

Molecular components and functionality of the GABA signaling during
development: a study on two model systems

Ph.D. Dissertation

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To my dear parents

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II LIST OF ABBREVIATIONS

AAH	artificial aqueous humor
ALP	alkaline phosphatase
AMPA	α -amino-3-OH-5-methylisoxazole-4-propionic acid
AMPA-R	AMPA-gated glutamate receptor channel
DAB	3,3'-diaminobenzidine
CNS	central nervous system
CP	cortical plate
E	embryonic day
EB	embryoid body
EGAD	embryonic glutamic acid decarboxylase
EPSC, EPSP	excitatory postsynaptic current, potential
ES cells	embryonic stem cells
FBS	fetal bovine serum
GABA	γ -aminobutyric acid
GABA _A receptor	GABA-gated receptor channel (type A)
GABA _B receptor	GABA-gated receptor channel (type B)
GABA _C receptor	GABA-gated receptor channel (type C)
GAD	Glutamic acid decarboxylase
GAT (1-4)	GABA transporters type 1-4
HEPES	<i>N</i> -hydroxyethylpiperazine- <i>N'</i> -2-ethanesulfonic acid
IP3	Inositol-3-phosphate
iPS	induced pluripotent cells
IZ	intermediate zone
K ⁺	potassium ion

KCC1-3	potassium chloride co-transporter type1-3
LVA Ca ²⁺ channels	low-voltage activated calcium channels
MZ	mantle zone
Na ⁺	sodium ion
NKCC1-3	Sodium potassium chloride co-transporter type 1-3
NMDA	<i>N</i> -methyl-d-aspartate
NMDA-R	NMDA-gated glutamate receptor channel
PAF	paraform-aldehyde
PBS	phosphate buffer saline
PCR	polymerase chain reaction
PSC	postsynaptic current
RT	room temperature
RT-PCR	reverse transcribed polymerase chain reaction
RYR	Ryanodine receptor
SB	standard buffer
SDS-PAGE	sodium dodecylsulphate-polyacrylamide gel electrophoresis
SVZ	subventricular zone
TBS	Tris buffered saline
VGAT	vesicular GABA transporter
VGCC	voltage-gated calcium channels
VZ	ventricular zone
Wt	wild type

“New idea: at first, they said, it can’t be true; when it became obvious it was true, they said it wasn’t important; and when its importance couldn’t be doubted, they said, it wasn’t new!”

Adopted from “How does a little acronym become a big transmitter?”, Krešimir Krnjević, 2006

III INTRODUCTION



1. γ - AMINO BUTYRIC ACID (GABA) SIGNALING

γ -aminobutyric acid (GABA) is a nonprotein amino acid that is conserved from bacteria through yeast and plants to mammals where it can exert diverse functions (Roberts, 1974; Work and Dewey, 1953; Marcus and Halpern, 1969; Bouche et al., 2003; Bouche and Fromm, 2004).

In the adult vertebrate central nervous system (CNS), GABA is mostly recognized as the principal inhibitory neurotransmitter and as such plays a key role in modulating neuronal excitability. It is estimated that, depending on the brain region, 20 to 50 % of all central synapses use GABA as their neurotransmitter (Bloom and Iversen, 1971; Docherty et al., 1985). The heterogeneity in types of GABAergic synapses and interneurons revealed in the last decade (Freund and Buzsaki, 1996; Parra et al., 1998; Cossart et al., 2006; Dumitriu et al., 2007) indicates that an array of GABAergic signaling functions might exist in the brain. GABA is directly involved in a variety of biological functions such as locomotor activity, learning, reproduction, circadian rhythms as well as cognition, emotions and sleep (Agmo et al., 1987; Paulsen and Moser, 1998; Mintz et al., 2002; Leventhal et al., 2003; Terasawa, 2005).

Furthermore, in the immature nervous system GABA serves as excitatory neurotransmitter and acting in a paracrine way (reviewed in Owens and Kriegstein, 2002a, 2000b; Ben Ari et al., 2002, 2007). Membrane and intracellular calcium rise

induced by GABA-mediated depolarization, activates second messenger-intracellular cascades involved in neuronal migration, growth, differentiation, plasticity of synaptic connections and establishment and patterning of neural networks (Owens and Kriegstein, 2002a, 2002b; Ben-Ari et al., 2002, 2007).

Proper function of the GABAergic system is crucial for the performing of higher brain functions. Various psychiatric and neurological disorders such as anxiety, depression, epilepsy, schizophrenia, and autism are partly caused by dysfunction of the GABAergic transmission and imbalanced excitation vs inhibition in the developing and/or mature brain (Wong et al., 2003).

Besides its function in the CNS, GABA has been also suggested to be involved in physiology of glia cells, digestive tract, pancreas, liver, osteoblasts, and chondrocytes (reviewed in Watanabe et al., 2002) and in modulating the proliferation of various cancer cells (reviewed in Watanabe et al., 2006).

1.1. Basic properties of GABA signaling

Molecular components and GABA neurotransmitter system is summarized in *Fig.1*. In the mammalian brain, **GABA** is synthesized primarily from glutamate in a reaction that is catalyzed by two glutamic acid decarboxylase (**GAD**) enzymes, GAD65 and GAD67. GABA is loaded into synaptic vesicles by a vesicular neurotransmitter transporter (**VGAT**) and is released from nerve terminals by calcium-dependent exocytosis. However, nonvesicular forms of GABA secretion (for example, by reversal of the membrane GABA transporter GAT) have also been described and might be particularly important during development (see below). The effects of GABA can be mediated by the activation of either ionotropic **GABA_A** or metabotropic **GABA_B** receptors, which can be localized both pre- or postsynaptically.

GABA signals are terminated by reuptake of the neurotransmitter into nerve terminals and/or into surrounding glial cells by the plasma membrane GABA transporters (GATs). Subsequently, GABA is metabolized by a transamination reaction that is catalysed by the mitochondrial GABA transaminase (GABA-T).

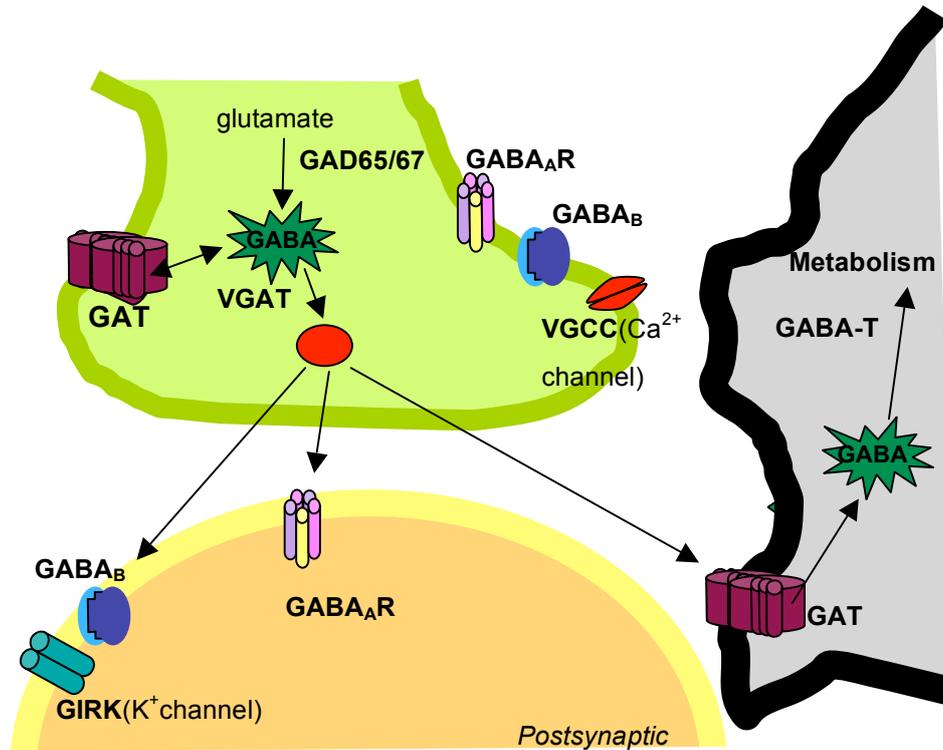


Fig.1. Molecular components of the GABA signaling. Schematic diagram of the synthesis and transport of GABA at synapses. GABA is synthesized primarily from glutamate in a reaction that is catalysed by GAD. GABA can be released either vesicularly (by VGAT) or nonvesicularly (by reverse transport). GABA receptors are located at pre- and postsynaptic sites. Reuptake of GABA by surrounding neurons and glia occurs through the activity of GABA transporters (GATs). GABA is metabolized by GABA transaminase (GABA-T). Modified from *Owens and Kriegstein, 2002*

1.1.1. GABA synthetic enzyme - the glutamic acid decarboxilase (GAD)

1.1.1.1. Two isoforms of GAD: GAD65 and GAD67

Following the successful purification of GAD in the laboratory of E. Roberts (Roberts, 1986), it has become possible to show that this enzyme catalyses the rate-limiting step in GABA synthesis from glutamate and that this is the predominate route for GABA synthesis in the CNS and periphery. These early studies have also shown that GAD is localized in the same neurons that use GABA and since then, it has been used as a morphological marker of GABAergic neurons.

Molecular cloning studies have shown that in the adult vertebrate brain GAD exists in two major isoforms, 65kD (**GAD65**) and 67kD (**GAD67**), which are products of two independently regulated genes located on chromosome 2 and 10, in humans and in the mouse, respectively (Soghomonian and Martin, 1998). Each of these forms is highly conserved among vertebrates and shares more than 95% identity at the amino acid level among different species such as rat, mouse, cat, and human (Kaufman et al., 1986; Kobayashi et al., 1987; Katarova et al., 1990; Wyborski et al., 1990; Erlander et al., 1991). In the same species, the two GAD proteins differ in amino acid sequence (with 65% identity in the rat), molecular sizes (65 kD and 67 kD), regional distribution in the brain, subcellular localization and regulation. Both forms are composed of two functional domains: a highly divergent N-terminal domain (with 23% identity between the two human GADs) that is involved in subcellular targeting, membrane association and heteromeric interactions, and a larger, more conserved (73% identity) C-terminal domain that contains the catalytic centre (Soghomonian and Martin, 1998; Varju et al., 2001). Experiments with recombinant GAD suggested that GAD65 is more responsive than GAD67 to the presence of the cofactor pyridoxal-P (PLP) (Erlander et al., 1991; Kaufman et al., 1991). This is consistent with the finding that the majority of apo-GAD (form without cofactor PLP) in brain is provided by GAD65 (Martin et al., 1991). Taking into account also the enrichment of GAD65 compared to GAD67 at synaptic vesicles of nerve terminals and its specific pattern of expression, GAD65 is thought to be preferentially responsible for the establishment of the vesicular GABA-transmitter pool. Thus this form is predominantly associated with stimuli-evoked, largely PLP-dependent inhibitory synaptic transmission. In contrast, GAD67 makes up most of the constitutively active holo-GAD (PLP bound GAD) pool and is enriched in the cytoplasm. The non-vesicular cytoplasmic GABA pool synthesized mostly by GAD67 is involved in tonic inhibition, which is less dependent on PLP availability

(Erlander et al., 1991; Martin and Rimvall, 1993). This form is mostly responsible for the synthesis of non-synaptic GABA pool that has metabolic functions (Martin and Rimvall, 1993; Esklapez et al., 1994; Soghomonian and Martin, 1998).

1.1.1.2. GABA metabolism

GABA is mainly metabolized through a short pathway called the “GABA shunt”, a closed loop, which acts in such a way as to secure the supply of GABA (Bouche et al., 2003; **Fig.2**). The first step in the shunt is the transamination of α -ketoglutarate to glutamic acid. Glutamic acid is then decarboxylated to form GABA. GABA is then transaminated to form succinic semialdehyde, but the transamination can only take place if α -ketoglutarate is the acceptor of the amino group. This transforms α -ketoglutarate into the GABA precursor, glutamate, thereby guaranteeing a continuity of supply. Thus, a molecule of GABA can only be degraded metabolically if a molecule of precursor glutamate is formed in its place. The succinid semialdehyde formed from GABA is rapidly oxidized to succinic acid, which is a normal constituent of the Krebs cycle.

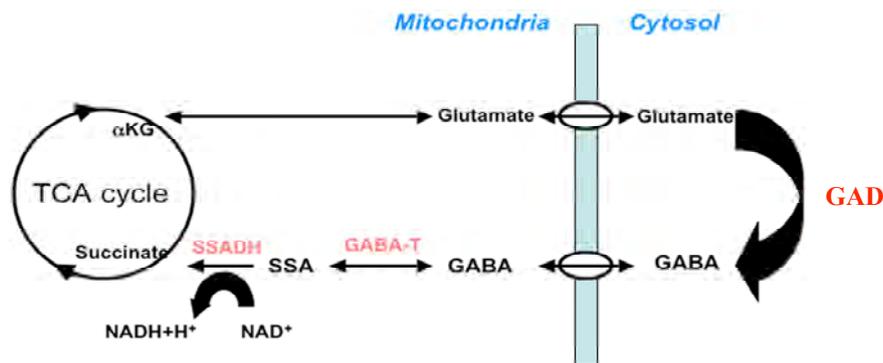


Fig.1. Schematic representation of the GABA shunt metabolic pathway. The GABA shunt is composed of three enzymes (red). Glutamic acid decarboxylase (**GAD**) is a cytosolic enzyme that catalyzes the irreversible decarboxylation of glutamate to produce GABA. GABA is transported into the mitochondria where it is converted into succinic semialdehyde (**SSA**) by GABA transaminase (**GABA-T**). **SSA** is then reduced by succinic semialdehyde dehydrogenase (**SSADH**) to form succinate, which enters the tricarboxylic acid (TCA) cycle. α -KG, α -ketoglutarate

Extensive studies on brain GABA have revealed the existence of two distinct GABA pools – synaptosomal, mediating synaptic inhibition and cellular, with a presumed

metabolic function (Martin and Rimvall. 1993; Martin and Barke, 1998). Roughly, less than 50% of neuronal GABA is engaged in the synaptosomal pool, consequently, there is a fairly large cytoplasmic GABA pool that could either be involved in a different functions, related to extrasynaptic cell-cell signaling or can serve as a reserve.

Finally, change in GABA homeostasis can have severe consequences on development of both humans and plants (reviewed in Bouche et al., 2003). For example, in humans, disruption of SSADH or GABA-T causes severe clinical manifestations associated with the accumulation of neurotransmitters in physiological fluids (Chambliss et al., 1998; Medina-Kauwe et al., 1999).

1.1.1.3. Alternatively spliced truncated GAD embryonic forms

During embryonic development, two additional truncated GAD proteins: GAD25 and GAD44 are also expressed, which are the products of alternatively spliced GAD transcripts encoded by the GAD67 gene.

The two distinct embryonic GAD transcripts include two almost identical alternatively spliced exons inserted into the coding region of GAD67 (**Fig.3**; Bond et al., 1990; Szabo et al., 1994). The bicistronic I-80 mRNA contains the embryonic exon “7/A” (filled boxes, **Fig.3**) with an overlapping translational stop-start codon (TGATG) and codes for both truncated enzymatically inactive GAD25 and the enzymatically active 44kD (**GAD44**) form, whereas I-86 mRNA containing the 86bp long exon “7/B” including an additional stop codon at the 3'-end codes only for GAD25 as the GAD44 open reading frame is interrupted in this message (**Fig.3**). GAD25 corresponds to the N-terminal regulatory domain of GAD67 and has been proposed to play mainly a regulatory role. GAD44 is roughly identical to the C-terminal catalytic domain of GAD67 and can synthesize GABA with putative developmental role (Szabo et al., 1994; reviewed in Varju et al., 2001).

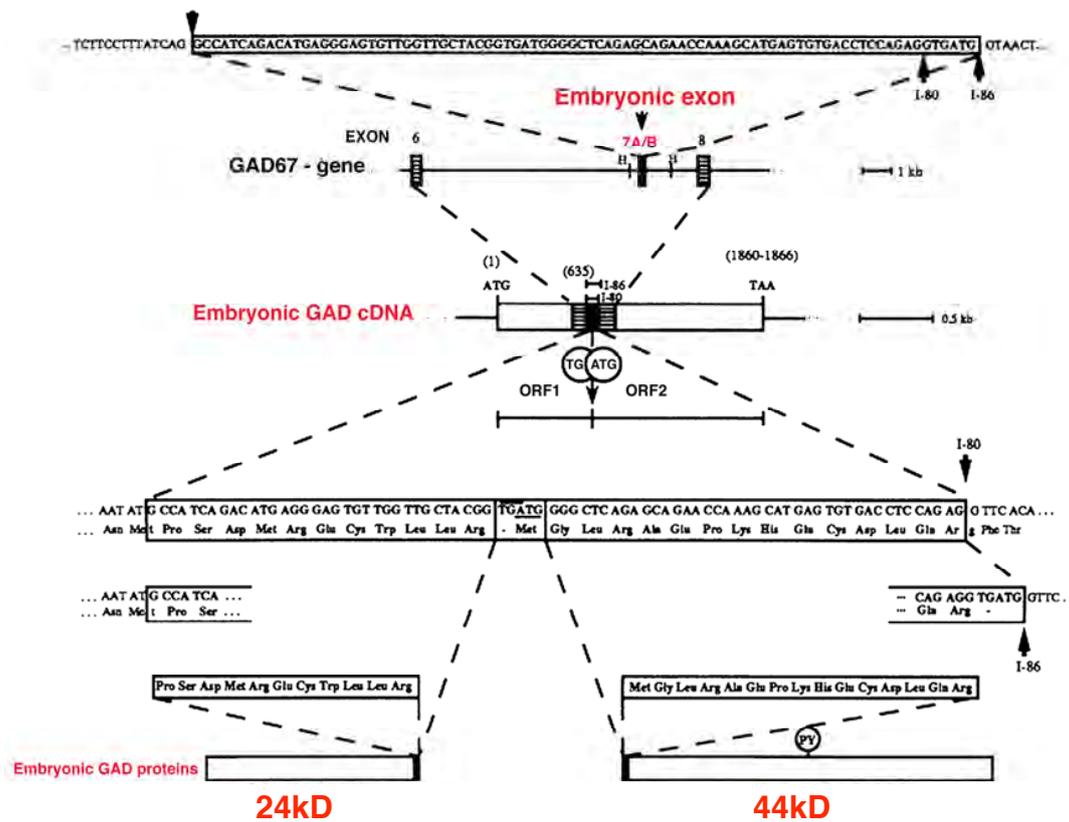


Fig.3. Schematic representation of the embryonic exon on GAD67 genomic and cDNA clones and the resulting putative protein products. GAD67 mRNA is rendered structurally and functionally bicistronic through the developmentally regulated alternative splicing of a single exon (7). The embryonic exon-derived TGATG overlapping codon splits the open reading frame of GAD67 into two (ORF1,2). Exons 6 and 8 are represented in striped boxes; The overlapping stop-start codon is circled. The corresponding amino acid sequences are given in a three letter code. Modified from Szabo *et al.*, 1994.

1.1.1.4. Upstream regulators of GAD genes: the Dlx gene family

The *Drosophila melanogaster* *Distal-less* (*Dll*) gene codes for a homeodomain protein (Cohen *et al.*, 1989) that is required for the proper development of the limb and for correct proximo-distal (PD) organization of the embryo (Cohen and Jurgens, 1989; Cohen *et al.*, 1989). Distal-less related genes named Dlx in vertebrates code for transcription factors that show a high degree of sequence similarity in their DNA-binding domains and the homeodomains (Liu *et al.*, 1997; Zhang *et al.*, 1997). Dlx genes found in mice and man are found linked in a paired tandem convergent configuration, in a following order: Dlx1 and Dlx2, Dlx5 and Dlx6, Dlx3 and Dlx7, respectively. (Bulfone *et al.*, 1993; Simeone *et al.*, 1994; Stock *et al.*, 1996; Anderson

et al., 1997a; Liu et al., 1997; Panganiban and Rubenstein, 2002; Zerucha et al., 2000). Recent work demonstrated that Dlx genes are expressed in spatially and temporally restricted patterns in craniofacial primordia, basal telencephalon and diencephalon, and in distal regions of extending appendages, including the limb and the genital bud (Merlo et al., 2000).

The expression pattern of Dlx genes in embryonic forebrain and some non-neuronal tissues is nearly identical to that of the GAD65 and GAD67 (Thomas et al., 1997; Katarova et al., 2000; Maddox and Condie, 2001; Depew et al., 2005; Tamayama et al., 2005; Levi et al., 2006). Forced expression of Dlx2 or Dlx5, but not Dlx1, in slice cultures of mouse embryonic cerebral cortex induced ectopic expression of GAD65 and GAD67 (Stühmer et al., 2002a) suggesting that Dlx2/5 are directly involved in activation of GAD genes and subsequently, in the establishment of the GABAergic phenotype (Stühmer et al., 2002a, 2002b).

1.1.2. Vesicular and membrane GABA transporters: GABA release and uptake

Following synthesis, GABA is loaded into synaptic vesicles by the vesicular GABA transporter (VGAT) directed by a Na⁺-independent proton-electrical gradient, which is generated at the cost of H⁺-ATPase. Upon neuronal stimulation GABA is liberated from the nerve terminals by calcium-dependent exocytosis (McIntire et al., 1997; Chaundry et al., 1998; Fon and Edwards, 2001). Similar mechanism is believed to apply to some non-neuronal GABA-synthesizing cells (Reetz et al., 1991; Geigerseder et al., 2003; Gammelsaeter et al., 2004), and indeed actual GABA release has been demonstrated recently in the β -cells of the pancreas (Braun et al., 2004).

The magnitude and duration of GABA action is dependent upon an effective mechanism by which the action of the transmitter can be cleared from the site of action. GABA is taken-up from the extracellular space by the membrane GABA-

transporters (GATs), members of the Na⁺- and Cl⁻- dependent transporters specialized in uptake of GABA in exchange for Na⁺ and a Cl⁻-ions. To date four GATs have been cloned, GAT1-4, respectively, which display distinct cellular and regional expression patterns and ionic requirements (Borden, 1996; Conti et al., 2004). **GAT-1** is the most abundant high-affinity neuronal GABA transporter found predominantly on GABAergic neurons and some astrocytes (Uhl, 1992). Beside its expression in the CNS, GAT-1 was described in the testis and sperm of rodents (Hu et al., 2000; Ma et al., 2000). Three other closely related transporters show specific pattern of distribution: **GAT-4** was detected only in the brain, while **GAT-2** and **GAT-3** are present in both the CNS (brain and retina) as well as a number of non-neuronal tissues such as liver and kidney (Liu et al., 1993; Borden, 1996; Jursky and Nelson, 1996, 1999).

Physiological and biochemical studies have demonstrated that GABA can be released by a reversal of GATs, a mechanism predominately utilized by the growing cones of neuronal precursors during synaptogenesis (Pin and Bockaert, 1989; Taylor et al., 1990; Taylor and Gordon-Weeks, 1991; Attwell et al., 1993; Levi and Raiteri, 1993; Wu et al., 2007).

1.1.3. GABA receptors as mediators of GABA action

GABA exerts its action by binding to the two main types of receptors: ionotropic, GABA_A and GABA_C, which mediate fast inhibitory neurotransmission, and metabotropic GABA_B, which is coupled to GTP-binding proteins and mediates slow inhibitory neurotransmission. In the adult brain, GABA_A and GABA_B receptors are abundantly expressed mainly in neurons but also in glia.

