis and put intraneuronal CA in a key position in GABAergic excitation (Kaila et al., 1997; Ruusuvuori et al., 2004). Moreover, these observations give important insights into the antiepileptic actions of CA inhibitors.

CONCLUSION

There is ample evidence that brain development and brain function critically depends on anion gradients. Whereas chloride has been in the focus of the neuroscientific community, much less is known about bicarbonate. With the development of several mouse models with targeted disruption of selected players of intraneuronal bicarbonate levels, some described in this review, first

clues how bicarbonate contributes to proper brain function like the production of the cerebrospinal fluid, neuronal excitability, and synaptic transmission evolved. The role of these processes for brain development is mostly unknown but it emerges that bicarbonate transporters modulate GABAergic transmission already in the developing brain. It will be essential to assess whether this reflects secondary effects in response to changes in pH or whether these effects rather reflect alterations of the existing anion gradients. Bicarbonate definitely plays an essential role for the GABAergic excitation observed upon massive GABAergic stimulation. This process is massively enhanced by CAs, which quickly replenish intraneuronal bicarbonate from P18

onwards. To understand the complex interplay of the different proteins in time and space is an emerging challenge for the future.

ACKNOWLEDGMENTS

No financial, commercial, or otherwise relationships that might be perceived by the academic community as representing a potential conflict of interest exist. This work was supported by a grant from the Interdisciplinary Centre for Clinical Research Jena to Christian A. Hübner and Knut Holthoff, funds of the DFG and the BMBF (01GQ0923) to Christian A. Hübner and Knut Holthoff. We thank Knut Kirmse for helpful discussions.

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Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Received: 25 July 2013; accepted: 21 September 2013; published online: 24 October 2013.

Citation: Hübner CA and Holthoff K (2013) Anion transport and GABA signaling. *Front. Cell. Neurosci.* 7:177. doi: 10.3389/fncel.2013.00177

This article was submitted to the journal *Frontiers in Cellular Neuroscience*.

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