1.1.3.1. Ionotropic GABA ($GABA_A$ and $GABA_c$) receptors

Most of the physiological actions of GABA are mediated by $GABA_A$ receptors. Functional $GABA_A$ receptors are heterooligomeric assemblies of distinct subunits that form a membrane channel permeable to chloride, and to a lesser extent to bicarbonate anions (Kaila, 1994). In mammals 19 receptor subunits have been identified that are grouped into 4 subgroups α (α_{1-6}), β (β_{1-3}), γ (γ_{1-3} and the γ substitutes δ , ϵ , π , θ). Additional forms derive from alternatively spliced variants (Barnard et al., 1993; Simon et al., 2004). All subunits are similar in size, containing roughly 450-550 amino acids and are strongly conserved among species. A high percentage of sequence identity (70-80%) is found between subunit isoforms and, to a lesser extent (30%) between subunit families (**Fig.4A**). The common elements of the subunit structure include a large N-terminal hydrophilic domain (containing a signal peptide, a cysteine bridge and glycosylation sites) exposed to the synaptic cleft and four hydrophobic segments named M1 to M4, each composed of approximately 20 amino acids forming four putative membrane-spanning segments (**Fig.4B**). In theory, a large number of various combinations could assemble from the numerous subunits and their splice variants. However, studies reveal that only a limited number of these combinations can be found on the cell surface (Turner, 1993; Fritschy and Mohler, 1995; Sieghart and Sperk, 2002). Most, in not all $GABA_A$ receptors form heteropentameric assemblies containing two α , two β and one γ (or δ , ϵ , π , θ) subunit (**Fig.4C,D**). In mammalian central and peripheral nervous system, activation of $GABA_A$ generates, in general membrane hyperpolarization and a reduction of action potential firing that leads to both phasic inhibitory postsynaptic currents (IPSCs) and inhibitory tonic currents (Kaneda et al., 1995; Nusser and Mody, 2002). The GABA binding sites are located at the α and β subunit interfaces (see **Fig.4C**), whereas receptor specific agonists and antagonists can bind at other locations (Baumann et al.,

2001). Due to the diversity in subunit composition, GABA_A receptor neurotransmission is characterized by high variability of synaptic responses. The kinetics and pharmacological diversity of GABA_A receptor response is particularly evident during development (Schousboe and Redburn, 1995; Costa, 1998; Cherubini and Conti, 2001).

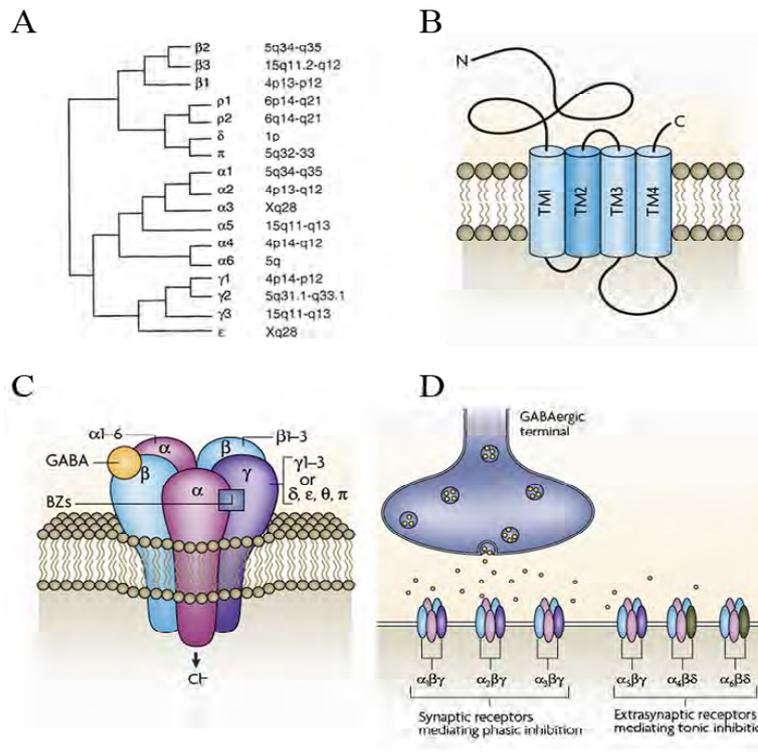


Fig.4. GABA_A receptor structure and localization. **A.** The human GABA_AR gene family. The dendrogram indicates the homologies between deduced amino-acid sequences of subunits. **B.** GABA_AR subunits consist of four hydrophobic transmembrane domains (TM1–4). The large extracellular amino terminus is the site of GABA binding, and also contains binding sites for psychoactive drugs such as benzodiazepines (BZs). **C.** Five subunits from seven subunit subfamilies (α , β , γ , δ , ϵ , θ and π) assemble to form a heteropentameric Cl⁻-permeable channel. Binding of the neurotransmitter GABA occurs at the interface between the α and β subunits and triggers the opening of the channel, allowing the rapid influx of Cl⁻ into the cell. **D.** GABA_ARs composed of $\alpha(1-3)$ subunits together with β and γ subunits are thought to be primarily synaptically localized, whereas $\alpha(4-6)\beta\gamma/\delta$ receptors are located largely at extrasynaptic sites Modified in part from Jacob et al., 2008.

GABA_C receptors are ligand-gated Cl⁻ channels assembled as homo- or heteropentamers from three distinct ρ ($\rho 1-3$) subunits (Bormann, 2000; Zhang et al., 2001). They are found primarily in the vertebrate retina and at restricted sites in the spinal cord and other parts of the CNS (Johnston, 1996; Lukasiewicz, 1996).

1.1.3.2. GABA_B receptor

Metabotropic GABA_B receptors were first described in 1981 as a bicuculline-insensitive, baclofen-sensitive GABA receptor type widely expressed in the mammalian central nervous system (Hill and Bowery, 1981). Functional GABA_B

receptors are heterodimers of GABA_{BR1} and GABA_{BR2} subunits (**Fig.5**; Bowery and Brown, 1997; Kaupmann et al., 1997). Heterodimerization of the two subunits is absolutely critical for efficient receptor trafficking to the cell membrane, formation of ligand-binding sites, and coupling to downstream effectors (Jones et al., 1998). Deletion of either of the subunits results in a complete loss of GABA_B receptor functions (Couve et al., 2000,2004; Bowery, 2006; Mombereau et al., 2004)

The cloning of GABA_{BR1} revealed the existence of several additional alternatively spliced forms in rodents and human (Couve et al., 1998). The first two initially discovered GABA_B receptor isoforms were termed GABA_{BR1a} and GABA_{BR1b} with molecular weights of 130 and 100kDa, respectively. These two isoforms differ by the insertion of a pair of tandem extracellular “Sushi” domains at the N-terminus of GABA_{BR1}, which have been previously shown to be involved in protein-protein interactions (Hawrot et al., 1998). Whereas the specific function of these different isoforms is unclear, they may provide a mechanism for differential targeting of GABA_B receptor within the cell or the coupling of these receptors to different effector systems (Vigot et al., 2006).

With respect to native receptors in the brain, GABA_B receptors can be separated into a postsynaptic population that controls postsynaptic excitability and a presynaptic population that regulates neurotransmitter release. Postsynaptic GABA_B receptors induce slow inhibitory postsynaptic currents (sIPSP) by gating (or activating of inwardly rectifying potassium channels (GIRK or Kir3-type K⁺ channels), which evokes hyperpolarizes action of the membrane and shunting of excitatory currents (Ong and Kerr, 2000). Presynaptic GABA_B receptors can function as autoreceptors and control GABA release or heteroreceptors that inhibit the release of other neurotransmitters, especially glutamate. These receptors mediate their presynaptic effects mostly by inhibition of high-voltage activated Ca²⁺ channels of N or P/Q-type

or through direct modulation of synaptic vesicle priming (Mintz and Bean, 1993; Poncer et al., 1997; Sakaba and Neher, 2003). Furthermore, activation of GABA_B receptors inhibits cAMP production (Hashimoto and Kuriyama, 1997; Simonds, 1999) predominately through G proteins of the G_{i/o} family (**Fig.5**; Bowery et al., 2000; Odagaki and Koyama, 2001; Bettler et al., 2002).

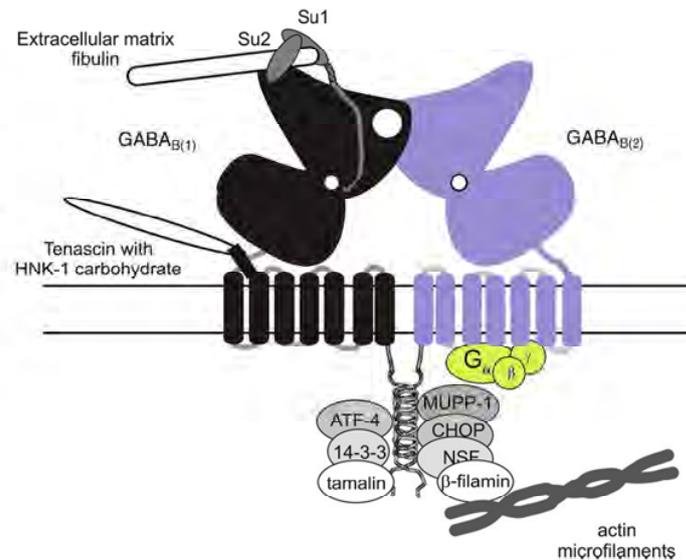


Fig.5. Schematic representation of GABA_B receptor structure. Through the activation of Gi / Go -type G proteins GABA_B receptors couple to pre- and postsynaptic effector systems. A number of proteins were found to interact with the cytoplasmic tails of GABA_B subunits. Among the interacting proteins are leucine-zipper transcription factors (CREB2/ATF-4, CHOP), suggesting that GABA_B receptors mediate long-term metabolic changes involving protein synthesis. Furthermore, a number of scaffolding and adaptor proteins such as 14-3-3, *N*-ethylmaleimide-sensitive factor (NSF), tamalin, and MUPP-1 were found to interact with the coiled-coil domains of either GABA_BR1 and/or GABA_BR2. It is proposed that these proteins regulate receptor dimerization, intracellular trafficking, and synaptic localization. -Filamin, an actin binding protein, is supposed to tether the GABA_B heterodimer to the cytoskeleton via interaction with the GABA_BR2 subunit. The extracellular sushi repeats (Su1 and Su2) of GABA_BR1a are proposed to interact with the extracellular matrix protein fibulin. HNK-1, a carbohydrate carried by extracellular matrix proteins such as tenascin, binds to the extracellular domains of both GABA_B receptor subunits. Modified from Bettler et al., 2004.

1.2. Mechanism of Depolarizing/Excitatory actions of GABA

The inhibitory role of GABA on membrane excitation in the adult brain is reversed in the embryonic nervous system, where GABA exerts mainly an excitatory role (Owens and Kriegstein, 2002; Ben-Ari, 2007). Numerous studies in rodents revealed that activation of GABA_A receptors, in immature neurons results in efflux, rather than influx of chloride ions and subsequently inward electric current producing membrane depolarization (Owens and Kriegstein, 2002; Ben-Ari et al., 2007) generating action

potentials. Activation of GABA_A receptors also increases intracellular calcium concentration ($[Ca^{2+}]_i$) due to the activation of voltage-gated calcium channels (**Fig.6**; Obrietan and van den Pol, 1995), attenuates the voltage-dependent magnesium block of NMDA channels (**Fig.6**; Leinekugel et al., 1997) and interferes with ionotropic glutamatergic transmission (Gao et al., 1998).

1.2.1. Developmental expression of KCC2

Members of cation chloride co-transporter family regulate ion homeostasis through inward or outward directed flux of ions, which depends on electrochemical gradients. The gradient of potassium (K^+) is regulated by the K^+/Cl^- co-transporters (KCC), which extrude Cl^- out of the cell whereas the sodium (Na^+) gradient is regulated by $Na^+/K^+/Cl^-$ co-transporters (NKCC) to accumulate Cl^- within the cell. Until now, four K^+/Cl^- co-transporters (KCC1-4) and two $Na^+/K^+/Cl^-$ co-transporters (NKCC1-2) have been described: **NKCC1** and **KCC1/3/4** are generally expressed in all tissues, **NKCC2** is restricted to kidney, while **KCC2** is only found in neuronal tissue. (Delpire and Mount, 2002; Payne et al., 2003).

The developmental shift of GABA effect on the cells from excitatory to inhibitory is thought to be caused by a decrease in the intracellular Cl^- concentration in developing neurons (**Fig.6**; Rivera et al., 1999; Payne et al., 1996,1997). This is supported by numerous studies that have demonstrated a correlation between the GABA reversal potential and expression level (Lu et al., 1999; Rivera et al., 1999; Coull et al., 2003; Shibata et al., 2004), membrane localization (Balakrishnan et al., 2003), phosphorylation (Kelsch et al., 2001), and/or activity of KCC2 (Woodin et al., 2003). Loss-of-function studies using antisense KCC2 RNA (Rivera et al., 1999), targeted gene knockdown (Hubner et al., 2001; Woo et al., 2002; Zhu and Delpire, 2005) or

over-expressing KCC2 in vitro studies (Lee et al., 2005) have shown that KCC2 is necessary and sufficient for establishing and maintaining a low $[Cl^-]_i$.

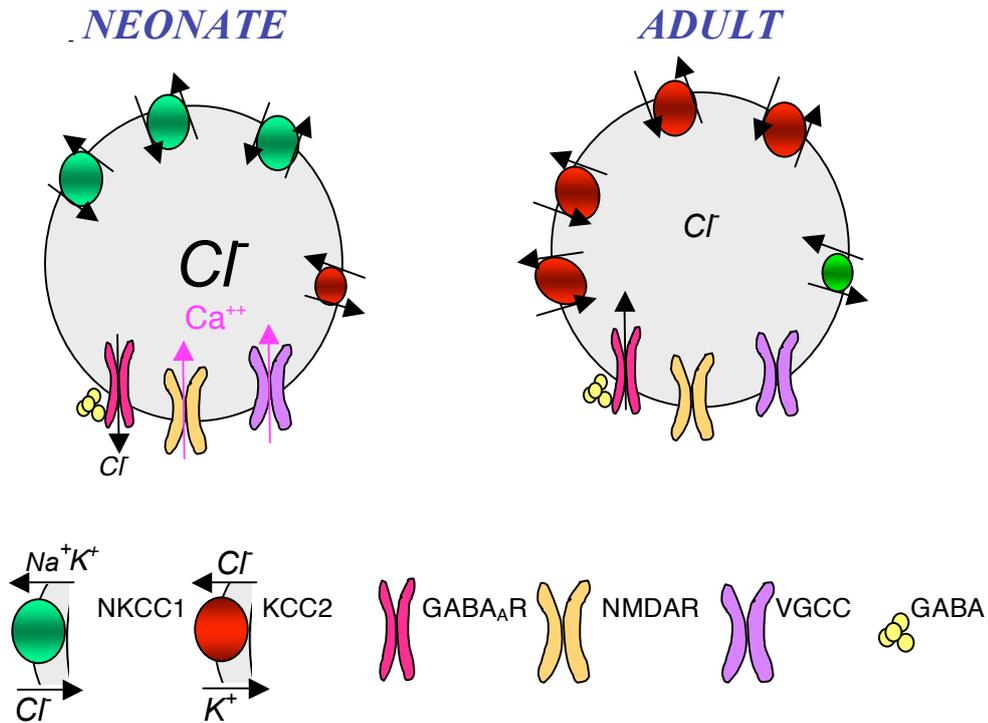


Fig.6. Schematic presentation demonstrating early expression of NKCC1 and late expression of KCC2 determines developmental changes in intracellular Cl^- concentration and in the GABA-induced membrane depolarization/hyperpolarization. Note that NKCC1 expression predominates in immature neurons, in which the intracellular concentration of chloride ($[Cl^-]_i$) is relatively high and sufficient to cause the GABA reversal potential to be either below or above the resting membrane potential and the threshold for action potential generation .

1.3. GABA signaling during development

In addition to being the principal inhibitory neurotransmitter in the adult nervous system, GABA is thought to play a signaling role during development acting in a paracrine/autocrine fashion (Manent et al., 2005; Manent and Represa, 2007) and regulating key developmental steps in the embryonic nervous system, such as proliferation, migration, differentiation, synapse maturation and cell death (LoTurco

et al., 1995; Barker et al. 1998; Katarova et al. 2000;Varju et al., 2000; Owens and Kriegstein, 2002; Ben-Ari, 2002).

1.3.1. Summary of nervous system development

During early development, the layer of **germinal neuroepithelium** consists of a monolayer of rapidly dividing neural stem cells. During the cell cycle, their nuclei undergo “interkinetic nuclear migration” migrating to the pial side of the neural tube during S-phase and back to the luminal side during M-phase (Gilbert, 2000). Extensive proliferation within the germinal neuroepithelium during early developmental stages leads to the generation of two different cell populations forming the **ventricular zone (VZ; Fig.7A)**: radial glia neural stem cells through symmetric, and neuroblasts through asymmetric cell division. The neural stem cells can divide asymmetrically to generate both neuronal progenitors and radial glia neural stem cells. At mid-neurogenesis a secondary proliferative zone is formed, the **subventricular zone (SVZ)** generating neurons and astroglia. After further thickening of the cerebral wall, neuronal progenitors migrate along radial glia fibres and settle above the SVZ, building the **mantle zone (MZ)** and below the newly forming **cortical plate (CP; Fig.7A)**. Early-born neurons settle in deeper layers and late born cells remain closer to the pial surface following an inside-out gradient of migration (Rakic, 1990). Cells stop migrating at CP/MZ interphase as they adopt their final position in the CP. There are two main neuronal types of the mammalian cerebral cortex: excitatory pyramidal cells and inhibitory non-pyramidal cells and they are generated in distinct proliferative zones. Pyramidal neurons arise from the germinal ventricular zone (VZ) and migrate along the radial glia processes of to take up their positions in the developing cortex (Rakic, 1990), while a diverse group of mostly γ -aminobutyric acid (GABA)-ergic interneurons originate predominantly in the **medial ganglionic eminence (MGE)** of

the ventral telencephalon and navigate along a multiple **tangential** migratory routes through the **intermediate zone (IZ)** to reach the cortex (**Fig.7B**, De Carlos *et al.*, 1996; Anderson *et al.*, 1997a,b, 2001)

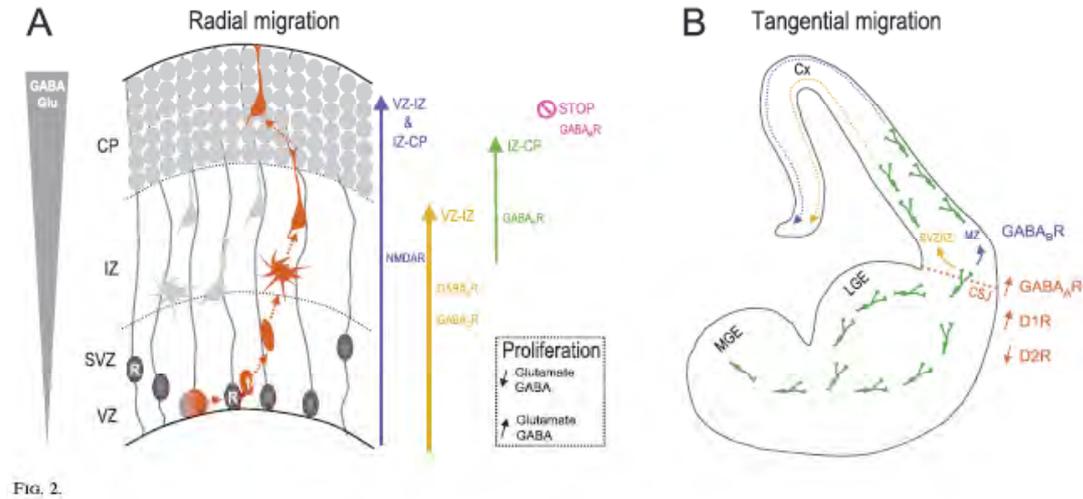


Fig. 2.

Fig.7. GABA and other neurotransmitters regulates neuronal migration in the developing telencephalon. (A) Schematic diagram showing the cortical wall of an E13.5 mouse brain. This diagram illustrates the functions undertaken by neurotransmitters and their cognate receptors in the regulation of progenitor proliferation and migration of newborn projection neurons within the cortical wall. γ -Aminobutyric acid (GABA) and glutamate (Glu) are distributed along an apical to basal concentration gradient. By activating GABA_AR or AMPA / kainate receptors, GABA and Glu regulate cell proliferation in opposite ways in ventricular zone (VZ) and subventricular zone (SVZ) progenitors (LoTurco *et al.*, 1995; Haydar *et al.*, 2000). While Glu stimulates radial migration along the entire cortical wall (Behar *et al.*, 1999), GABA triggers the ionotropic receptors GABA_AR, GABA_CR, which in turn promote the migration of projection neurons from the VZ to the intermediate zone (IZ). Importantly, activation of the metabotropic GABA_BR is specifically required for the transition of cortical projection neurons from the IZ to the caudate putamen (CP; Behar *et al.*, 2000). Finally, activation of GABA_AR also acts as a stop signal for projection neurons in the CP (Behar *et al.*, 1998). (B) GABAergic interneurons (green cells) are mainly generated in the medial ganglionic eminence (MGE) and follow distinct tangential migratory routes to integrate into the cerebral cortex (Cx). After having avoided the presumptive striatum, interneurons cross the corticostriatal junction (CSJ), which corresponds to the interface between the subpallium and the pallium, and enter the cortical anlage. These cells then follow distinct routes within the cortical wall (the VZ / SVZ and the MZ at early stages of corticogenesis). GABA promotes the crossing of the CSJ by triggering interneuron GABA_AR activation (Cuzon *et al.*, 2006), and influences the choice of cortical migratory route by acting on the metabotropic GABA_BR (Lopez-Bendito *et al.*, 2003). LGE, lateral ganglionic eminence; NMDAR, N-methyl-d-aspartate receptor. Heng *et al.*, 2007.

1.3.2. Molecular components of the GABA signaling pathway are expressed during neuronal development in a strict spatio-temporal manner

Components of GABA signaling system, GABA and GABA-synthesizing enzymes, GABA receptors and GABA transporters appear transiently during development (Buznikov *et al.* 1996; Zecevic and Milosevic 1997; Barker *et al.* 1998; Verney 2003; Lujan *et al.* 2005). The embryonic and adult GAD mRNAs and protein forms are expressed in the embryonic and postnatal development of the nervous system in a

highly specific temporal fashion (Behar et al., 1994; Ma et al., 1994; Szabo et al., 1994; Katarova et al., 2000, reviewed in Varju et al., 2000, 2001). The GAD25 form is more abundant in midgestation mouse embryos (E10.5-E12.5), while GAD44 is characteristic for later developmental stages (E12.5-E15.5) (Szabo et al., 1994). Concomitant with neuronal maturation and synaptic formation embryonic forms are gradually replaced by the adult GAD forms (Szabo et al., 1994; Somogyi et 1995; Varju et al., 2001). These results led to the speculation that the embryonic GAD proteins may be involved in the synthesis of morphogenic GABA while adult GADs provide GABA for synaptic and metabolic action in the mature GABAergic neurons (Szabo et al., 1994; Varju et al., 2000).

Studies of neurons in culture suggest an earlier presence of GABA_A receptors and currents than other transmitters, including glutamate and glycine (Lujan et al., 2005; Owens and Kriegstein, 2002; Ben-Ari et al., 2008). Several types of GABA_A receptor subunits have been detected and proved functional in rat embryo as early as E10-E11 by in situ hybridization, immunocytochemistry and pharmacological studies (Laurie et al. 1991, 1993, Bettler, Haydar et al., 2000, Behar et al., 1993). Whole-cell recordings made from embryonic and early postnatal cells have shown that early expressed GABA_A receptors change kinetic properties during development (Owens and Kriegstein, 1999). Differences in receptor properties might be due to the differences in subunit composition, intracellular calcium concentrations, or degree of receptor phosphorylation and association with cytoskeletal anchoring proteins (Lujan et al., 2005; Owens and Kriegstein, 2002; Ben-Ari et al., 2007).

Even though functional GABA_B receptor requires heterodimerization of GABA_BR1 and GABA_BR2 subunits, they display different expression patterns during embryonic development, indicating they are regulated independently. As revealed by in situ hybridization, GABA_BR1 receptor subunit mRNA is intensively expressed by E11,

and from E12 on is detected in the rat hippocampus, cerebral cortex, intermediate and posterial neuroepithelium (Kim et al., 2003; Martin et al., 2004; Li et al., 2004). In contrast, a weak expression of GABA_B R2 mRNA in the same brain areas could be detected only from later developmental stages (Li et al., 2004).

Both membrane (GAT-1, GAT-3 and GAT4) and vesicular transporter (VGAT) mRNAs have been detected at early stages of mouse forebrain development however their function in differentiating neurons has not been studied in detail yet (Jursky et al., 1996; Oh et al., 2005; Fukui et al., 2008).

1.3.3. GABA as a modulator of neurogenesis

1.3.4.1. GABA modulate proliferation and survival of neuronal progenitor cells

Emerging evidence indicates that GABA exerts modulatory, often contrasting effect on cell proliferation depending on the cell and GABA receptor type (**Fig.7A**; Owens and Kriegstein, 2002; Lujan et al., 2005; Ben-Ari et al., 2007). Estimation of [³H]-thymidine or bromodeoxyuridine (BrdU) incorporation in cells derived from the E16-E19 cortex has revealed that GABA, activating GABA_A receptors inhibited DNA synthesis in proliferating cells. This cell cycle arrest was associated with depolarization and Ca²⁺ entry as high potassium and glutamate treatments had the same effects. (LoTurco et al., 1995; Owens et al., 1996). Conversely, in rat immature cerebellar granule cells GABA promoted proliferation by indirect activation of the mitogen-activated protein kinase (MAPK) cascade (Fiszman et al., 1999). A marked difference in the rate of proliferation in response to GABA between cells isolated from the different germinal layers of embryonic mouse brain was also demonstrated. In cortical slices, GABA_A receptor activation led to an increase in the number of BrdU-labelled cells in the VZ, but to a decrease in the SVZ (Haydar et al., 2000). Both effects seem to be mediated by GABA_A receptor since treatment with GABA_A-receptor agonists mimics this action, whereas antagonists completely abolish it.

1.3.4.2. GABA regulates neuronal migration

Once neuroblasts are generated in the germinal layers, they undergo terminal mitosis and migrate to their final destination in the cerebral tissue. In cultured rat brain slices, activation of GABA_B- and GABA_C receptors stimulate the migration from the VZ and IZ (intermediate zone), respectively, while activation of GABA_A receptors acts as a „stop” signal for neurons that reached the CP (**Fig. 7A**; Behar et al., 1998). Blocking of GABA_A receptors resulted only in a delay of cell movements, but not a complete arrest of migration (Behar et al., 2000). Furthermore, in the developing rat cortex, GABA is transiently expressed in the CP, near target destinations for migrating neurons (Lauder et al., 1986; Barker et al., 1998). In vitro, femtomolar GABA or baclofen stimulated gradient-dependent or direct neuronal migration from VZ, while micromolar levels induced random migration of neurons isolated from CP. Migration was completely blocked by saclofen, GABA_B receptor antagonist (Kerr and Ong, 1992; Behar, 2001) indicating that GABA act as chemoattractants for neurons during rat cortical histogenesis via mechanisms involving GABA_B receptor (Behar, 2001). Recently, several studies demonstrated modulatory role of GABA during tangential migration supporting the notion of involvement of GABA_B receptors in neuronal migration (**Fig. 7B**; Heng et al., 2007).

1.3.4.3. GABA as a modulator of neuronal differentiation

In addition to proliferation and migration, a number of studies demonstrated that GABA-mediated signaling is important during synaptogenesis and neuronal differentiation (Redburn and Schousboe, 1987; Lujan et al., 2005). In pioneer studies by Wolff and colleagues demonstrated that continuous application of GABA to the rat superior cervical ganglion *in vivo* stimulated dendritic growth and sprouting (Wolf et

al., 1978, 1987). In cultured embryonic hippocampal and neocortical neurons, GABA_A receptor activation promoted neurite outgrowth and maturation of GABA interneurons (Barbin et al., 1993; Marty et al., 1996; Maric et al., 2001; reviewed in Lujan et al., 2005).

1.4. GABA signaling outside the nervous system

As mention above, GABA as well as different molecular components of GABA signaling have been detected in many tissues and cell types throughout the body including the pancreas, kidney, lung, liver, adrenal gland, platelets and erythrocytes, endocrine cells of the pineal and pituitary, oocytes, male and female reproductive organs, chondrocytes of the growth plate, intestinal epithelial cells and keratinocytes (Lancaster et al., 1975; Kataoka et al., 1984; Erdo and Amenta, 1986a, 1986b; Garry et al., 1986,1987,1988; Erdö et al., 1989; Gilon and Remacle, 1989; Erdö and Wolff, 1990; Parducz et al., 1992; Tillakaratne et al., 1992; Gilon et al., 1997; Calver et al., 2000; Erlitzki et al., 2000; Jin et al., 2005; Tamayama et al., 2005; Xiang et al., 2007; Mizuta et al., 2008a, 2008b).

In contrast to CNS, the role of GABA in these tissues is not so well defined. However, there is plenty of evidence of diverse GABA functions in non-neuronal tissues such as modulation of proliferation in pancreatic islet cell and tibial growth plate, regulation of hepatic regenerative activity, motility and agglutination of spermatozoa, stimulation of testosterone production, regulation of uterotubal and gut motility, and control of hormone release in ovary, adrenal medulla, pancreas and gastrointestinal tract and differentiation of epithelial cells of the jejunum (Kataoka et al., 1984; Louzan et al., 1986; Racagni et al., 1986; Gilon et al., 1991; Tillakaratne et al., 1995; Tamayama et al., 2001; Wang et al., 2005; Ligon et al., 2007).

Furthermore, GADs were shown to be expressed in non-neural tissues of the mouse embryo such as the tail bud mesenchyme, the pharyngeal pouches and arches, the ectodermal placods, the apical ectodermal ridge, mesenchyme and ectoderm of the limb buds, olfactory epithelium, heart, endothelial cells of blood vessels and lens fibers indicating that GABA might have a role also in the development of a variety of tissues (Li et al., 1995; Katarova et. al., 2000; Maddox and Condie, 2001). The cleft palate phenotype, of the GAD67 knock-out mutants and mice with a deletion or targeted mutation in the $\beta 3$ subunit of the GABA_A receptor, showed the involvement of GABA-mediated signals in the normal development of the palate (Maddox and Condie, 2001; Hagiwara et al., 2003). In addition, cleft palate phenotype was observed in the offspring of mice treated during pregnancy with GABA_A antagonist (and agonists) or GAD inhibitor (Ding et al., 2004).

1.4.1. GABA influences cancer cell proliferation and metastasis

Recently, several studies demonstrated potential role of GABA in cancer cell proliferation (Watanabe et al., 2006). Increased GAD activity and GABA content have been reported in colon cancer (Kleinrok et al., 1998; Maemura et al., 2003), breast cancer (Opolski et al., 2000), gastric cancer (Matuszek et al., 2001), and glioma (Bianchi et al., 2004). Increased expression of different GABA receptor subunits have been detected in neuroblastomas (Roberts et al., 2004), liver cancer (Biju et al., 2002), and breast cancer (Jiang et al., 2002). Increased GABA and GAD expression was found in prostate cancer patients with metastasis. In the latter, authors demonstrated that GABA promoted cancer cell invasion via GABA_B receptor pathway, and an increase in MMP (matrix metalloproteinase) activity was an underlying mechanism of action (Azuma et al., 2003).

These findings imply that GABA in concert with other signaling components could play important roles in cancer development and progression, and that this pathway can be a promising molecular target for the development of new therapeutic strategies.

2. EMBRYONIC STEM (ES) CELLS

2.1. Origin of mouse ES cells

Stem cells are a unique for their ability for substantial self-renewal and the potential to differentiate into various cell types. Stem cells that are capable of differentiation into cells of all three germ layers – ectoderm, mesoderm and endoderm are defined as **pluripotent stem cells**. Three types of pluripotent stem cells have been described so far; i) embryonic germ (EG) cells of the gonads of a postimplantation embryo, ii) embryonic carcinoma (EC) cells, isolated from embryo-derived teratocarcinomas, and iii) embryonic stem cells, derived directly from the blastocyst-stage embryo without use of immortalizing or transforming agents (Gottlieb, 2002).

Embryonic stem (ES) cell lines were originally isolated from in vitro outgrowths of inner cell mass (ICM) of mouse blastocysts (Evans and Kaufman 1981; Martin 1981). To date, various ES cell lines have been generated from other species including humans (Thomson *et al.*, 1998). Although a layer of mitotically inactivated mouse fetal fibroblasts, also called feeder cells, was necessary for the initial isolation of ES cells (Martin, 1981), it was subsequently shown that for maintenance of an established culture in its undifferentiated state a soluble factor identified as leukemia inhibitory factor (LIF) was required, which then became the standard ingredient of ES cell medium.

In cultured ES cells, STAT3 activation by the cytokine LIF is required to sustain the self-renewal (Niwa *et al.*, 1998; Matsuda *et al.*, 1999), however it is not required for normal mouse development (Stewart *et al.*, 1992).

More recently, other factors in addition to LIF were shown to contribute to maintenance of ES cell self-renewal. These include members of BMP (Ying *et al.*, 2003) and the Wnt families (Sato *et al.*, 2004).

2.2. Properties of undifferentiated ES cells

ES cells multiply by symmetrical cell divisions, which are not restricted by contact inhibition (Fuji-Yamamoto et al., 2005). It has been demonstrated that their cell cycle is characterized by a virtual lack of the G1 and G2 gap phases, thus approximately 50% of the ES cells in culture sojourn in the S phase at any given time-point. (Jirmanova et al., 2002). Furthermore, mouse ES cells can proliferate for years in an undifferentiated state while maintaining the potential to give rise to the majority of cell types found throughout the whole body. Intact embryos do not normally maintain unlimited proliferation of undifferentiated cells meaning that ES cells could be considered *in vitro* culturing artifacts (Martin, 1981; Martin and Evans, 1980). However, these cells provide a stable model of embryonic growth and development and a tool for investigation specific signaling systems.

Identification of cell surface and molecular markers has proven critical to define the molecular basis of stem cell identity. It is now well established that mouse ES cells express specific cell surface antigens (SSEA-1) and membrane-bound receptors (gp130) and possess enzyme activity for alkaline phosphatase (**ALP**) and telomerase (**Tert**) (Wobus et al., 2005; Armstrong et al., 2000). To date, there are only a few transcription factors whose appropriate level of expression is necessary to maintain pluripotency of cell population in pre- and early postimplantation embryos. **Oct4** is developmentally regulated cell-restricted transcription factor. In the early embryo it is abundantly expressed in the inner cell mass, and down-regulated in the trophoectoderm (**Fig.8**; Pesce and Scholer, 2001). Oct3/4-deficient embryos form a blastocyst, but the internal cells of the prospective ICM are unable to produce extraembryonic endoderm or other differentiated derivatives and rather become diverted into the trophoectoderm lineage (Nichols et al., 1998). The recently identified homeodomain protein **Nanog** is another key regulator of ES cell pluripotency (Mitsui et al., 2003; Hart et al., 2004). In

preimplantation embryos, its expression is restricted to and required in epiblast cells from which ES cells can be derived (**Fig.8**). Other molecular markers potentially defining pluripotency include Rex-1, **Sox-2**, Genesis, GBX2, UTF1, Pem and L17. All of these are expressed in the ICM of the blastocyst and are downregulated upon differentiation, but can be found in other cell types as well (Niwa et al., 1998, 2000, 2001, 2007). The ES cell property of self-renewal therefore depends on the stoichiometric balance among various signaling molecules, and an imbalance in any one can cause ES cell identity to be lost (**Fig. 7**).

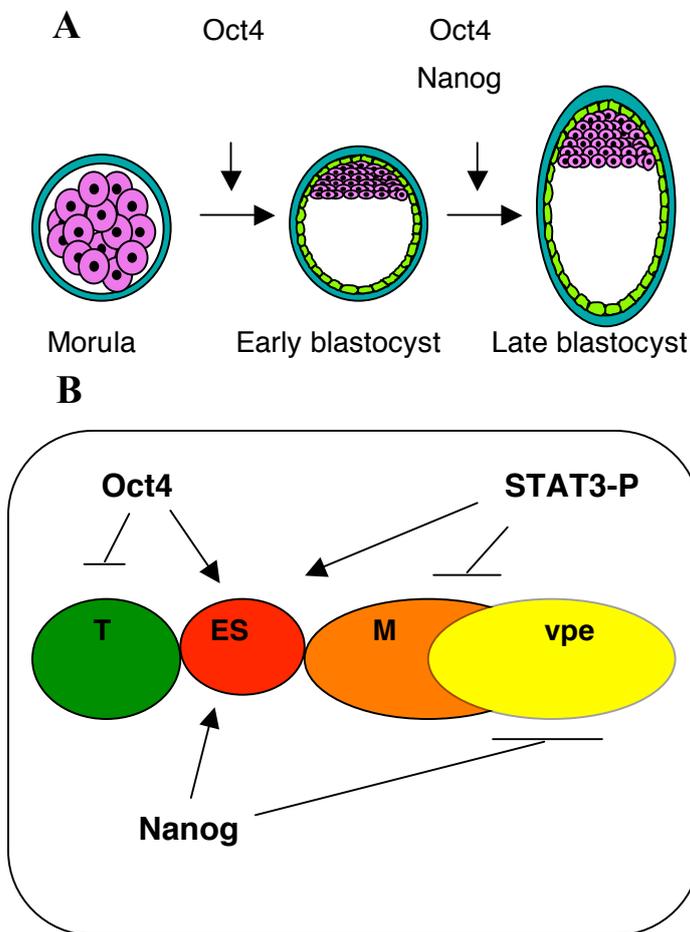


Fig.8. Model of transcription factor operating system in pluripotent lineages of the mouse embryo. (A) Pluripotent stem cells (pink) are imaged in a morula as the inner cells, which then form the inner cell mass (ICM) of the blastocyst. Oct4 is crucial for the first embryonic lineage specification, and Nanog is crucial for the second. (B) Maintenance of the pluripotent epiblast of postimplantation embryos requires Oct4, Sox2, and FoxD3. In the ES cell Oct4, Sox2, Stat3, and Nanog are essential for selfrenewal:the pools of target genes controlled by each transcription factor or combination of factors are shown in color. T, trophoectoderm; M, mesoderm; vpe, visceral/pariental endoderm. Modified from Niwa et al.2007

3. THE OCULAR LENS

3.1. An overview of mouse lens development

The lens is a highly specialized, transparent structure inside the eye that, along with the cornea, helps to refract light so as to be focused on the retina. Moreover, by changing shape, lens can change the focal distance of the eye so that it can focus on objects at various distances, thus allowing a sharp real image of the object.

The establishment of the unique architecture of the lens in the embryo is thought to depend on a series of inductive interactions that initiate its differentiation (reviewed in McAvoy, 1999). The current view is that lens induction begins during mid to late gastrulation, when involuting endoderm and mesoderm interacts with the adjacent prospective head ectoderm to give a lens-forming bias. The activation of the head ectoderm's latent lens-forming ability and the positioning of the lens in relation to the retina is accomplished by the **optic vesicle (Fig.9)**. The optic vesicle bulges out of the diencephalon, and induces the formation of the **lens placode (Fig.9)** from the overlying head ectoderm. Subsequent invagination of the placode forms the **lens pit**, which later closes to form the **lens vesicle (Fig.9)**. Cells in the posterior segment of the lens vesicle, next to the optic cup, elongate to form the **primary fibers**, whereas cells in the anterior segment of the vesicle become the lens epithelial cells. These divergent fates of embryonic lens cells give the lens its distinctive polarity. From this stage onwards, the lens grows by continued proliferation and differentiation of epithelial cells into fiber cells. Proliferation initially occurs throughout the lens epithelial compartment but at later developmental stages becomes progressively restricted to a band of cells above the equator, known as **germinative zone (Fig.9)**.

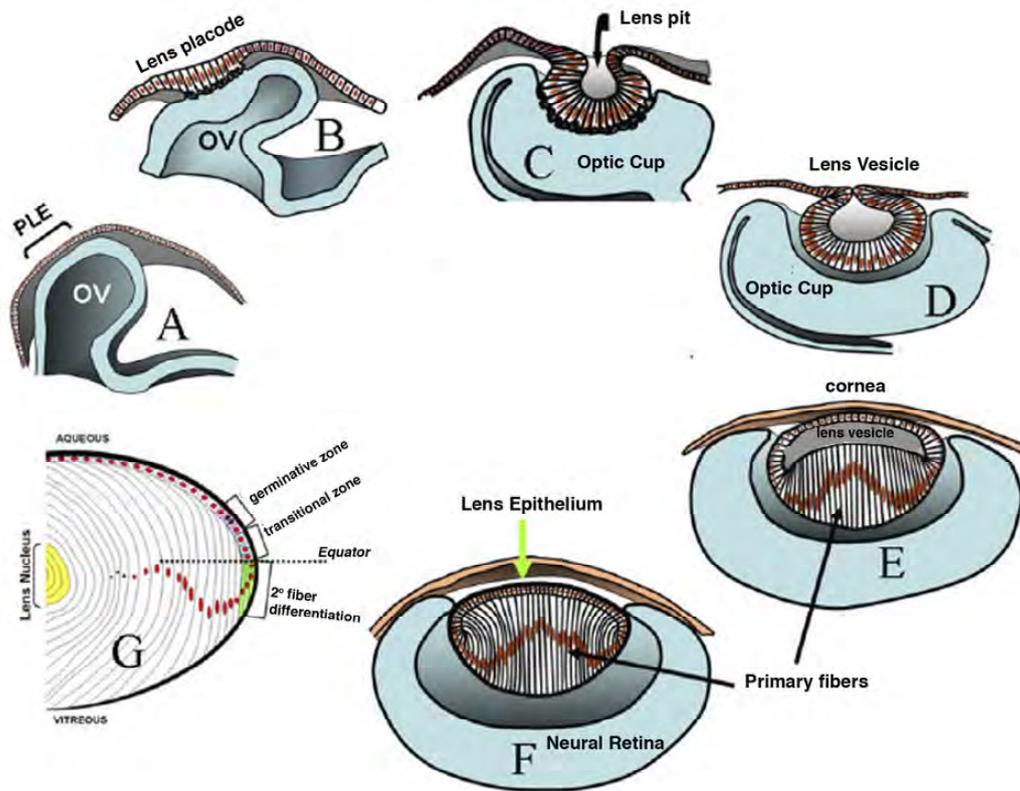


Fig. 9. Schematic view of a developing mouse eye. A. Morphological development of the lens begins as the optic vesicle (OV) approaches the presumptive lens ectoderm (PLE). B. PLE thickens to form the lens placode (E9.5). C. The lens placode invaginates forming the lens pit and the OV evaginates forming the optic cup (E10.5). D. The lens pit deepens and close to form the lens vesicle. E.-F. Once the lens vesicle has formed (E11.5) the primary lens fibers elongate from the posterior epithelium of the lens vesicle and fill its entire lumen. G. The mature lens consists of an anterior epithelial layer composed of non-proliferating central lens epithelial cells and a narrow band of proliferating cells known as the germinative zone (pink cells). Just posterior to the lens equator transitional zone (blue cells) epithelial cells begin to elongate forming secondary fiber cells (green cells). The secondary fiber cells start to elongate at the lens bow region (equator) at E14.5. The lens nucleus (yellow) is composed of primary fiber cells. Modified by *Robinson, 2006*

The lens epithelial cells are sequestered into four regions known as:

- a) a central zone (cz) – a broad polar cap of the LE covering 80% of anterior surface of the lens arrested in the G₀ stage of the cell cycle (Harding et al., 1971)
- b) pregerminative zone (pgz) – comprising 5% of LE and “daughter cells” add to the population of the lens epithelium as the anterior surface of the lens increases in size during aging (Rafferty and Rafferty, 1981);
- c) germinative zone (gz) – stem cells, which undergo mitotic division, then migrate towards the equator where they elongate adding to already existing differentiated fibers (*Fig.9*) and

d) transitional zone (tz)– nascent fibers having already begun the process of elongation (comprises approximately 5% of LE ; **Fig.9**) towards both lens poles.

The ocular lens grows throughout life in both diameter and cell number as equatorial epithelial cells elongate, lose all their intracellular organelles, synthesize crystallins- the major lens cytoplasmic proteins (α , β , γ in mammals), and an integral membrane protein known as the major intrinsic protein (MIP) (Alcala et al., 1975; Bloemendal et al., 1989,2004; Bassnett et al., 2002, 2008; Bhat, 2003)

3.2. Multiple factors controlling lens development

Distinct transcriptional regulators and growth factors are implicated in the initiation and maintenance of lens structure (**Fig.10**). A number of recent transgenic studies provided evidence that retinoic acid and members of BMP and FGF family genes play important roles as inductive signals in the lens at early stages. For example, expression of BMP genes are essential for the proper formation of lens placode (Furuta and Hogan, 1998; Wawersik et al., 1999) while the fibroblast growth factor (FGF), appears to be a key initiator during later developmental stages- lens fiber differentiation and maintenance (**Fig.10**; Robinson et al., 1995a, 1995b; Stolen and Griep, 2000). Other factors, such as transforming growth factor (TGF), platelet-derived growth factor (PDGF-A), insulin-like growth factor (IGF-1), epidermal growth factor (EGF) and hepatocyte growth factor (HGF) are also involved in lens cell proliferation and differentiation (reviewed in Lovicu and McAvoy, 2005).

A number of transcription factors have recently been identified as regulators of eye and lens development, including homeobox genes Pax6, Six3, Prox-1 and Sox2 (**Fig.10**: Ogino and Yasuda, 2000). Finally, most transcription factors involved in lens development have additional roles, in regulating other aspects of eye morphogenesis (Jean et al., 1998).

Although eye development is one of the most extensively studied processes in the embryo, the identity and specific roles of many regulatory factors controlling lens differentiation are still obscure.

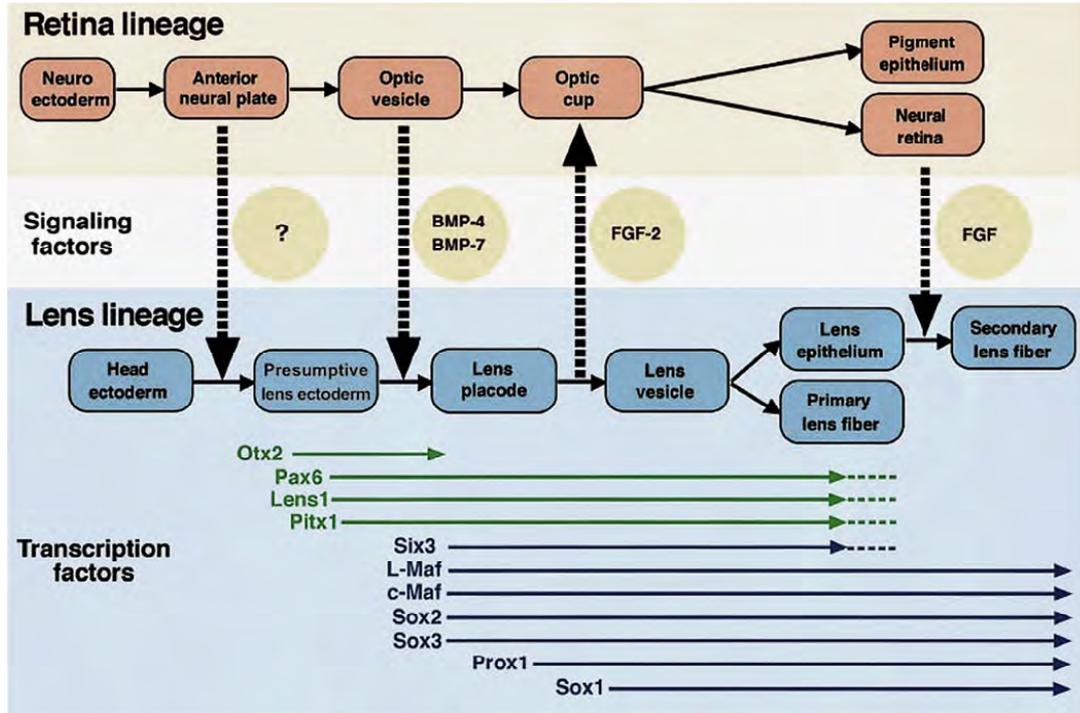


Fig.10.Diagram showing the link between inductive interactions, expression of transcription factors during lens development. Developmental stages (at the top) and major inductive events between the retina- and lens-forming tissues (bold, dotted arrows) are illustrated according to the currently favored model (Grainger 1992; Pittack et al., 1997). Signaling factors implicated in the respective inductive events are shown in yellow circles. The green and blue arrows represent the expression profile of transcription factors that were analyzed mainly at the messenger ribonucleic acid (mRNA) level in *Xenopus*, chicken, and/or mouse embryos. Only factors whose expression has been fairly well characterized are shown. Modified from Ref.: Ogino and Yasuda, 2000.

IV OBJECTIVES/AIMS

The major objective of this thesis was to characterize the GABA signaling components and functionality during differentiation using two different model systems: mouse embryonic stem (ES) cells and the ocular lens. Using these two model systems will allow us to uncover the role of GABA in developmental processes such as proliferation, differentiation and migration and the general mechanism of action, which could also apply to the nervous system.

The specific aims followed in the course of this work could be summarized as follows:

- (1) Identification and expression characterization of the GABA signaling components in undifferentiated ES cells and in the course of retinoic acid-induced neuronal differentiation: GAD forms and the two transcriptional regulators of GAD - Dlx2 & 5, GABA_A and GABA_B receptor subunits, vesicular and membrane GABA transporters.
- (2) To probe the functionality of the GABA signalling in the ES cells by studying the effect of GABA_A and GABA_B receptor activation on the intracellular calcium levels using digital calcium imaging and confocal microscopy.
- (3) To determine the effect of GABA and GABA receptor ligands on ES cell proliferation and differentiation.
- (4) Characterization of the spatio-temporal expression pattern of GABA signalling components in the developing mouse lens: GAD forms, Dlx2&5, GABA, GABA_A and GABA_B receptor subunits, vesicular and membrane GABA transporters;
- (5) Using calcium-imaging to test the functionality of the GABA signalling in the lens and in parallel, in primary lens epithelial cell (LEC) cultures prepared from mouse lenses.

V METHODS

1. ANIMAL HANDLING

Wild type C57Bl6 and FVB/Ant mice (Charles Rivers, Hungary) ranging from 6-12 weeks of age used in this study were housed in the SPF animal facility of the Institute of Experimental Medicine, under regulated light-cycle with water and food supplied ad libitum. For embryo collection, 60-150 days old females were caged with males at late afternoons and checked for vaginal plugs next morning. The presence of a vaginal plug was considered as embryonic day 0.5 (E0.5). Pregnant mice, at various gestational days, were killed by cervical dislocation and embryos were recovered by Caesarean section.

All experiments were conducted in compliance with NIH (NIH Publication #85-23, 1985) and EC (86/609/EEC/2) guidelines and approved by in-house and national committees.

2. CELL CULTURE

2.1. Preparing mouse embryo fibroblasts (MEF)

Primary culture of mouse embryonic fibroblasts was prepared using standard protocols. Briefly, 14-15 days after mating time pregnant females (C57Bl6) were killed by cervical dislocation and embryos were collected and washed in sterile phosphate buffer saline (**PBS**). Embryo head, limbs, and internal organs were removed under dissecting microscope in a sterile hood. Remaining embryo tissue was washed once in Dulbecco's Modified Minimum Medium (**DMEM**) and incubated for 45 min in 0.025% trypsin/EDTA solution at 37°C in a flask with gentle stirring. Subsequently, cell suspension was washed twice with DMEM supplemented with 10% fetal calf serum (**FCS**), centrifuged at 200g for 5 minutes and plated on tissue culture dishes (5×10^6 cells per 150-mm dish). After reaching confluence, cells were split 1:6 and

grown in culture for another 4 days. Exponential growing fibroblasts were frozen in the growing medium and kept under liquid nitrogen prior to use.

2.2. ES cell culture

Mouse embryonic stem cell line R1 (Nagy et al., 1993) were routinely grown on mitomycin C (100 μ g/ml, Sigma-Aldrich Co.) - treated or irradiated primary embryonic fibroblasts in high-glucose DMEM supplemented with sodium pyruvate, 20% heat-inactivated ES cell-qualified FBS, 2 mM L-glutamine, 1% non-essential amino acids (all from Invitrogen), 100 μ M β -mercaptoethanol (Sigma-Aldrich Co.) and 1000 U/ml of leukemia inhibitory factor (ESGRO, Chemicon International). The cultured cells were checked visually, culture medium was changed daily and cells were passaged every 24-36 hours after plating. For passaging, medium was removed and cells were incubated in 0.02% trypsin in ethylenediaminetetraacetic acid (**EDTA**) for 5 min. Subsequently, trypsin was inactivated, cells were centrifuged at 1000g rpm for 5 minutes and plated on previously prepared fibroblast monolayer.

2.3. ES cell neuronal differentiation procedure

In vitro differentiation of ES cells to neurons was promoted with all trans-retinoic acid (RA) using modified 4-/4+ induction protocol (Bain et al., 1995). Briefly, after trypsinization fibroblasts were removed from ES cells by panning - three rounds of successive differential adherence steps to tissue culture dish. The remaining cell suspension (\leq 90% ES cells) was seeded in EB medium (ES medium without LIF and reduced serum to 15%) into bacterial grade culture dish allowing ES cells to aggregate and form embryoid bodies (EBs). Medium was changed every day by transferring the EBs to a 15-ml centrifuge tube; after sedimentation of the aggregates, the supernatant was replaced with fresh EB medium. At the end of day 4, growing medium was

replaced with a fresh one containing retinoic acid (**RA**). The RA stock solution (10^{-2} M in dimethyl-sulphoxide – **DMSO**; Sigma-Aldrich) was diluted directly into the culture medium to obtain the desired concentration, 2×10^{-6} M. The medium containing RA was prepared in the dark and it was replaced every day.

At the eighth day of induction period, intact or trypsinized aggregates were seeded on poly-L-lysine (20 μ g/ml; Sigma-Aldrich Co.)-coated tissue culture dishes or coverslips and twelve hours later medium was changed to DMEM supplemented with 10% FBS. This day was considered as “day 0 after induction” (dai 0). Four sets of experimental cultures generated at different time were used for this study.

2.4. Primary lens epithelial culture

The lenses were dissected from deeply anaesthetized newborn (P0) mice in ice-cold Tyroide buffer (Sigma-Aldrich Co.) with fine forceps under a dissecting microscope. Ten lenses were incubated in 500 μ l of 0.25% EDTA solution (Sigma-Aldrich Co.) at 37°C for 10 min. Following the addition of 1ml Medium 199, supplemented with 10% FBS, to slow the reaction, the suspension was triturated 10-20 times to dislodge cells from the lens capsula. Cells were pelleted at 200g for 2 min and plated on (approximately 2 lenses per 1.88 cm²) matrigel-coated coverslip (BD Biosciences, 0.4 mg/ml) in growing medium. Medium was changed every 3-4 days and 2 hours before recordings.

Both primary cultures (fibroblast and LEC) and ES cells used in this study were kept at 37°C in a humidified atmosphere containing 5% CO₂.

3. GENE EXPRESSION ANALYSIS

3.1. RNA isolation from cells and tissues

Cells were washed twice with **DEPC** (diethyl pyrocarbonate, Sigma-Aldrich Co.) - treated PBS and lysed with TRI-reagent (1ml/90mm² surface, Sigma-Aldrich Co.) for 5min at room temperature (**RT**).

Lenses were dissected from embryos or deeply anaesthetized mice of various ages in ice-cold DEPC-treated PBS. The ciliary epithelium, vitreous and retina remnants were removed with fine forceps under a dissecting microscope. Lenses were collected in RNA-later (Sigma-Aldrich) and stored in the same solution at 4°C.

Embryonic and adult mouse brains were homogenized with TRI-reagent, and stored in liquid nitrogen.

Total RNA was isolated from cells and tissue using TRI-Reagent, following the manufacturer's protocol or by the method of Chomczynski and Sacchi (Chomczynski and Sacchi, 1987). Briefly, chloroform was added to the samples and mixed for 1min by inversion. After incubation at RT for 15 min, the samples were centrifuged at 12000 g for 15 min. The aqueous phase was transferred to a new tube and an equal volume of isopropanol was added. Carrier t-RNA or glycogen (20µg) was added to lens (E14.5, E15.5) samples to facilitate RNA recovery. The content was mixed for 1 min by inversion and then incubated at RT for 10 minutes or -20°C for overnight. RNA was recovered by centrifugation at 12000 g for 20 min. The pellet was washed in 70% ethanol, dried at RT and then dissolved in RNase-free water. The concentration and purity of the RNA preparations were determined by measuring optical density at 260 and 280 nm, the ratio of OD 260/280 being ≥ 1.8 for all preparations. The integrity of RNA samples were evaluated by running aliquots on 1.2% formaldehyde gels.

3.2. First-strand cDNA synthesis

Three micrograms of RNA were reverse-transcribed using RevertAidTM H Minus First Strand cDNA Synthesis Kit (Fermentas) with random hexamer primers or Oligo-dT primers (Invitrogen) according to the manufacturer's instructions. 200 units of RevertAidTM H Reverse Transcriptase or SUPERScript III (Invitrogen) and 0.5 µg of 12- to 18-mer primers were applied in 20µl mixture. Until use, reaction mixtures were stored at -20°C.

3.3. PCR amplification

PCR amplification was performed on one tenth of the first strand cDNA. Conditions for PCR reactions were as follows: initial denaturation for 5 min at 95°C, followed by 12-40 cycles of amplification as follows: 60 sec at 94°C; 60 sec at 55- 68°C; 60-90 sec at 72°C followed by a final extension for 10 min at 72°C. For all RT-PCR assays the following control reactions were always carried out: (a) mouse β-actin was used as a positive control for the effectiveness of RNA extraction and RT-PCR reactions and as an internal standard for quantification; (b) direct amplification of RNA without reverse transcription served to exclude the presence of contaminating genomic DNA in the samples; (c) to test for contamination with exogenous DNA first strand cDNA template was replaced with water. Primer pairs for all studied genes were selected from different exons. Specific primers were also designed to identify all alternatively spliced transcripts encoded by the mouse GAD67 gene and to avoid the direct amplification of the GAD67 pseudogene may be present as contaminating DNA in the RNA samples. Sequences, annealing temperatures and extension times for each primer pairs are listed in Table 1. PCR products were separated on 1.5 -2% agarose gels containing ethidium bromide and run in **TBE** buffer (0.089M Tris, 0.089M boric acid and 0.002M EDTA). 100 bp DNA ladder (Fermentas) was used as marker to estimate the size of the amplified products.

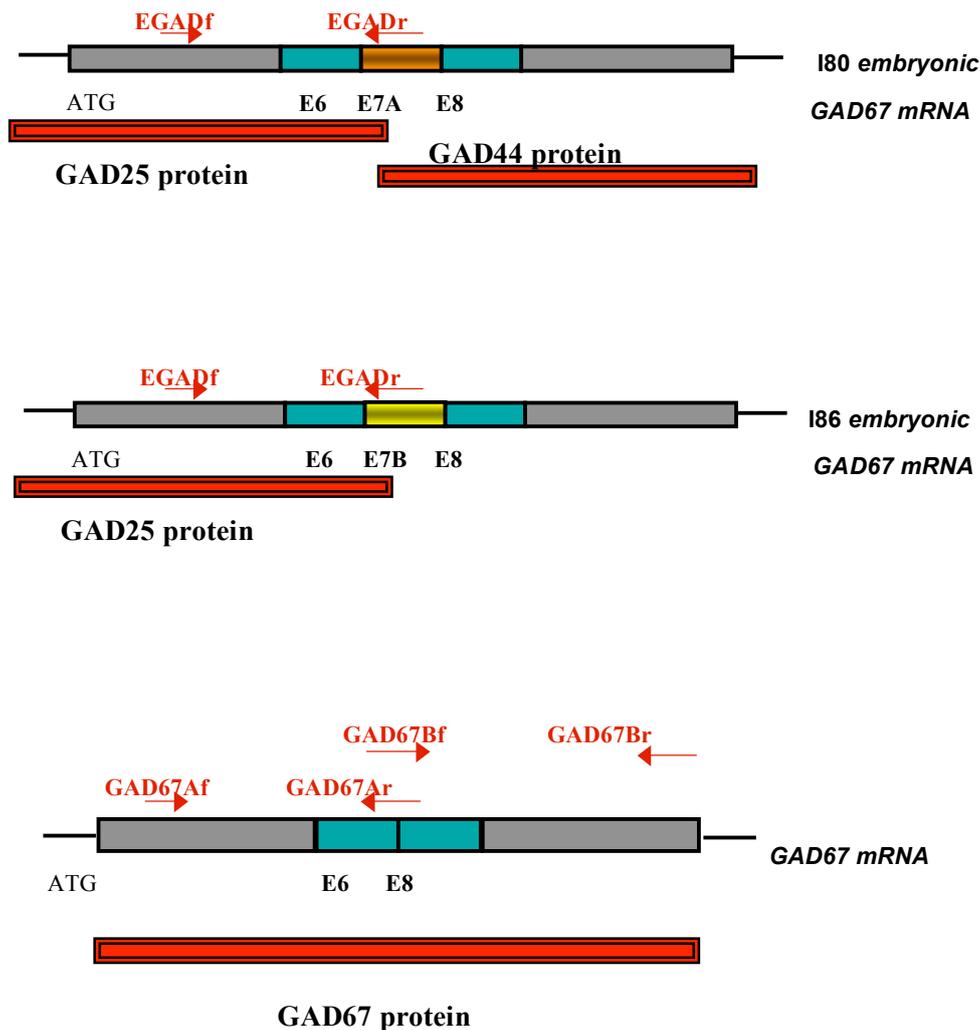


Fig. 11. Schematic representation of GAD67-family mRNAs with encoded proteins, and positions of PCR primers used for RT-PCR amplification. A1,A2. Embryonic GAD transcripts include either exon 7A (I-80, orange-filled box) or exon 7B (I-86, yellow-filled box). Primers used for PCR amplification are also indicated. Primer *EGADr* recognizes the embryonic exons (7A and 7B), while *EGADf* primer binds to an upstream region of GAD67 cDNA. This primer set co-amplifies both I-80 and I-86 transcripts. **B.** The embryonic exons (7/A and 7/B) are spliced out from the adult-specific GAD67 mRNAs, bringing together exon 6 and exon 8 (green boxes). We used two different primer pairs to amplify adult GAD67 mRNA (1) primer *GAD67Af* was selected from a region that is not represented in the GAD67-related pseudogene (Brilliant et al. 1990). Using this forward primer with corresponding reverse primer only GAD67 cDNA, but not contaminating genomic sequences derived from the pseudogene were amplified. Primer *GAD67Ar* spans a region shared by both exons 6 and 8 and therefore this reverse primer with the corresponding forward primer amplifies only the adult GAD67, but not the embryonic transcripts. (2) primer *GAD67Bf* also spans the border between exons 6 and 8 and *GAD67Br* binds to GAD67 coding region downstream of exon 8. Red boxes represent the protein products of different GAD mRNAs. Positions of the primer sets are indicated by arrows and their corresponding sequences are listed in Table 1.

TABLE 1. Primers and amplification parameters used in the semi-quantitative RT-PCR analyses

mRNA	Position	Primer Sequence	Product Size (bp)	Anneal. Temp (°C)	Exten. Time (sec)	Ref.
Actin	609	F 5'-AGCTGAGAGGGAAATCGTGC-3'	499	55-62	60-90	*
	1107	R 5'-GATGGAGGGGCCGACTCAT-3'				
Dlx2	138	F 5'-CAACAGCAGCCTGCACAA-3'	351	60	60	*
	488	R 5'-GAGTAGATGGTGGCTGGTTTC-3'				
Dlx5	186	F 5'-CTCTGCTTCTTATGGCAAAGC-3'	350	58	60	*
	535	R 5'-TTTGCCTCAGTCTAGAGAGG-3'				
EGAD	438	F 5'-GCTGGAAGGCATGGAAGGCTTTA-3'	244	58	60	*
	681	R 5'-TGAGCCCCATCACCGTAGCA-3'				
GAD67A	618	F 5'-GACTGCCAATACCAATATGTTTCAC-3'	1105	60	90	*
	1722	R 5'-GGCGCTGGGTTAGAGATG-3'				
GAD67B	85	F 5'-GATACTTGGTGTGGCGTAGCC-3'	575	62	90	*
	659	R 5'-ACCCATCTTCTGCTACAACCACTG-3'				
GAD65	13	F 5'-GGCTCCGGCTTTTGGTCTTC-3'	438	58	80	*
	450	R 5'-TGCCAATTCCAATTATACTCTTGA-3'				
GABA _A α1	51	F 5'-TCTGAGCACACTGTCGGGAAG-3'	584	62	60	Modified from (2)
	634	R 5'-ACCCATCTTCTGCTACAACCACTG-3'				
GABA _A α2	798	F:5'-TCTCTCCCAAGTGTCTTCTGGCTG-3'	583	58	60	*
	1380	R:5'-GCCCAAAGTAACCAAGTCTA-3'				
GABA _A α3	217	F 5'-CGGCTTTGGATGGCTATG-3'	737	60	90	*
	953	R 5'-ATGGTGAGAACAGTGGTGACA-3'				
GABA _A α4	1045	F 5'-CAGAAAGCCAAAAAGAAGATA-3'	550	58	90	*
	1594	R 5'-TAAATGCTCCAAATGTGACTG-3'				
GABA _A α5	164	F 5'-GACTCTGGATGGCTATGAC-3'	560	58	90	*
	723	R 5'-TGTGCTGGTGTGATGTTCTC-3'				
GABA _A α6	52	F:5'-CAAGCTCAACTTGAAGATGAAGG-3'	420	58	60	(3)
	419	R:5'-TCCATCCATAGGGAAGTTAACCC-3'				
GABA _A β1	581	F:5'-ATGGAGGAGAGGGAGCAGTAA-3'	806	55	90	(4)
	1386	R:5'-AGAAAAGGTGATGGGGAAGAA-3'				
GABA _A β2	856	F:5'-ACCAATCAACACCCACCT-3'	299	58	60	*
	1154	R:5'-CCCATTACTGCTTCGGATG-3'				
GABA _A β3	934	F 5'-GGCTGCTTGTCTTTGTTATTC-3'	337	55	60	(4)
	1270	R 5'-TGTGCGGGATGCTTCTGTCTC-3'				
GABA _A γ1	1077	F 5'-CAACAATAAAGGAAAAACCACCAGA-3'	293	58	60	Modified from (2)
	1369	R 5'-CCAGATTGAACAAGGCAAAAGC-3'				
GABA _A γ2	553	F:5'-ATTGATGCTGAGTGCCAGTTGC-3'	412	60	60	*
	964	R:5'-TGGCTATGGTGTCTTAAAGTTGTC-3'				
GABA _A γ2S	1048	F 5'-GTGGAGTATGGCACCTGCATTATTTTGTG-3	219+	65	90	(5)
	GABA _A γ2L	1362				
GABA _A γ3	646	F 5'-TGGCGGCTCTATCAGTTTG-3'	225	60	60	*
	870	R 5'-GATGCCTAATGTTGTTCTTGC-3'				
GABA _A δ	1079	F 5'-TCGTCTTTTCTCCCTCTCAG-3'	454	58	60	*
	1532	R 5'-AGCCATCCTGTCCATCTA-3'				
GABAAε1	1935	F 5'-CAATGCGAAGAACAACACTG-3'	459+	55	60	(6)
	GABAAε2	2393				
GABA _B R1	859	F:5'-ACGCATCCATCCGCCACAC-3'	347	58	60	(7)
	1183	R:5'-AACCAGTTGTCAGCATACCACCC-3'				
GABABR1a	161	F 5'-ATTCCTGCCTGTGGACTATG-3'	281	58	60	*
	441	R 5'-GCCCTGACTACAGATGCTCC-3'				
GABABR1b	88	F 5'-TCTCACTCCCTCATCTCCC-3'	485	58	60	*
	572	R 5'-TTCTCCAGCCCCACTTTTC-3'				
GABA _B R2	2171	F 5'-CCCTGGTCATCATCTTCTGTAG-3'	230	58	60	*
	2400	R 5'-CTTCATTGTCAGGCGGTGG-3'				
GABA _C p1	-25	F:5'-GAATCTATGTTGGCTGTCCAGA-3'	775	60	90	(8)
	751	R:5'-TGGTGTGGAATCTTGAAT GAG-3'				
GAT1	147	F:5'-ACGCTTCGACTTCCTCATGTCTGTC-3'	699	62	90	(9)
	845	R:5'-GAATCAGACAGCTTTCGGAAGTTG-3'				
GAT2	294	F:5'-CTGTGGCATCCAGTGTTC-3'	313	60	60	*
	606	R:5'-GACAGGCGAGGTAAAGTTCTC-3'				
GAT3	517	F:5'-AATGTGACCTCCGAGAATGC-3'	533	62	90	*
	1049	R:5'-ACCTCAGATATGGGCACACC-3'				
GAT4	1130	F:5'-ACCCAAGGCTGTCACTATG-3'	734	62	90	*
	1864	R:5'-CTGTGATGGCAGAGATGGTG-3'				

VGAT	-31 541	F:5'-TTCTGTCCCTTTCTCCCGCCCGCCG-3' R:5'-GCACCACCTCCCCGTCTTCGTTCTCCT-3'	572	68	180	(10)
NKCC1	2561 3309	5'-TGCAGAGAAGGTGCACAATAC-3' 5'-TCTGTACGGCTCGATCATGT-3'	749	58	90	*
KCC2	4 402	5'-CTCAACAACCTGACGGACTG-3' 5'-ACAGAAGGACTCCATGATGCCCGCG-3'	399	62	90	*
Oct3/4	496 806	5'-GTTCTCTTTGGAAAGGTGTC-3' 5'-ACTCGAACCACATCCTTCTC-3'	311	62	60	*
Nanog	59 297	5'-CTGGGAACGCCTCATCA-3' 5'-CATCTTCTGCTTCTGGCAA-3'	239	63	60	(11)
SOX2	240 790	F: 5'- GTGGAAACTTTTGTCCGAGAC-3' R: 5'-TGGAGTGGGAGGAAGAGGTAAC-3'	551	58	60	(12)
mTert	2079 2544	F:5'-ACCCAGGATGTACTTTGTTAAGG-3' R:5'-AGCAAACAGCTTGTTCATGTC-3'	466	58	60	(13)
RyR1	1506 1861	F:5'-CCACTTTGCCGAGTTTGC-3' R: 5'-GGTCTGTTGGAGCGTA-3'	356	60	60	(14)
RyR2	2058 2390	F: 5'-GACGGCAGAAGCCACTCACCTGCG-3' R: 5'-CCTGCAGAGAACTGACAACCTGG-3'	333	60	60	(14)
RyR3	210 482	F: 5'-GGAACCCACCTCAGAAGC-3' R: 5'-TCCCGCAGACTACATCA-3'	273	60	60	(14)
VGCC-L	3246 3336	F 5'-TCTTTATAAGGACGGCGATGTTG-3' R 5'- AGCCGAAAGGACATTGTCAAAA-3'	96	55	60	(15)
VGCC-N	3613 3740	F 5'- CGGGACCTGTGGAACATTCT-3' R 5'- CGCAGGACTCTCAGAGACTTGAT-3'	128	55	60	(15)
VGCC-R	5903 6035	F 5'- CTCTGGTGGTAACAGACCCTAGCT-3' R 5'- CTGCGCCGCGACTTG-3'	133	55	60	(15)
VGCC-T _G	2891 3166	F 5'-TGGAGGGTTCCAGGCAGAG-3' R 5'-TGCTGGAGCTCTTGGGCAG-3'	276	55	60	*
VGCC-T _H	3106 3414	F 5'-GAGGAAGATTTCGATAAGCTC-3' R 5'-CTGTTCCAGCTGGAGCGC-3'	309	55	60	*
GIRK1	435 876	F 5'-CTATGGCTACCGCTACATCACAG-3' R 3'-CTGTTTCAGTTTGCATGCTTCGC-3'	442	59	60	(16)
GIRK2	-97 5	F 5'- CGTGGAGTGAATTATTGAATCT-3' R 5'- GTCATTCTCTTTGTGCTTTT-3'	113	60	60	(17)

The positions are relative to the ATG codons of the corresponding cDNAs. Primer pairs for all studied genes were selected from different exons. F: forward, R: reverse,

(*) Gene-specific primers were designed for this study according to the published sequences as appeared in Ensembl (www.ensembl.org). (1) Varju et al., 2002 ; (2) Ma et al., 1993; (3) Liu et al., 1998; (4) Drescher, et al., 1993; (5) Homanics et al., 1999; (6) Kasparov et al., 2001; (7) Behar et al., 2001; (8) Chen et al., 2007; (9) Voutsinos et al., 1998; (10) Westmorelan et al., 2001; (11) Ginis et al., 2004; (12) Kania et al., 2005; (13) Armstrong et al., 2000; (14) Yu et al., 2008; (15) Etheredge et al., 2007; (16) Plummer et al., 2004; (17) Aguado et al., 2008.

3.4. Semi-quantitative RT-PCR analysis

For quantification of RNA expression, β -actin was generally co-amplified with the target gene for 20 cycles. When the target gene(s) and the β -actin were separately amplified due to the interference of the two simultaneous amplifications, the same master mix containing the first strand with the corresponding primers was used. RT-PCR reactions were performed in triplicate for each gene using RNA from two independent tissue preparations. In some cases the amplification reactions were trace-labeled with [³²P] dCTP (3000Ci/mmol; Institute of Isotopes, Budapest, Hungary),

aliquots were run on 1.5-2% agarose gels, and then capillary blotted in 0.4MNaOH onto nylon membranes (Hybond-N+, Amersham Pharmacia Biotech, Buckinghamshire, England, UK). Blots were exposed and quantified with Phosphor Imager and values were normalized to β -actin. The results were averaged from triplicate reactions, and presented as \pm SEM.

Quantification of data obtained by densitometry of ethidium bromide stained gel images was carried out using the public domain NIH Image program (free software developed at the U.S. National Institutes of Health available at <http://rsb.info.nih.gov/nih-image>). For each experimental group β -actin densities were normalized to the maximal actin expression level. The intensities of the PCR bands for the studied genes were then normalized to the corresponding equalized β -actin densities. Relative expression levels for each transcript were represented by using a gray scale or plotted in a histogram.

4. WESTERN BLOT ANALYSIS

Cells were harvested from the culture dishes in Ca^{2+} -. Mg^{2+} -free PBS containing 0.2%EDTA and homogenized in standard buffer (SB) containing 1mM aminoethylthioammonium bromide (AET), 0.2mM pyrodoxal 5-phosphate (PLP), 0.1mM phenylmethylsulfonyl fluoride (PMSF), 10mM benzamidine, 1mg/ml pepstatine, 1mg/ml leupeptin as protease inhibitors followed by brief sonication.

Lens protein extracts were prepared by homogenizing freshly isolated lenses in SB buffer containing 1 mM ethylene glycol tetraacetic acid (**EGTA**; Sigma).

Brain homogenates containing detectible amount of both embryonic and adult GAD forms, prepared from the newborn (P0) mice, were loaded to the same gels as controls.

Protein concentration was determined according to Bradford and samples were

adjusted to 2µg/µl in 2X Laemmli sample buffer (125M Tris-Cl, pH 6.8, 4 % SDS, 20% (v/v) glycerol, 0.2 M DTT, 0.02% bromphenol blue). Denatured protein samples were subsequently run in 10 or 12% sodium dodecylsulphate-polyacrylamide gels (SDS-PAGE; 20µg protein/lane) in mini Bio-Rad apparatus according to the manufacturer instructions.

Blotting was initially performed at 60V for 20 min, followed by 30 min at 80V and 5min at 100V to ensure blotting of larger proteins also.

The membranes were blocked and then were incubated overnight at 4°C with the primary antibodies diluted in TBST containing 1% bovine serum albumin (50 mM Tris±HCl pH 7.4, 100 mM NaCl; 0.05% Tween-20). The following antibodies were used: polyclonal rabbit sera no. 6799 (1/2000) recognizing all GAD forms also including GAD25 and GAD44 (Katarova et al. 1990; Szabo et al. 1994), polyclonal rabbit sera no. 8876 (1/1000) specific for GAD25 (Szabo et al. 1994) and monoclonal anti-βactin antibody (Sigma-Aldrich). After incubating with the appropriate alkaline phosphatase-conjugated secondary antibodies, blots were stained with NBT/BCIP (Katarova et al. 1990).

The molecular sizes of protein were estimated on the basis on molecular weights standards (BioRad) run on the same gels.

5. GABA MEASUREMENT BY HIGH PERFORMANCE LIQUID CHROMATOGRAPHY

These experiments were performed in collaboration with Mária Baranyi from Department of Pharmacology, Institute of Experimental Medicine Hungarian Academy of Sciences.

Frozen lens tissue was homogenized in 50 µl 0.2 M ice-cold perchloric acid, cleared by centrifugation (3510 x g for 10 min at 4°C) and neutralized by addition of 2M KOH (25:7 ratio). Derivatized 1-(alkylthio)-2-alkylisoindol amino acid adducts were

separated on a 5 μm Discovery HS C-18 (150x4.6 mm) analytical column that was equilibrated with 11.25% methanol-acetonitrile (3.5:1 v/v) in 0.01 M PBS, pH 7.2. The mobile phase was 22.2% acetonitrile in methanol and the profile of the gradient was as previously described (Mally et al., 1996). Amino acid derivatives were detected by filter fluorometer at 340 nm excitation and 455 nm emission wavelengths. Concentrations were calculated by a two-point calibration curve internal standard method: $(A * f * B)/(C * D * E)$ (A: Area of GABA; B: Sample volume; C: Injection volume; D: Response factor of 1 nmol GABA standard; E: Protein content of sample; f: factor of Internal Standard (IS area in calibration/IS area in actual)). The data was expressed per μg protein.

6. ELECTRON MICROSCOPY

Transmission electron microscopy sample preparation and photomicroscopy in collaboration with Dr. József Takács and Dr. Csaba Vastagh from Neurobiology Research Group, United Research Organization of the Hungarian Academy of Sciences and Semmelweis University .

ES cells grown for one day on a feeder layer in Petri dishes were fixed in 2% paraformaldehyde and 2% glutaraldehyde in 0.1 M PB pH 7.4 at RT for 20 min, then extensively washed and post-fixed with 1% osmium-tetroxide (0.1 M PB, pH 7.4) for 1 hour. All the following steps were carried out in the same plastic Petri dishes that cells were grown on, with cells fixed on the surface. Dehydration process was performed through serial incubation of graded series (50, 70, 90% and 100%) of ethanol. Cells were then sequentially infiltrated with mixture of 100% ethanol and epoxy resin (DurcupanTM ACM, Fluka) at various (2:1, 1:1, 1:2, repectively) ratios, followed pure resin, O/N. From the next day on, samples were incubated at 56°C for 72 hour for appropriate degree of polymerization. Series of ultrathin sections were cut

at 60-70 nm thickness and mounted on copper grids, counter-stained with uranyl-acetate and lead-citrate, finally examined and microphotographed in a transmission electronmicroscope at 80 kV.

7. IMMUNOHISTOCHEMISTRY

7.1. Tissue preparation for immunohistochemistry

Whole embryos or heads were fixed in 4% (w/v) PFA in PBS for 1 hr at RT or in 4% PFA-0.1% glutaraldehyde-PBS overnight at 4°C for GABA detection. The embryonic tissue was immersed in 20% sucrose-PBS solution at 4°C, embedded in Cryo-gel (Instrumedics, Inc.), frozen in dry ice and sectioned at 25 µm on a cryostat (Microm HM550).

Postnatal eyes were removed with curved forceps, placing them behind the globe and gently pulling forward. The eyes were placed in a beaker containing 20 ml ice-cold 4% (w/v) PFA in PBS containing 0.1% Tween-20 (**PBT**) and fixed in a microwave (MW) oven. This method ensures rapid and deep penetration of the fixative inside the lens and provides good ultrastructural preservation. The optimal conditions of MW exposure for the P7 eyes was 2x10 sec; for P14 eyes: 3x8 sec; for P30 eyes: 3x10 sec, respectively, at 25-30°C. The temperature of the fixative was maintained by cooling in ice. Eyes were post-fixed in 4% (w/v) PFA in PBT up to 1 hr at RT, and then subsequently immersed in ascending grades of sucrose-PBS solution at 4°C, embedded in Cryo-gel, sectioned at either 16 or 25 µm and collected on gelatin-coated glass slides.

7.2. GAD immunohistochemistry

For GAD staining, sections were briefly dried, washed in PBS and dehydrated in graded methanol in PBS for 15 min/step, stored in 100% methanol overnight at -20°C and finally rehydrated through descending methanol series into PBS. Subsequently,

sections were incubated in 1% H₂O₂ in PBS for 30 min, washed 3x10 min with PBT and blocked for 2 hrs in 0.5% Blocking Reagent (Roche Applied Sciences) in PBS. The sections were then incubated overnight at 4°C with rabbit anti-GAD serum #6799 (Katarova et. al., 1990) at a dilution of 1:500 or rabbit anti-GAD65 serum N65 at 1:500 (a kind gift from Pietro de Camilli, Yale University; Solimena et al., 1993) in 1% bovine serum albumin (BSA, Sigma-Aldrich CO) in PBT. Sections were washed 3x10 min at RT with PBT, then incubated for 2 hrs with a biotinylated anti-rabbit antibody (1: 500 in PBT, Vector Laboratories), followed by 2x10 min PBT and 1x10 min PBS washes. Sections were subsequently incubated with ExtrAvidin-HRP (Sigma-Aldrich Co.) diluted 1:500 in 1% BSA-PBS for 90 min at RT. Finally sections were reacted with 0.3 mg/ml diaminobenzidine (DAB, Sigma-Aldrich CO) in TBS (Tris-buffered saline, 50 mM Tris-Cl, pH 7.4, 0.15 M NaCl) in the presence of 0.001% H₂O₂.

7.3. Immunofluorescent labeling

For immunofluorescent detection of GABA, GABA_Aβ₃, GABA_BR2 receptor subunit, VGAT, GAT-1 and GAT-3, slides with serial cryostat sections were dried, washed with TBS for 5 min. They were subsequently blocked in 1% BSA, 0.02% saponin (Sigma-Aldrich Co.) in TBS for 2 hours at RT. The sections were incubated with primary antibodies: rabbit anti-GABA (Sigma-Aldrich Co., 1:5000), rabbit anti-GABA_Aβ₃ (Novus Biologicals, 1:1000), rabbit anti-GABA_BR2 (BD Biosciences, 1:500), guinea pig anti-VGAT (Calbiochem, 1:5000), guinea pig anti-mouse GAT1 (gift from N. Nelson, 1:1000), rabbit anti-GAT3 (Abcam, 1:1000), diluted in 1% BSA, 0.005% saponin in TBS, overnight at 4°C. After washing with TBS containing 0.005% saponin, the sections were incubated for two hours with biotinylated secondary antibody (1:500, Vector Laboratories) in 1% BSA, 0.005% saponin in

TBS, followed by three washes with TBS, 0.005% saponin. For visualization, Streptavidin-conjugated Cy3 (1:2000, Jackson Res.) was applied for two hours at room temperature. Cell nuclei were labeled for 3min with 0.8 µg/ml DAPI (Sigma-Aldrich Co.). Sections were washed with TBS, and mounted with Mowiol (Calbiochem) supplemented with 1,4-diazabicyclo-[2,2,2]-octane (DABCO, 2,5% (w/v), Sigma-Aldrich Co.).

8. IMMUNOCYTOCHEMISTRY

At various times after plating on coverslips, the cells were fixed in 4% PFA-0.1% glutaraldehyde in PBS (for GABA detection) or 4% PFA in PBS (for all other antibodies) for 20min at room temperature. Cells were permeabilized with 0.2% Triton X-100 for 5min and blocked in 0.5% blockig reagent (Roche Applied Science) or in 10% goat serum for 1-2 hours at RT. Cells were then incubated overnight at 4°C with primary antibodies in 1% BSA PBT or in TBST as follows: rabbit anti-GABA (Sigma-Aldrich Co., 1:5000), rabbit anti-GAD serum #6799 (1/1000, Katarova et al. 1990), rabbit anti GAD-65 serum N65 (1/500, Solimena et al. 1993), goat anti- α A-crystallin (Santa Cruz Biotechnology, 1:500), rabbit anti- α B-crystallin (Biomol International Inc., 1:500), rabbit anti-GABA_A β 3 (Novus Biologicals, 1:1000), rabbit anti-GABA_BR2 (BD Biosciences, 1:500), guinea pig anti-VGAT (Calbiochem, 1:5000), guinea pig anti-mouse GAT1 (gift from N. Nelson, 1:1000), rabbit anti-GAT3 (Abcam, 1:1000), mouse anti-uvomoruline (E-cadherin, Sigma-Aldrich Co., 1:1000), mouse anti-N-cadherin (BD Transduction Laboratories, 1:1000), and goat anti-Oct4 (Abcam, 1:1000).

For light-microscope visualization, appropriate biotinylated secondary antibodies in dilution 1/500 were applied for 1.5-2 hours at RT, followed by incubation in with avidin-biotin complex in dilution 1/500 for 1hour at RT. Staining was visualized by

incubating the cultures in 0.05% 3,3'-diaminobenzidine (DAB) containing 0.003% hydrogen peroxide in 50mM TBS (pH 7.6). For fluorescence detection, fluorescein isothiocyanate (FITS)-conjugated anti-rabbit or Texas red-conjugated anti-mouse secondary antibody (both from Sigma Aldrich) was diluted in the blocking solution at 1/500 and were applied for 1 hour at RT. Cell nuclei were labeled for 3min with 0.8 μ g/ml DAPI (Sigma-Aldrich Co.). In each experiment, control staining was performed following the same procedure but omitting the primary antibody.

Specimens were examined under Zeiss Axioscop-2 microscope supplied with AxioCam HRc using AxioVision 4.6 software or confocal laser scanning microscope Zeiss LSM 510 META and Olympus FV500, respectively.

9. $[Ca^{2+}]_i$ IMAGING STUDIES

9.1. Measurements of change of intracellular Ca^{2+} concentration in the undifferentiated ES cells

ES cells line R1 grown for one day on a layer of primary embryonic fibroblast plated on the previous day on glass coverslips were loaded with 2.5 mM Fluo-4AM (Molecular Probes, Eugene, OR, USA) in perfusion buffer containing 0.2% pluronic acid (Sigma) for 45 min at 37°C. The buffer composition was (in mM): NaCl, 154; KCl, 5.6; $CaCl_2$, 2.3; glucose, 5.6; HEPES, 10, pH. 7.4. No specific measures were taken to remove fibroblasts from the cultures. Their distinct morphology allowed their identification, and Ca^{2+} measurements were only obtained within ES colonies. Cells were, mounted in a submerge type perfusion chamber onto the stage of an upright microscope (Olympus BX61WI,) equipped with the FluoView300 confocal laser-scanning system (Olympus). Serial scanning of cells was carried out at 488 nm excitation wavelength and fluorescence monitoring at 530 nm through a 20x water immersion objective. Whole field images were acquired every 3 s. Fluorescence

intensity was followed over time within regions of interest (ROI), covering a single ES cell. Fluorescence changes were expressed as the percentage of the average fluorescence value of the first 30 images. GABA (1mM; Sigma-Aldrich), GABA_AR agonist muscimol (30μM; Sigma-Aldrich) and GABA_BR agonist baclofen (20μM; Sigma-Aldrich) were applied in the absence and presence of the corresponding antagonists: bicuculline methiodide (20μM; Sigma-Aldrich), CGP 55845 or CGP 54626 (20μM; Tocris) in a randomised order. Cyclopiazonic acid (10μM; Tocris) and cadmium chloride (100μM; Tocris) were always applied after the application of the tested agonist alone.

9.2. Measurements of change of $[Ca^{2+}]_i$ in the lens cells

9.2.1. Intact lens

Freshly isolated lenses from neonatal mice were washed in HEPES-buffered saline solution (HBSS) containing (in mM): 137 NaCl, 5 KCl, 20 HEPES, 10 glucose, 1.4 CaCl₂, 3 NaHCO₃, 0.6 Na₂HPO₄, 0.4 KH₂PO₄ at pH 7.4. Intact lenses were loaded for 30 min at 37°C with 5 μM Fluo-4/AM in the presence of 0.1 % pluronic acid F-127 (both from Molecular Probes, OR, USA). A stock solution containing 1 mM Fluo-4/AM, 20 % (w/v) pluronic acid in DMSO was prepared in advance. Subsequently the lenses were secured by means of a mesh to the bottom of a plastic perfusion chamber mounted on a laser scanning confocal microscope (LSM, Olympus FV-500). Recordings were made under excitation light emitted by an argon laser at 488 nm using a 10x optical and 5x digital magnification. Image acquisition frequency was set at one image/1.8 sec. Cells were superfused with HBSS buffer at a rate of 2 ml/min using a peristaltic pump (Gilson, Inc., WI, USA) initially for 2 min with HBSS to wash out excess dye. Additional 2 min perfusion with HBSS was performed to obtain a steady baseline, then 1 mM GABA was applied for 2 min, followed by 2 min HBSS and finally HBSS-50 mM KCl. Analysis of fluorescence intensity was

performed using the imaging software of the Olympus FV500 (FluoView™).

9.2.2. Primary LEC culture

Primary LEC cultures were loaded with Fluo-4/AM (2.5 μ M Fluo-4/AM in 0.1 % pluronic acid) in growth medium at 37°C for 45 min, in a humidified atmosphere of 5 % CO₂/ 95 % O₂. Subsequently the medium was changed and de-esterification of the dye was allowed to proceed for another 15 min. The glass coverslips with attached cells were placed in a POC-R type submerged chamber maintained at 24°C (PeCon GmbH, Germany). In each experiment cells were perfused for 2 min at 2 ml/min with artificial aqueous humor (AAH) of the following composition (in mM): NaCl, 130; KCl, 5; CaCl₂, 1; MgCl₂, 0.5; D-glucose, 5; NaHCO₃, 5; HEPES, 10 (pH 7.25) to obtain a steady baseline. GABA (1 mM), muscimol (100 μ M) or baclofen (20 μ M), all from Sigma-Aldrich were applied for 2 min, followed by a 10 min perfusion with AAH.

Whenever GABA_A or GABA_B receptor antagonists were used, cells were initially perfused for 2 min with the antagonist: 20 μ M of bicuculline (Sigma-Aldrich; a GABA_AR antagonist) or 20 μ M of CGP55845 (Tocris Bioscience, UK; a GABA_BR antagonist) followed by 2 min with 100 μ M muscimol + 20 μ M bicuculline or 20 μ M baclofen + 20 μ M CGP55845, respectively. Perfusion was continued for another 10 min with AAH then cells were perfused for 2 min with AAH containing only the agonist (muscimol or baclofen, respectively). Cultures were observed under 10x (1.3 NA) or 20x (0.7 NA) objectives of a laser confocal scanning microscope (Zeiss LSM 510 META, Carl Zeiss Inc., Germany) and scanned every second. All settings of the laser, optical filter and microscope as well as data acquisition were controlled by LSM 510 META software. Excitation was at 488 nm and emitted light was read at 515 nm. Analyses were performed at the single-cell level and expressed as relative

fluorescence intensity.

10. CELL PROLIFERATION STUDIES

For these experiments ES cells were plated at density of 1000/well on gelatin-coated 96-well assay plates (Corning, Inc). Twelve hours following cell seeding, cultures were treated with muscimol, SR95531, and baclofen, with final concentration 100, 50, 50 μ M, respectively and cultured for another 12 or 36 hours. Medium was replaced every 12 hours. Cellular proliferation was first measured using a fluorescence-based assay (CyQUANT Cell Proliferation Assay Kit; Molecular Probes, Invitrogen) following manufactures's instructions. Fluorescence was measured by a multilabel microplate reader (Victor 3V; Per kin Elmer) using excitation and emission wavelengths of 480 and 550nm, respectively. A histogram was created by converting sample fluorescence values into percentage and normalized to the ones that were measured under control conditions (100%). Cellular proliferation: cell number and ES cells clone size were also determined after crystal violet (CV) staining and images of the entire growth area (from three different wells) were quantified in the ImageJ software (ImageJ; www.nih.gov).

11. DATA ANALYSIS

Data are presented as either representative single experiments or mean \pm S.E.M.. Statistical significance was estimated by unpaired two-tailed Student's t test between control and treated cells (GraphPad InStat version 3.0a for Macintosh, GraphPad Software, San Diego California USA, www.graphpad.com). The level of significance was expressed as follows: * $p < 0.05$, ** $p < 0.01$, and *** $p < 0.0001$.

VI RESULTS

1. GABA SIGNALING IN THE MOUSE EMBRYONIC STEM CELLS

1.1. *Characterization of mouse ES cells, line R1*

1.1.1. *ES cells of line R1 express markers of pluripotency*

Pluripotency can be defined as the ability of a cell to differentiate into the multiple cell types of the embryo and adult (Solter, 2006; Niwa et al., 2007). In this study we worked with mouse embryonic stem cells line R1 (Nagy et al., 1993), widely used by many laboratories as they can contribute with high percentage to all tissues (including the germline) in chimaera production. In the presence of LIF and embryonic fibroblasts the undifferentiated ES cells form typical spindle- to round-shaped colonies (**Fig.12A**). The pluripotency of cells was monitored by the expression of Oct3/4, Nanog and Sox-2, key players in maintenance of the undifferentiated state in ES cells and early embryos (see Introduction) and the telomerase (mTert) which is highly expressed in stem cells but reduced upon differentiation (Amstrong et al., 2000). Our RT-PCR results showed that Oct3-4, Nanog, Sox-2 and mTert mRNAs were abundantly present in ES cells after one day in culture and nearly absent, with exception of Sox-2, in the newborn mouse brain (MB; **Fig.12B**). Furthermore, immunocytochemical staining with Oct3/4 specific antibody revealed an intense and uniform nuclear labeling of all ES cell colonies (**Fig.12C**).

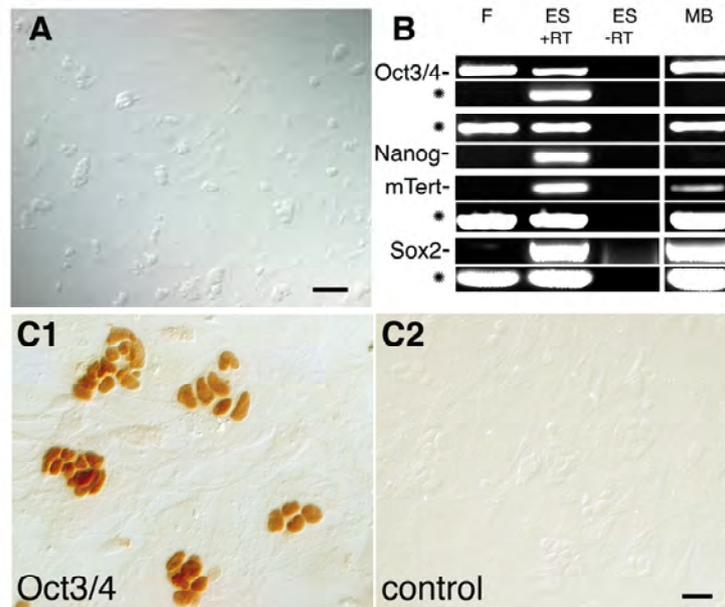


Fig.12. Mouse embryonic stem cells line R1 express markers characteristic for pluripotent phenotype. **A.** Phase-contrast image of one-day-old culture of ES cells grown on primary mouse fibroblasts. **B.** Expression of Oct3/4, Nanog, Sox-2 and mTert mRNAs in one-day-old R1 ES cell culture as revealed by semi-quantitative RT-PCR analysis. F- mitomycin-treated fibroblasts; ES- embryonic stem cells; - RT – reverse transcriptase omitted from first strand synthesis reaction; MB – mouse brain; * - actin; **C.** Images of ES cell culture immunostained for transcription factor Oct3/4. Scale bar: 100µm in A, 20µm in C1,C2.

1.1.2. Ultrastructural features of cultured R1 ES cells show high level of similarity with other ES cells

The fine structure of mouse ES cells was analyzed after one day in culture by transmission electron microscopy (TEM). All of the cells, appeared single or in compact colonies, had large nuclei and scanty cytoplasm with fewer organelles. The appearance of nuclei was mostly euchromatic with small regions of electron dense area of perinuclear heterochromatin (**Fig.13A**). Some of the ES cells were mitotic displaying characteristic midbody/interbody formations (**Fig.13B**). Furthermore, cells had close contacts with each other, connected by putative adherent and apparent gap junctions (**Fig.13D**). Occasionally we could detect clathrin-coated vesicles (CCV) near the membrane (**Fig.13C**).

These features are consistent with the ultrastructural characteristics of undifferentiated mouse ES cells (Niwa et al, 2007; Todorova et al., 2007).

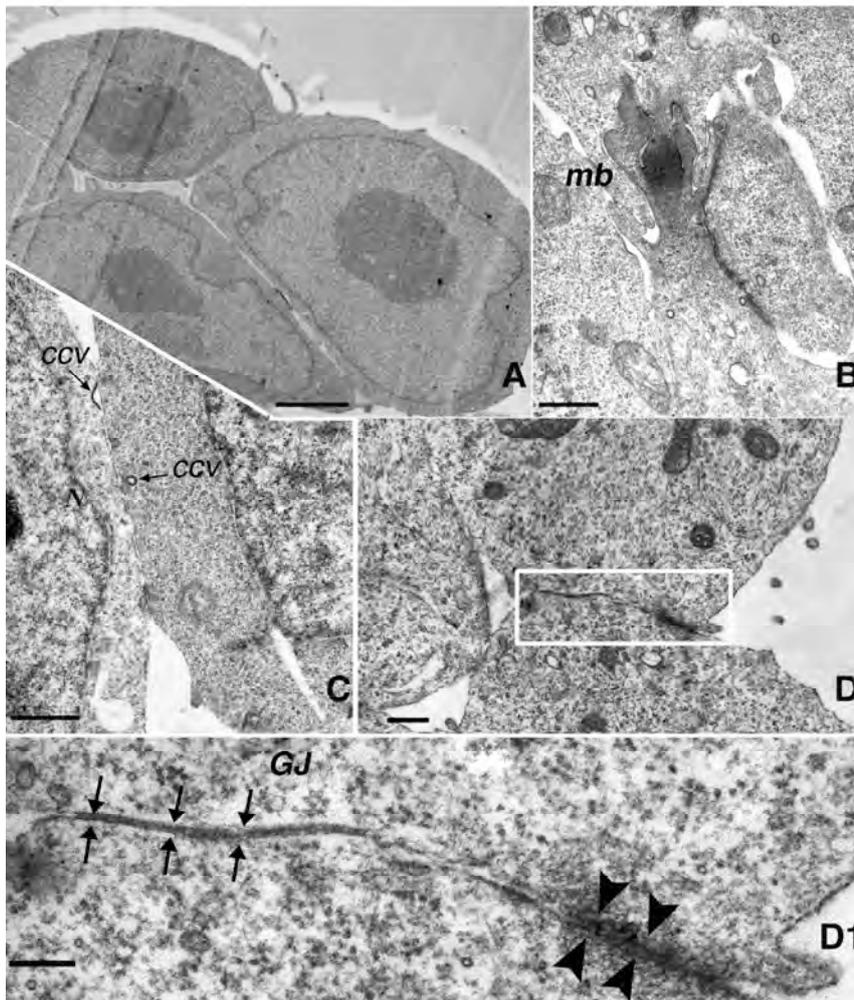


Fig.13. Ultrastructure of the undifferentiated ES cells. (A) Low magnification TEM image of an ES cell colony. Note the large mostly euchromatic nuclei. (B) Midbody/interbody formations between cells. (C) Clathrin-coated vesicles near the membrane. (D) Cells were tightly connected with apparent gap junctions (GJ) and adherent junction (arrows and arrowheads, respectively in enlarged image of boxed area in D1) mb, midbody; ccv, clathrin coated vesicle; GJ, gap junction. Scale bars: 2 μ m in A, 0.5 μ m in B, 500nm in C, 400nm in D and 200nm in D1. TEM study was done in collaboration with Dr. Jozsef Takacs and Dr. Csaba Vastagh.

1.2. Undifferentiated mouse ES cells express different GAD isoforms and contain γ -aminobutyric acid (GABA)

Expression of different forms of GABA synthesizing enzyme GAD and their upstream transcription factors, *Dlx2* and *Dlx5*, in the ES cells was examined by RT-PCR and immunocytochemistry. As outlined in the Methods Chapter, we identified the embryonic and adult transcripts encoded by the *GAD67* gene, as well as the *GAD65* transcript with specific primer sets. Adult *GAD67* amplification was performed using a reverse primer spanning the exon 6-8 border region, thus

precluding amplification of exon 7A/B-containing embryonic cDNAs. For simultaneous amplification of the two embryonic GAD transcripts I-86 and I-80, (EGAD) reverse primer was derived from embryonic exon 7A/B. As shown in **Fig.14A**, high mRNA levels of Dlx2, EGAD and GAD65 and only traces of Dlx5 and adult GAD67 transcript were found in undifferentiated ES cell but not in the embryonic fibroblasts - feeder cells.

The presence of GABA in ES cells was investigated by immunocytochemistry. As shown in **Fig.14B** relatively strong GABA immunoreactivity was observed in all ES cells while underlying fibroblasts were not labelled. All ES colonies showed consistently strong staining.

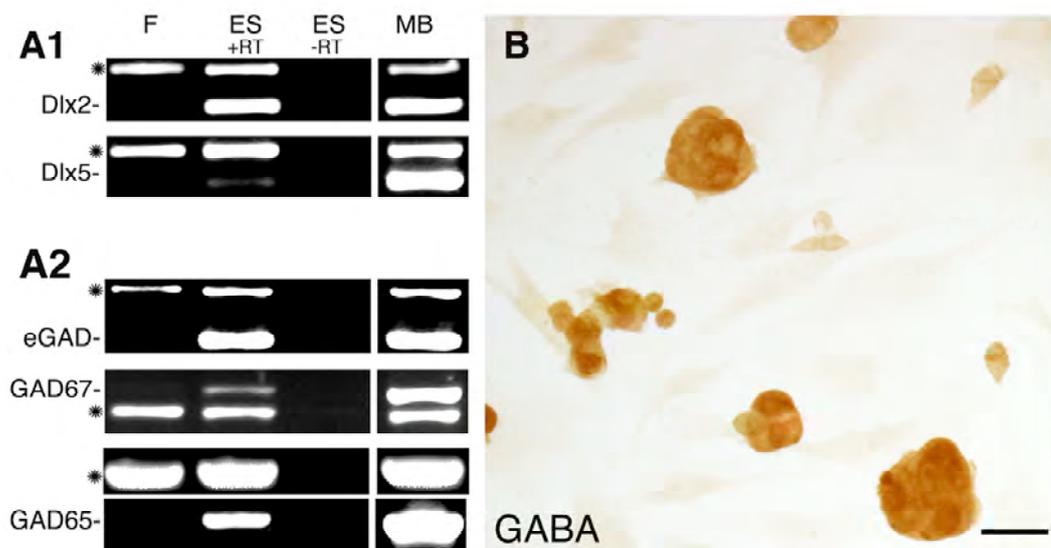


Fig.14. Expression of different Dlx and GAD forms and GABA in ES cells as revealed by RT-PCR (A1, A2) and immunocytochemistry (B), respectively. F- mitomycin-treated fibroblasts; ES- embryonic stem cells; - RT – reverse transcriptase omitted from first strand synthesis reaction; MB – mouse brain; * - actin; Scale bar: 20µm.

1.3. ES cells express both membrane and vesicular GABA transporters

We investigated the expression of all four membrane transporters (GAT1-4) and the vesicular GABA (VGAT) transporter in undifferentiated ES cells both at RNA and protein levels using RT-PCR and immunocytochemistry, respectively. The results shown in **Fig.15A** demonstrate that only GAT-1 and VGAT are expressed in ES cells.

Furthermore, stainings with specific antibody demonstrated, especially in the case of GAT1, preferential membrane localization, although the fluorescence intensity varied from cell to cell (**Fig.15B**).

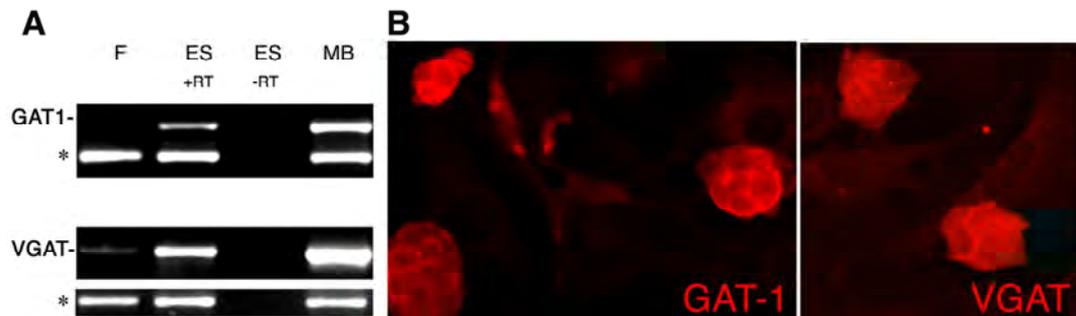


Fig.15. Expression of membrane and vesicular GABA transporters in undifferentiated ES cells, as revealed by RT-PCR (A) and immunocytochemistry (B). F- mitomycin-treated fibroblasts; ES- embryonic stem cells; - RT – reverse transcriptase omitted from first strand synthesis reaction ; MB – mouse brain; * - actin; Scale bar: 20 μ m.

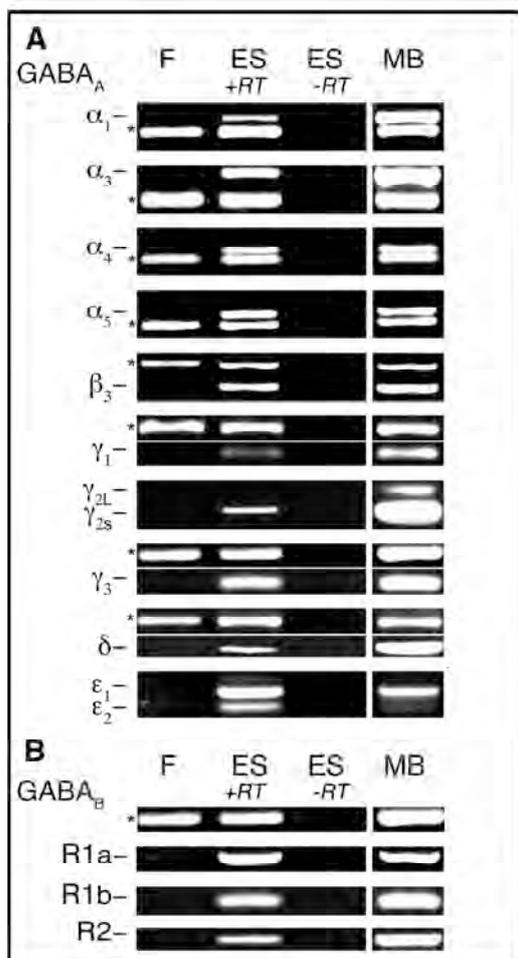
1.4. GABA_A and GABA_B receptor subunits are expressed in ES cells and transported to the cell membrane

First, we have studied the expression of different GABA_A and GABA_B receptor subunits in undifferentiated ES cells cultures by RT-PCR. We found that some, but not all, mRNAs for GABA_A receptor (GABA_AR) subunit polypeptides are expressed in undifferentiated ES cells. We could detect transcripts of α 1, α 3, α 4, α 5, β 3, γ 1, γ 2S, γ 3, δ as well as both splice forms of ϵ subunit in the ES cells, but not in fibroblasts. No α 2, α 6, β 1 or β 2 transcripts were detected even after 35 amplification cycles. Noteworthy is the presence of the shorter version of γ 2 subunit - γ 2S (**Fig.16A**) and the absence of γ 2L. Moreover, mRNAs coding for the GABA_C receptor subunits (ρ 1 and ρ 2) were not detected in the ES cells.

We found also abundant expression of both GABA_BR1 subunits (GABA_BR1a and GABA_BR1b,) and GABA_BR2 (**Fig.16B**), therefore providing the structural basis of a functional receptor assembly.

The ratio between R1a and R1b variants showed opposite tendency compared to mouse brain - there is a clear prevalence of the sushi domain- containing R1a subunit in ES cells, whereas the shorter R1b is the predominant form in the brain (Fritschy et al., 1999). These results were highly reproducible in independent experiments using RNA obtained from different ES cell cultures.

For immunofluorescent detection of GABA_A receptors in the ES cells, we used an antibody that recognizes the GABA binding β₃ subunit of the GABA_A receptor, a component of most abundant functional GABA receptors in the developing and mature brain. **Fig.17a** shows β₃ immunolabeling in all ES cells with more intense staining at the borders of neighboring cells (arrows).



The subcellular distribution of GABA_B receptors in ES cells was examined by immunofluorescent labeling of cultures with GABA_BR2 specific antibody, as this subunit is shown to be essential for correct trafficking of the receptor to the cell membrane and interaction with its cognate G protein. As shown **Fig.17b** positive staining was observed in all ES cells, while underlying fibroblasts were not labeled. High magnification revealed a pattern of bright “vesicular” appearance along the border of neighboring cells.

Fig.16. Expression of GABA_A(A) and GABA_B (B) receptor subunits in undifferentiated embryonic stem cells as revealed by RT-PCR. F- mitomycin-treated fibroblasts; ES- embryonic stem cells; - RT – reverse transcriptase omitted from first strand synthesis reaction; MB – mouse brain; *-actin.

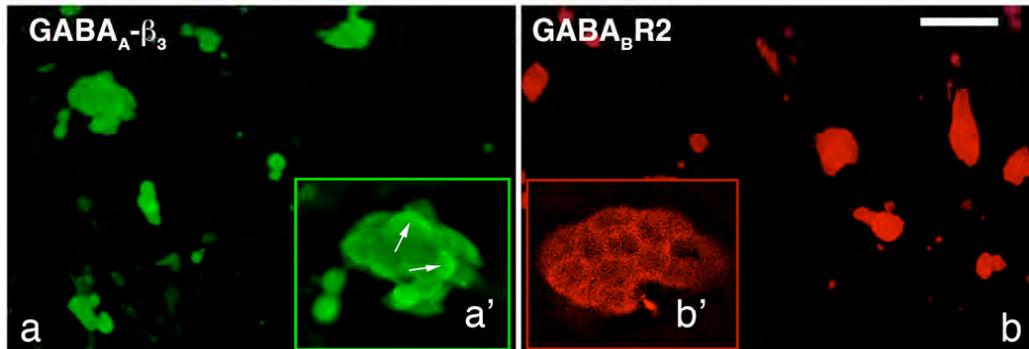


Fig.17. Immunofluorescent localization of GABA_A-β₃ (a) and GABA_BR2 (b) receptor subunits in mouse ES cell colonies. Insets a' and b' represent enlarged single clones of ES cells from a and b, respectively. Arrows in a' and arrowheads in b' mark the intensive signal at cellular borders. This enrichment of GABAR subunits at sites of putative contacts of adjacent cellular membranes suggests that both types of GABAR are engaged in intensive cell-cell communications within the ES colony. Scale bar: 50μm

1.5. ES cells express functional GABA_A and GABA_B receptors

In order to test the functionality of the GABA receptors, we examined the effect of receptor activation on the intracellular calcium levels, since we postulated that GABA action on ES cells could be mediated through calcium signals similar to immature neurons.

1.5.1. Expression of calcium signaling related molecular components in ES cells

The regulation of calcium (Ca²⁺) signaling depends on the interplay between intracellular and extracellular calcium sources (Berridge et al., 2003). Several molecules related to calcium signaling such as IP3 receptor (IP3R)-regulated intracellular Ca²⁺-stores, store-operated Ca²⁺-channels, and both plasma membrane Ca²⁺-pumps and Na⁺/Ca²⁺-exchangers have been described recently in undifferentiated ES cells (Yanagida et al., 2004; Kapur et al., 2007). Several other components, including G protein coupled inwardly rectifying potassium channels (GIRK1 and GIRK2) and voltage gated calcium channels (VGCC) have not yet been identified in ES cells, however they are known to be involved in GABA induced Ca²⁺ signaling in

the immature neurons (Owens and Kriegstein, 2000). We have checked for the presence of the above molecular components at mRNA level in undifferentiated ES cells. As shown in Fig.18, we could detect by RT-PCR the transcripts encoding Kir 3.1(GIRK1) and Kir3.2 (GIRK2) potassium channels (**Fig.18A**) as well as L, N, R, and T-type Ca^{2+} -channels (**Fig.18B**) in ES cells. Moreover, mRNA for two out of three members of ryanodine receptors, RyR1 and RyR2, mediators of intracellular calcium stores-dependent calcium efflux in early stages of embryonic development (Rossi and Sorrentino, 2002), were also detected (**Fig.18C**).

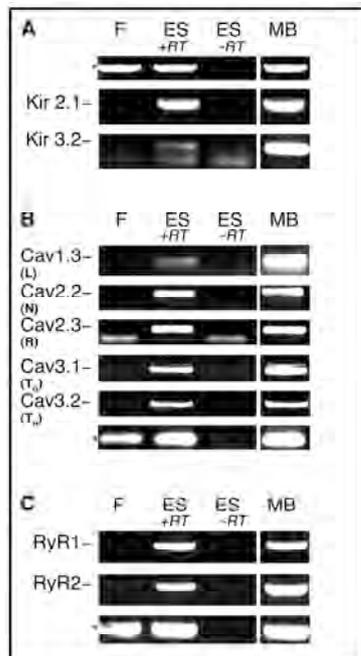


Fig.18. Expression of different inwardly rectifying potassium channels (A), voltage-gated calcium channels (B) and ryanodine receptors (C) in undifferentiated ES cells as revealed by RT-PCR. F- mitomycin-treated fibroblasts; ES- embryonic stem cells; -RT – reverse transcriptase omitted from first strand synthesis reaction ; MB – mouse brain; * - actin;

1.5.2. Activation of $GABA_A$ and $GABA_B$ receptors triggers rise in the intracellular calcium levels in ES cells

The presence of multiple subunits of $GABA_A$ R and the two subunits of $GABA_B$ R in undifferentiated ES cells indicated that the assembly of functional receptors is possible. To test this assumption we monitored the changes in $[Ca^{2+}]_i$ by confocal calcium imaging in ES cells upon application of GABA and selective receptor agonists/antagonists.

After 16-18 hours in culture, ES cells formed multicellular colonies exhibiting a distinct round boundary without signs of outgrowth. After loading of the Ca^{2+} -sensitive dye Fluo-4AM, we monitored in the presence of GABA or selective GABA_A and GABA_B receptor agonists/antagonists the changes of fluorescence intensity in individual cells within a colony, which correspond to changes of $[\text{Ca}^{2+}]_i$. Addition of GABA to the recording chamber (1 mM) for 90 s induced a sharp rise in $[\text{Ca}^{2+}]_i$ in a significant number of cells ($198.99 \pm 47.85\%$, $n = 38$ in 13 coverslips; **Fig.19**).

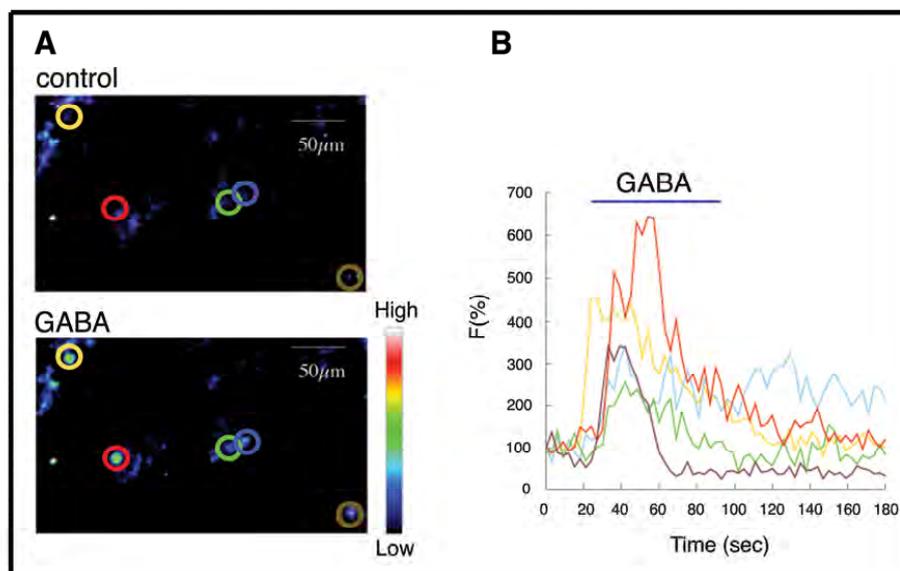


Fig.19. GABA induces calcium transients in undifferentiated mouse ES cells. **A:** Confocal images were taken at different time points of the recording shown in panels B. **B2** Representative recording of GABA evoked changes in fluorescence of Fluo4/AM-loaded ES cells **B:** Graph that shows changes in fluorescence intensity in cells circled on images in A panel. F(%): the change in intracellular Ca concentration is given as the change in fluorescence relative to the basal fluorescence at the beginning of experiment, which was normalized to 100%.

To characterize the GABA receptor type linked to the $[\text{Ca}^{2+}]_i$ increase evoked by GABA, the effect of selective GABAR agonists and antagonists were tested. As expected, muscimol (30 μM) a selective GABA_A receptor agonist induced sharp rise of $[\text{Ca}^{2+}]_i$ ($n = 31$ in 14 coverslips) with a $147,98 \pm 26,6\%$ increase in the amplitude of the response. This effect was reversally blocked by the selective GABA_A receptor antagonist, bicuculline (20 μM) ($n=0$ in 14 coverslips, $p < 0.05$), confirming that the effect of muscimol is due to GABA_A receptor activation (**Fig.20: A,D**).

To test whether the $[Ca^{2+}]_i$ increase induced by activation of the GABA_AR was due to Ca^{2+} release from intracellular Ca stores or/and influx through VGCC, we applied specific inhibitors to block each pathway.

Application of sarcoplasmic-endoplasmic reticulum Ca^{2+} -ATPase blocker CPA (10 μ M) resulted in Ca^{2+} transients when it reached to the perfusing chamber (n = 26 in 7 coverslips) and after 120.3 \pm 4.2 s-long incubation activated a capacitive Ca^{2+} entry (n = 15 in 7 coverslips, and data not shown), therefore the effect of muscimol was tested 60 s after CPA was applied. In the presence of CPA the occurrence of muscimol-evoked Ca^{2+} transients was greatly reduced (n = 4; p < 0.05), while their amplitude remained the same (126.55 \pm 48.11 %; **Fig.20: B,D**).

Initial bath application of the voltage-gated calcium channel (VGCC) blocker cadmium (100 μ M) significantly reduced the amplitude of the Ca transients (by 37%) but not the number of cells displaying calcium transients induced by subsequent addition of muscimol (93.14 \pm 13.68; 1.83 \pm 0.31; n=6; **Fig.20: C,D**).

The above findings show that GABA_A receptor activation evokes calcium release from internal stores and influx through VGCC.

Strikingly, GABA_BR agonist baclofen (20 μ M) evoked a sharp and significant increase in fluorescence in ES cells (2.70 \pm 0.32, n = 23; 177.35 \pm 29.88 %), that, in most cases, was wave-propagated to neighboring cells (**Fig.21: A,E**). Surprisingly, we could not block the Ca^{2+} transients induced by baclofen with two known GABA_BR antagonists- CGP 55845 (20 μ M) or CGP 54626 (10 μ M), which are widely used to block neuronal GABA_BR signalling. CGP 55845 affected neither the number of cells showing baclofen-induced $[Ca^{2+}]_i$ increase nor the amplitude of the evoked transients (3.0 \pm 0.71 and 210.84 \pm 47.73, respectively; n = 5, p > 0.05; **Fig.21: A,E**).

The amplitude of the calcium transients, induced by the application of KCl were further increased by addition of baclofen (**Fig.21B**) supporting the notion that in

contrast to neurons, the predominant effect of GABA_BR activation in undifferentiated ES cells is $[Ca^{2+}]_i$ rise.

GABA_BR have previously been shown to suppress (Bowery et al., 2002) or facilitate Ca^{2+} influx via several sub-types of VGCCs (Parramon et al., 1995; Carter and Mynlieff, 2004). Pretreatment of ES cultures with Cd^{2+} (100 μ M) a potent blocker of the VGCC reduced the amplitude of the baclofen-induced $[Ca^{2+}]_i$ transients, but not the number of responding cells (91.03 \pm 14.92 and 3.2 \pm 0.2, respectively, n=5, **Fig.21: C,E**). Furthermore, baclofen-evoked Ca^{2+} responses were essentially unaffected when Ca^{2+} -free buffer was used (2.33 \pm 0.67; 206.12 \pm 53.94; n=3), which clearly demonstrates that GABA_BR- induced Ca^{2+} transients are largely independent of extracellular Ca^{2+} (**Fig.21: D,E**).

In contrast to muscimol-induced calcium signals, those induced by the baclofen were nearly completely omitted in the presence of the endoplasmic-sarcoplasmic reticulum Ca^{2+} -ATPase blocker CPA (0.33 \pm 0.33; n = 3 p < 0.05; **Fig.21E**), indicating they originate from intracellular calcium stores.

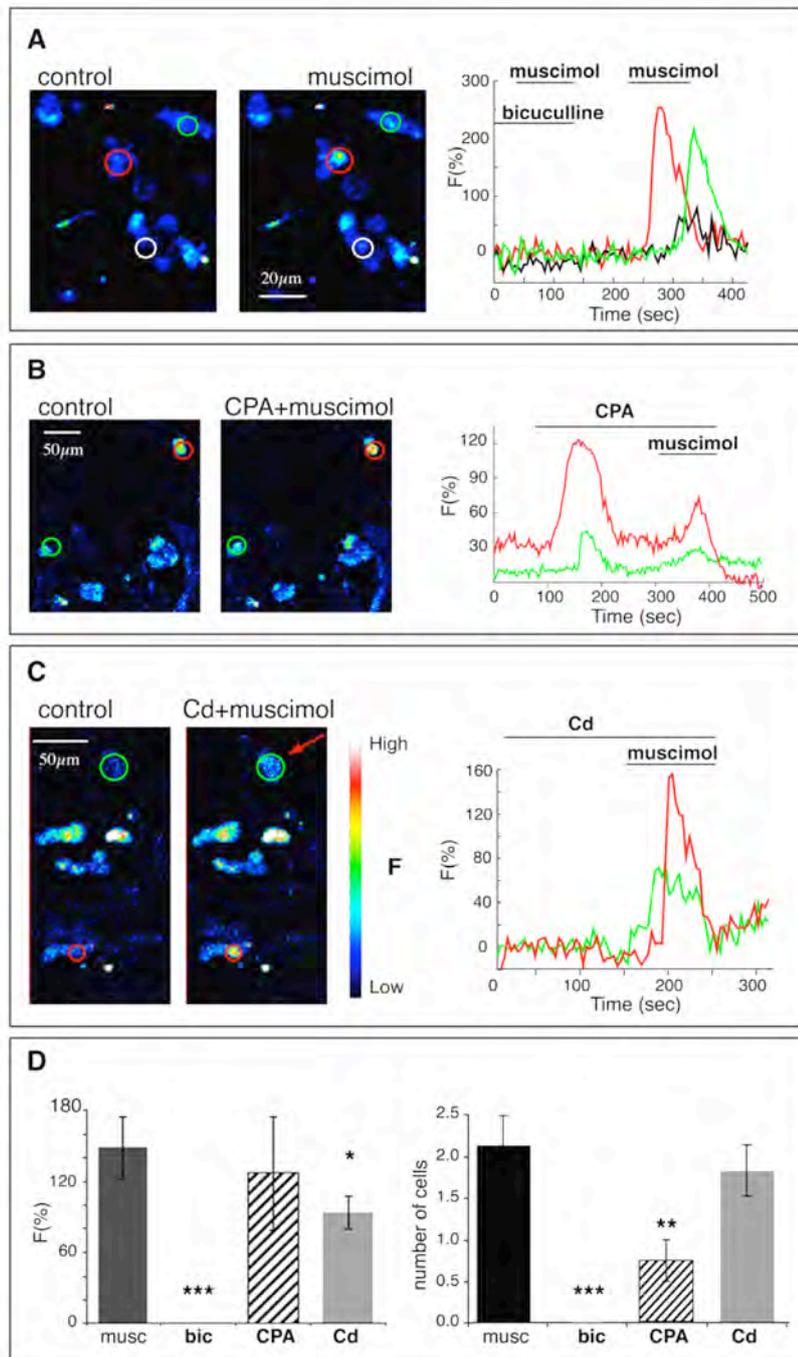


Fig.20. Modulation of agonist-induced GABA_A responses in ES cells. **A:** Representative recordings of GABA_A receptor-evoked changes in fluorescence of Fluo4/AM-loaded ES cells. Cultures were initially perfused with 30 μ M bicuculline, then 30 μ M muscimol and 20 μ M bicuculline were co-administered followed by application of 30 μ M muscimol (Muscimol) alone **B:** Effect of intracellular Ca²⁺ store blockage on muscimol-induced rise in [Ca²⁺]_i in ES cells. Cells responded to muscimol in the presence of cyclopiazonic acid (CPA), but with lower amplitude. **C:** Example of muscimol-induced [Ca²⁺]_i in ES cells after 3min pre-incubation with VGCC blocker cadmium (Cd). **Left panels:** Cells manifesting [Ca²⁺]_i transients are circled on the confocal images taken at different time points of the recording. **Right panels:** Change in fluorescence intensity (F(%)) in cells circled on corresponding images **D:** Comparison of GABA_A induced [Ca²⁺]_i responses (**musc**) in the presence of bicuculline (**bic**), SERCA blocker CPA (**CPA**) and VGCC blocker cadmium (**Cd**). Bar charts illustrating the amplitude of the responses (F(%) (left panel) and the average number of cells responding to the treatment (right panel). F(%): the change in [Ca²⁺]_i is given as the change in fluorescence relative to the basal fluorescence at the beginning of experiment (100%).

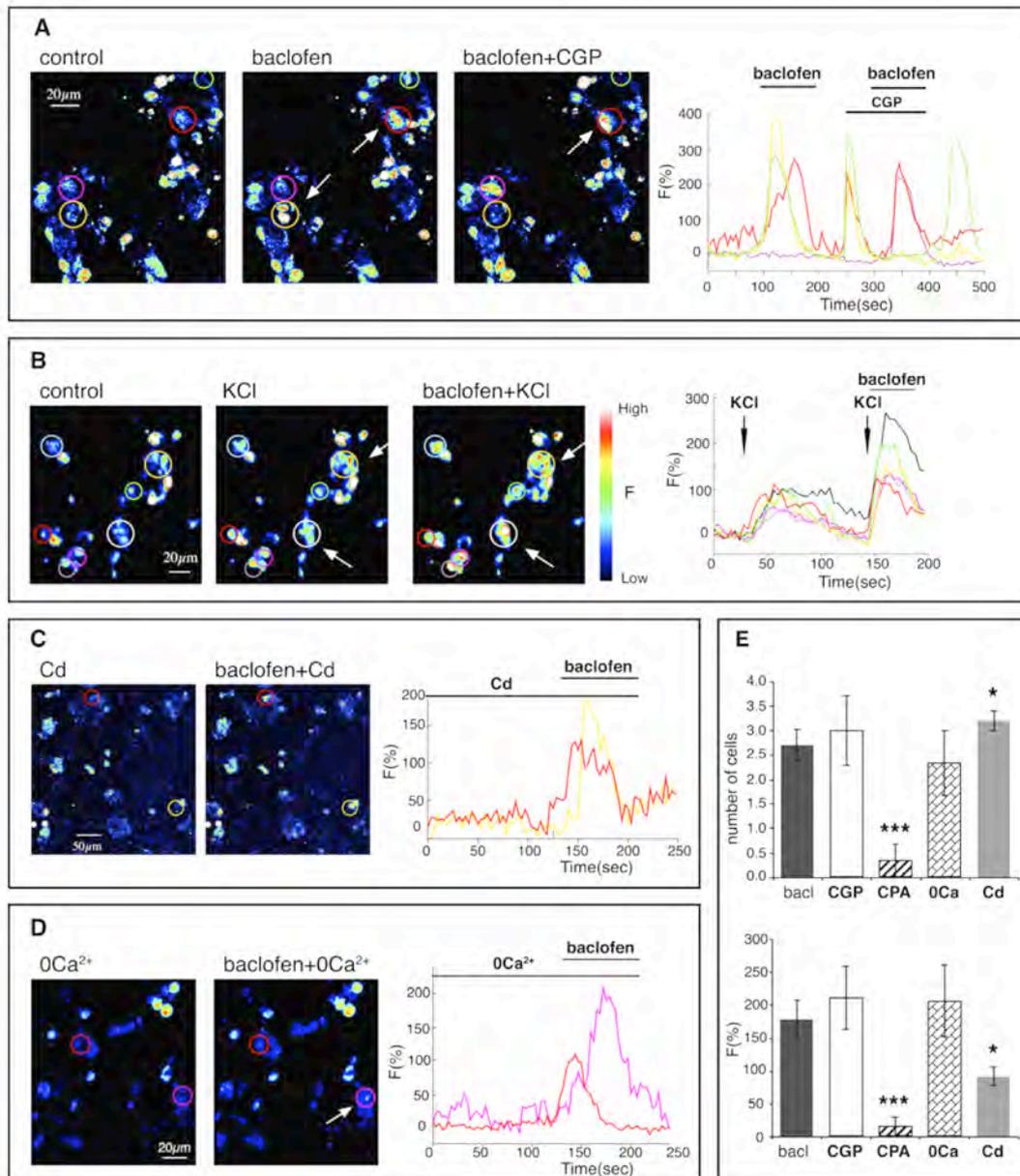


Fig.21. Modulation of agonist-induced GABA_B responses in ES cells **A:** Representative recordings of GABA_B receptor-evoked changes in fluorescence of Fluo4/AM-loaded ES cells. Cultures were initially perfused with 30 μ M baclofen, then 20 μ M CGP and finally baclofen and CGP were co-administered (baclofen plus CGP). **B:** Seven sample mean time courses of GABA_B - induced changes in [Ca²⁺]_i in ES cells after addition of KCl (KCl) followed by perfusion with KCl-baclofen (KCl + Baclofen); note that baclofen further increased the KCl-evoked responses. **C:** Example of baclofen-induced calcium transients in ES cells after 3min pre-incubation with VGCC blocker cadmium (Cd). **D:** Example of baclofen-evoked Ca²⁺ responses in Ca²⁺-free buffer. **Left panels:** Cells manifesting [Ca²⁺]_i transients are circled on the confocal images taken at different time points of the recording. **Right panels:** Change in fluorescence intensity (F(%)) in cells circled on corresponding images. **E:** Bar charts comparing the amplitude of the responses (left panel) and the average number of cells responding to the treatment (right panel). The change in intracellular Ca concentration is given as the change in fluorescence relative to the basal fluorescence at the beginning of experiment (100%).

1.6. GABA receptors modulates ES cell proliferation

Previously, it has been shown that GABA and different GABA receptor ligands can modulate the proliferation of neural stem cells of the embryonic ventricular zone (VZ) (Lo Turco et al., 1995; Haydar et al., 2000, Fukui et al., 2008). To test the effect of GABA activation on ES cell proliferation we plated ES cells at low numbers in multi-well plates, grew them under standard conditions for 18 hours to allow for stable membrane expression of the GABA receptor, thereafter added different agonists/antagonists to ES growth medium. It should be stressed that omitting of either LIF or serum from the medium, or use of dialyzed serum for prolonged incubation without application of drugs had an inhibitory effect over the ES cell growth and proliferation (data not shown).

Cell numbers were estimated 24/48 hrs following addition of the drugs using two different approaches. In proliferation tests, no statistically significant differences in the fluorescence intensity (proportional to cell number; see Materials and Methods) were encountered after one day-treatment with the muscimol, SR95531, baclofen (**Fig.22Aa**) or CGP (not shown). Exception was the GABA_AR blocker bicuculline, which proved highly toxic to the ES cells at concentration applied for blocking the GABA_A receptor (10-30 μ M), therefore we excluded it from later analysis and used instead the highly specific GABA_AR antagonist SR95531. Treatment of ES cells with these drugs for a further 24 hrs revealed a much more complex picture. While the GABA_AR agonist muscimol slightly, but not significantly decreased ES cell numbers compared to control, the GABA_BR agonist baclofen greatly increased the total cell numbers and size of clones. The effect of simultaneous treatment with both muscimol and baclofen was cumulative of the two drugs alone, i.e. cell numbers were close to control. Addition of the highly specific GABA_AR antagonist SR95531 to the media produced the strongest proliferative effect by nearly doubling the cell number

(average clone size) compared to control (**Fig.22Aa**). These results indicate that GABA affects proliferation of undifferentiated ES cells acting through both, GABA_A and GABA_B receptors.

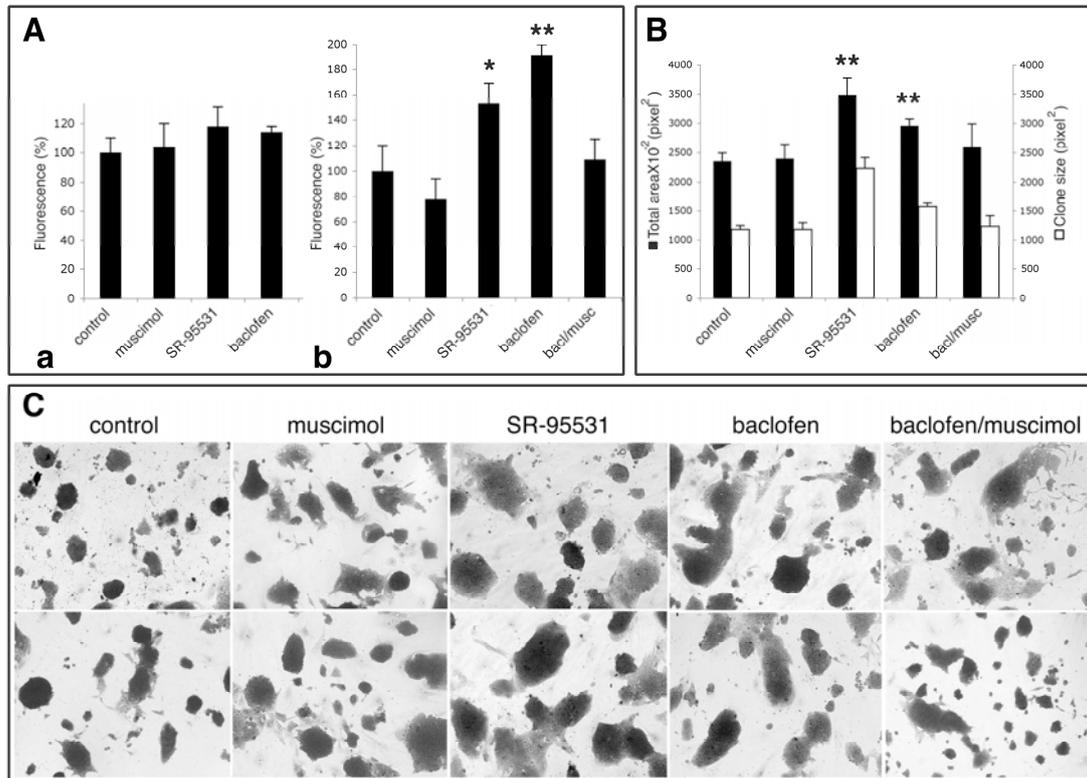


Fig.22. Effects of selective GABA receptors on the proliferation of undifferentiated ES cells. **A:** ES cells were incubated in the standard ES medium supplemented with 100 mM muscimol (Muscimol), 50 mM SR-95531 (SR-95531), 50 mM baclofen (Baclofen) or baclofen plus muscimol (Baclofen/Muscimol) for 18(a) and 24(b) hours and cell viability was determined using a fluorescence-based assay (CyQUANT Cell Proliferation Assay Kit; Molecular Probes, Invitrogen). Fluorescence in ES medium without additional drug treatment (Control) was considered to be 100%. **B, C:** Cellular proliferation was determined by measuring the size and total area of ES clones, which correspond to cell numbers. Cells were visualized with cresyl violet, images were captured with digital camera (Zeiss) and evaluated using NIH image software. Bar charts (B) and representative whole-field images of cells (C) treated as described above.

2. SEQUENTIAL EXPRESSION OF MOLECULAR COMPONENTS OF GABA SIGNALLING DURING IN VITRO NEURONAL DIFFERENTIATION OF MOUSE EMBRYONIC STEM CELLS RECAPITULATES EMBRYONIC DEVELOPMENT

In the present study, we examined the expression dynamics of different GAD forms GABA receptor subunits and GABA transporters during in vitro neuronal differentiation of mouse embryonic stem cells. We also characterized GABA synthesizing cells by immunocytochemistry. Our results demonstrate that neuronal progenitors synthesize different forms of GAD and contain GABA during earliest days of in vitro differentiation. Furthermore, the temporal expression profile of molecular components of GABA signalling machinery was highly consistent in our experiments and resembled its embryonic counterpart.

2.1. *Expression of different GAD isoforms during in vitro differentiation of ES cells*

According to the currently used induction protocol for *in vitro* neuronal differentiation (4-/4+; Bain et al., 1995), ES cells are initially cultured in suspension, in the absence of LIF, to induce ectodermal differentiation. The subsequent addition of high concentration of retinoic acid (RA; 10^{-7} - 10^{-6}) to the ES cell aggregates (embryoid bodies; Yamada et al., 1994; Leahy et al., 1999), induces rapid neuronal, and repress mesodermal gene expression in cells (Bain et al., 1996).

We have analysed the temporal expression pattern of different mRNAs encoded by GAD and *Dlx* genes during defined phases of ES cell neuronal differentiation by semi-quantitative RT-PCR analysis. Total RNA was isolated from ES cells, cell aggregates grown in suspension for 8 days (EB; 4-/4+ RA), and from attached EBs differentiated for 2, 4, 6, 8 and 10 days following induction period. The average of gene expression, relative to the actin expression, was calculated from at least three RT-PCR obtained from three independent in vitro differentiation experiments.

We detected a considerably high level of EGAD transcript in cells from all stages that was slightly increased on day 4 (dai4; **Fig.23**). In contrast, GAD67 mRNA was first detected in the cells grown for two days following RA (dai2) and gradually increased during maturation. Using the same PCR protocol condition as in the case of other genes, the expression of GAD65 mRNA was first seen in cells from one week old cultures (dai6), thereafter increased exponentially as they proceeded toward neuronal differentiation (**Fig.23**). However, GAD65 transcript could be amplified from ES cells (see Chapter 1.2.) to dai4 stage (not shown) after 35 amplification cycles.

In our cultures, both genes, Dlx2 and Dlx5 were expressed transiently, with partially overlapping patterns. The Dlx2 transcript was expressed, at a considerably high level, in ES cells and EBs, then increased between dai2-dai4, and decrease after dai6, being undetectable beyond the 8th day after RA-induction. Expression of Dlx5 mRNA was detectable for the first time at dai2, slightly increased by dai4 and remained practically unchanged until dai8 after which sharply declined.

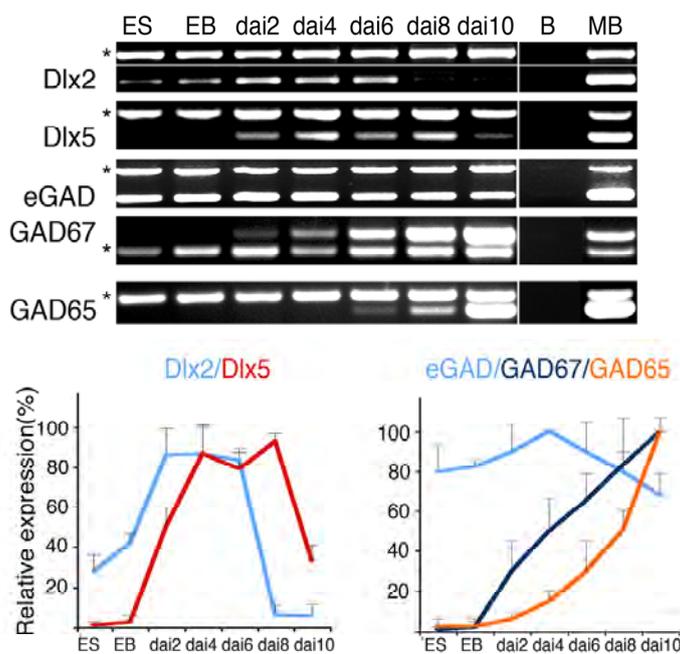


Fig.23. Temporal expression patterns of Dlx2, Dlx5 and GAD gene family in ES cells toward neuronal differentiation as revealed by semi-quantitative RT-PCR. A: Representative RT-PCR amplifications of Dlx2 and Dlx5 and different GAD (EGAD, GAD67 and GAD65) transcripts using cDNA synthesized from total RNA preparations of undifferentiated (ES) and RA-induced ES cells (EB, dai2-dai10), and adult brain (MB). Expression levels were normalized to β -actin, (*) and represented on a grey scale as % of max level. B: Plots (means \pm SEM) of relative expression level of each gene calculated from three independent experiments normalized to the maximum expression level of co-amplified β -actin (100%).

The GAD proteins were identified on Western blots of cell homogenates from ES cells and in RA-induced ES cells. The GAD25 protein, previously shown to run as

two closely spaced bands (32-32kD), (Szabo et al,1994), were expressed, at considerably high level in ES and EBs, up-regulated from dai2 to dai8 and thereafter declined but still detectible at dai 10 and beyond showing similitatities to embryonic expression (**Fig.24**). The GAD44 protein was expressed at very low level in ES cells, but then increased gradually and remained stably expressed throughout the studied period. The translated GAD67 and GAD65 were first detected, at the protein level, at later differentiation periods, around dai6 and dai8, respectively, which matches with the RT-PCR results.

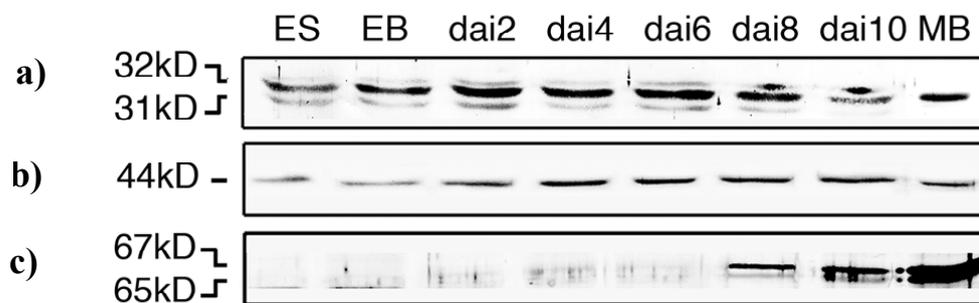


Fig.24. Western blot analysis of GAD forms toward neuronal differentiation of ES cells: protein homogenates from cells isolated from the defined stages of neuronal differentiation (EB, dai2-dai8) were run on 12% SDS-PAGE, blotted and stained with an anti-GAD rabbit serum #8876 (a) and #6799 (b,c).

2.2. Expression of different GABA receptor subunits and GABA transporters

We found all examined GABA_A receptor subunits to be expressed during *in vitro* neuronal differentiation of ES cells. The mRNAs for α 3 and α 5 subunits could be detected already in ES and EBs, whereas α 2, α 4, and α 1 subunits were detectible after induction of neuronal differentiation (dai2), respectively (**Fig.25**). The β 1 subunits showed a low level of expression in the EBs, and gradually increased from day 4, while β 2 could be detected for the first time at dai2 with reasonable high level of expression that stayed practically unchanged until day 10. β 3 transcript, which is the predominant β subunit expressed during embryonic development (Laurie et al.,1992; Wisden et al.,1992; Serafini et al.,1998) had high level of expression in ES cells and

EBs and remained constant until dai10 (**Fig.25**). γ 1 and γ 2L expression paralleled the neuronal differentiation from dai2 on. γ 2S and γ 3 mRNA displayed relatively high levels of expression at all stages and was increased after RA-treatment.

Using specific primer pairs, designed for each subunit type, we investigated the expression of GABA_BR1(GABA_BR1_a and GABA_BR1_b) and GABA_BR₂ receptor subunits during neuronal differentiation. All three mRNAs were detected already in ES cells and EBs and gradually increased during differentiation (**Fig.25**). The sushi-domain containing GABA_BR1a subunit is the most abundant of all three.

Next, we examined the dynamics expression of different plasma membrane (GATs) and the vesicular (VGAT) GABA transporters. GAT-1 and VGAT are the first transporters to be induced as they are expressed in ES cells and parallel the neuronal differentiation while GAT4, GAT3 and GAT2 were detected after dai2 and throughout the differentiation period (**Fig.25**).

The developmental switch of GABA action from excitation to inhibition is due to reduction in intracellular chloride concentration that correlates with the induction of KCC2, the chloride potassium K⁺-Cl⁻ cotransporter, and down regulation of NKCC, the sodium potassium chloride Na⁺K⁺Cl⁻ exchanger (Ben-Ari., 2002). As expected, NKCC1 was detected in all stages studied here but gradually increased during differentiation, while KCC2 transcript could be hardly detected initially, but was upregulated in cultures after RA treatment (**Fig.25**).

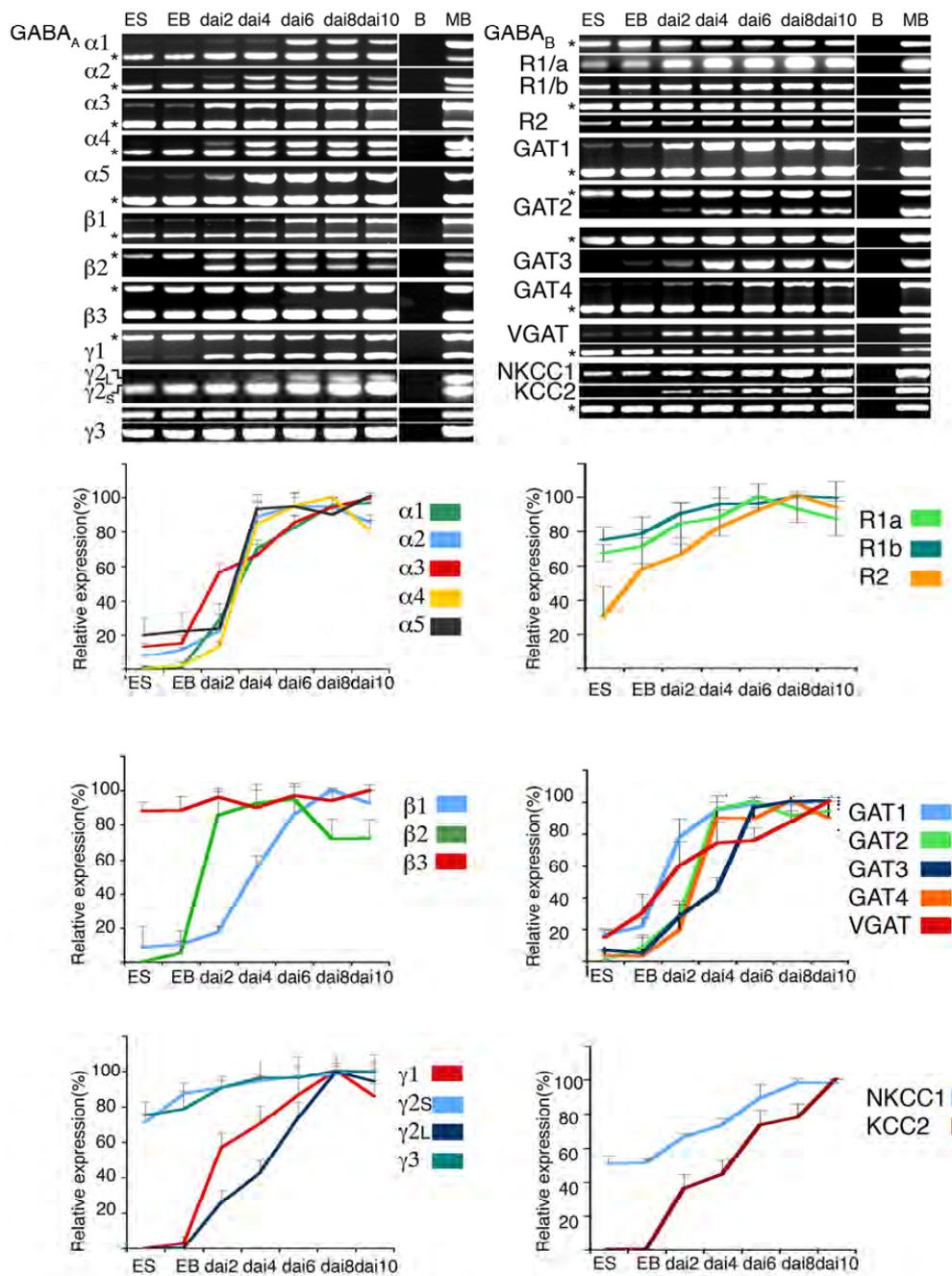


Fig.25. Temporal expression profile of GABA signaling components during neuronal differentiation of ES cells. A: Expression of GABA_A and GABA_B receptor subunits and GABA transporters as revealed by semi-quantitative RT-PCR. **Upper panel:** Representative gel images showing the PCR products amplified from reverse-transcribed total RNA that was isolated from from total RNA preparations of undifferentiated (ES) and RA-induced ES cells (EB, dai2-dai10), and adult brain (MB). Specific primer pairs derived from different exons and product size for each cDNA are listed in Table 1. β-actin (*) was co-amplified with the target genes. **Lower panel:** Relative expression levels of GABA receptor subunits and GABA transporters. Expression levels were normalized to β-actin, and represented on a grey scale as % of maximum level. **B:** Charts represent comparison of developmental expression profiles of different GABA signaling components. Expression levels were calculated from three independent experiments and normalized to β-actin and were expressed as a % of the maximum expression in each group.

2.3. Development of GABA immunoreactivity in ES-derived neurons during *in vitro* differentiation

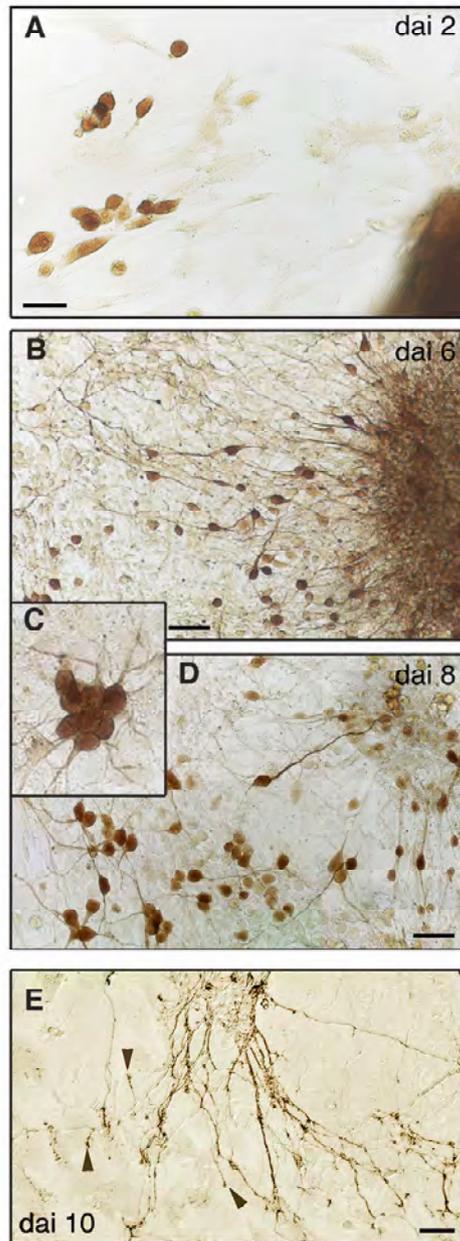


Fig.26. Immunocytochemical detection of GABA in cells at different stages of neuronal differentiation. Images were taken on second (dai2; A), sixthth (dai6;B), eighthth (dai8;D) and tenth (dai10;E), respectively. Arrows show GABA positive growth cones. Scale bars:20µm in A, 40µm in B,D and E.

GABA immunoreactivity was examined during different stages of *in vitro* neuronal differentiation of ES cells. At the earliest time point used, two days after plating, cultures consisted mostly of flat, fibroblast-like cells and few long neurites protruding from EBs that were not immunopositive for GABA. However, in some areas we detected single cells, with migratory appearance, displaying strong staining in both perikaria and short, branched processes (**Fig.26A**). In the following days, cells start to migrate in large numbers from EBs, GABA immunoreactivity increased drastically in both cell bodies and neurites. By day 6, most of immunopositive cells, ranging from bipolar to multipolar neurons, formed long dissimilar neurite-like processes radiating from EBs, also GABA⁺ (**Fig.26B**), while some of the young neurons formed tight clusters containing 10-30 cells (**Fig.26D**) or fascicles (not shown).

In cultures fixed 8 days after induction period we could not detect any GABA immunoreactivity in the EBs. The labeled neurons with morphology of migrating cells had longer neurites and became progressively spread out in the culture (**Fig.26D**). By day 10 after RA induction the morphology of GABA-containing neurons changed dramatically by appearance of long, bundled processes (**Fig.26E**). Growth cones were also strongly labeled for GABA: some of them had a large number of filopodia, others had multiple long extensions arising from a long neurites (arrows). Flat cells with appearance of fibroblasts or glia in our cultures were not immunoreactive for GABA.

3. GABA SIGNALING IN THE DEVELOPING MOUSE LENS

Earlier work from this laboratory demonstrated, by using GAD *in situ* hybridization, that mRNAs for both GAD65 and GAD67 are present in the elongating lens fiber cells of mid-gestation mouse embryo (Katarova et al., 2000). Here we examined the temporal and spatial expression of different GAD forms, GABA receptor subunits and GABA transporters (vesicular VGAT and membrane GATs) in the developing mouse lens and in primary lens epithelial cultures (LEC) and present evidence about the functionality of GABA signaling in the developing mouse lens.

3.1. GAD mRNAs are sequentially expressed in the developing lens

The temporal expression pattern of different mRNAs encoded by the two GAD genes was determined during lens development by semi-quantitative RT-PCR analysis using first-strand cDNA from embryonic (E14.5, E15.5, E16.5, E17.5) and postnatal (P0, P7, P14, P30) mouse lenses. Embryonic stage E14.5, when the lens is composed of epithelium and predominantly primary lens fibers, was chosen as a starting point due to technical limitations.

We found that all three types of GAD mRNAs were already present in E14.5 lenses, but had distinct expression profiles during later stages of lens development (**Fig.27**). The embryonic GAD transcripts encoded by the GAD67 gene (EGADs) were already expressed at a considerably high level at E14.5, then slightly increased in intensity with a maximum level of expression at E16.5, declining sharply thereafter and were barely detectable postnatally. On the contrary, GAD67 was expressed at a constant low level until birth (P0), with a slight peak at E16.5, and then gradually increased, peaking postnatally at P14-P30. Similar to EGADs, GAD65 mRNA level was already high at the primary fiber stages (E14.5) peaked at E15.5 then decreased with a sharp decline from E17.5 and was undetectable at P14 and beyond (**Fig.27**).

We also examined for the presence of Dlx2 and Dlx5, two transcription factors that

overlap with expression domains of GAD (Katarova et al., 2000) and considered to be the upstream regulators of GAD. Our semi-quantitative RT-PCR showed that similarly to the forebrain, Dlx2 expression in the developing lens was induced before Dlx5. As shown in Fig.23, Dlx2 mRNA levels were high and remained almost unchanged throughout E14.5, E15.5 and E16.5, after which Dlx2 expression declined and was undetectable in P7 and older lenses. Dlx5 mRNA, on the other hand, was barely detectable at E14.5 then gradually increased between E15.5 and P0, reaching a maximum at around birth (E17.5-P0). In the postnatal lens Dlx5 was down - regulated and was not detected beyond P30 (*Fig.27*).

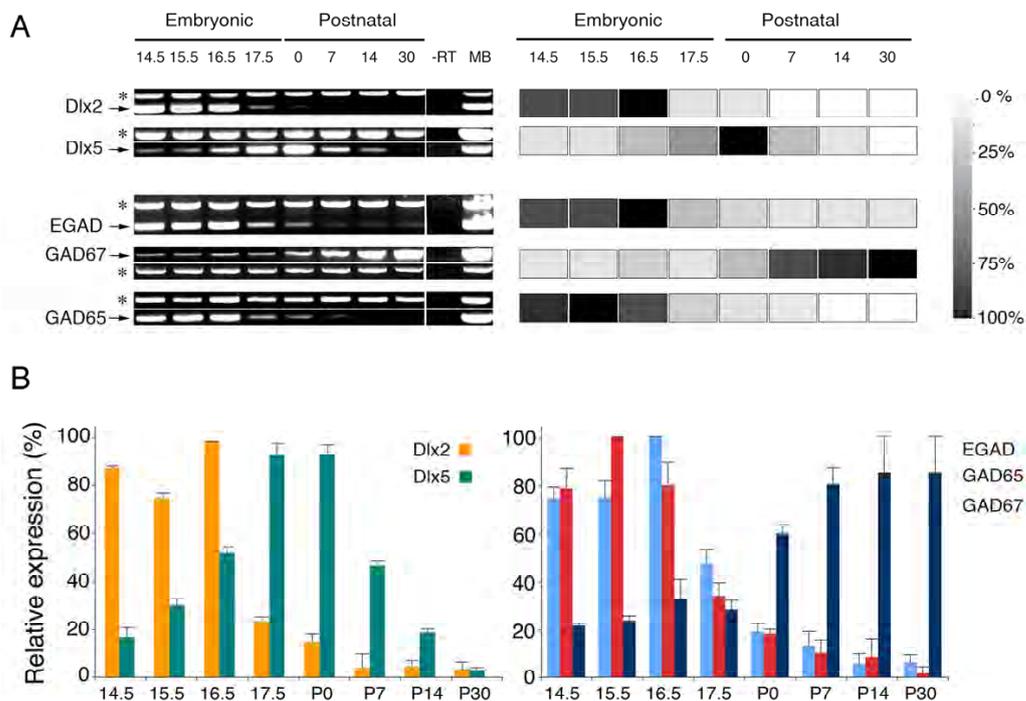


Fig.27. Temporal expression patterns of Dlx2, Dlx5 and GAD gene family in developing mouse lens as revealed by semi-quantitative RT-PCR. A: Representative RT-PCR amplifications of Dlx2 and Dlx5 and different GAD (EGAD, GAD67 and GAD65) transcripts using cDNA synthesized from total RNA preparations of E14.5-P30 lenses and adult brain (MB). Expression levels were normalized to β -actin (*), and represented on a grey scale as % of max level. B: Histograms (means \pm SEM) of relative expression level of each gene calculated from three independent experiments normalized to the maximum expression level of co-amplified β -actin (100%).

3.2. *GABA_A and GABA_B receptor subunits and GABA transporters are abundantly expressed in the developing lens*

We determined the temporal expression profile of mRNAs encoding different GABA_A subunits, the two GABA_B receptor subunits, the vesicular and membrane GABA transporters (VGAT and GAT1-4, respectively) during lens development (stages E14.5-P30) by semi-quantitative RT-PCR (**Fig.28**). Out of the six α subunits (α_{1-6}) amplified using specific oligo-pairs, we failed to detect α_5 and α_6 at any stage during lens development (not shown). α_2 and α_3 subunits were detected at the earliest stages (E14.5-E15.5) and were abundantly expressed until P0, thereafter α_2 was present at low levels. Subunits α_1 and α_4 were virtually undetectable at E14.5-E15.5. While α_1 expression level rose continuously from E16.5, α_4 elevated steeply at E16.5 and remained at constant high-level expression thereafter.

Of the three known β subunits, β_1 displayed only slightly over-background expression throughout all studied stages with the exception of P0 when it was up-regulated to a small degree. This unusual profile was consistent in multiple experiments (data not shown) and may reflect a special need for this subunit at birth. β_3 was by far the most abundant subunit at embryonic stages E14.5-E17.5, but was dramatically reduced after P0. The GABA_AR β_2 subunit showed low expression at earlier (E14.5-E15.5) and late (P14-P30) stages and a slightly higher expression at E16.5-P7 (**Fig.28**).

All three GABA_AR γ subunits and the δ subunit (a substitute for γ) were expressed from the earliest studied stages. γ_1 is predominantly prenatal and almost undetectable beyond P0 with the peculiar exception of P30, while γ_2 , γ_3 and δ were present at all studied stages. Interestingly, γ_3 subunit is by far the most abundant of all γ subunits during all stages of lens development, which contrasts its highly restricted expression in the brain (Laurie et al., 1992). We failed to detect the retina-specific ρ_1

and $\rho 2$ subunits (not shown), which demonstrates that all RT-PCR reactions are strictly lens-specific.

The functional GABA_B receptor is an obligatory heterodimer comprised of two subunits: GABA_BR1 harboring the GABA binding site and GABA_BR2, responsible for the dimer formation (Bowery and Brown, 1997; Kaupmann et al., 1997). While both R1 and R2 were expressed in the developing lens, they displayed strikingly different expression profiles (**Fig.28**). GABA_BR1 showed strong and uniform expression throughout all stages, whereas GABA_BR2 subunit was expressed at moderate levels at E15.5-E17.5 peaking at E16.5 and was hardly detectable at earlier and later stages.

Transcripts for all GATs and VGAT were abundant and showed distinct profiles during different stages of lens development (**Fig.28**). The predominantly neuronal-specific GAT1 showed a much higher expression around birth (E17.5-P0) than at earlier or later stages. GAT2 and GAT3 found in both neurons and glia were hardly detectable at E14.5-E15.5, but were upregulated at the later stages. The GAT4 was nearly uniformly expressed throughout all stages being slightly increased at E15.5-E16.5. VGAT was much more abundant before birth, showing a transient peak at E17.5, but was greatly reduced after birth. These patterns clearly followed different stages of fiber cell differentiation.

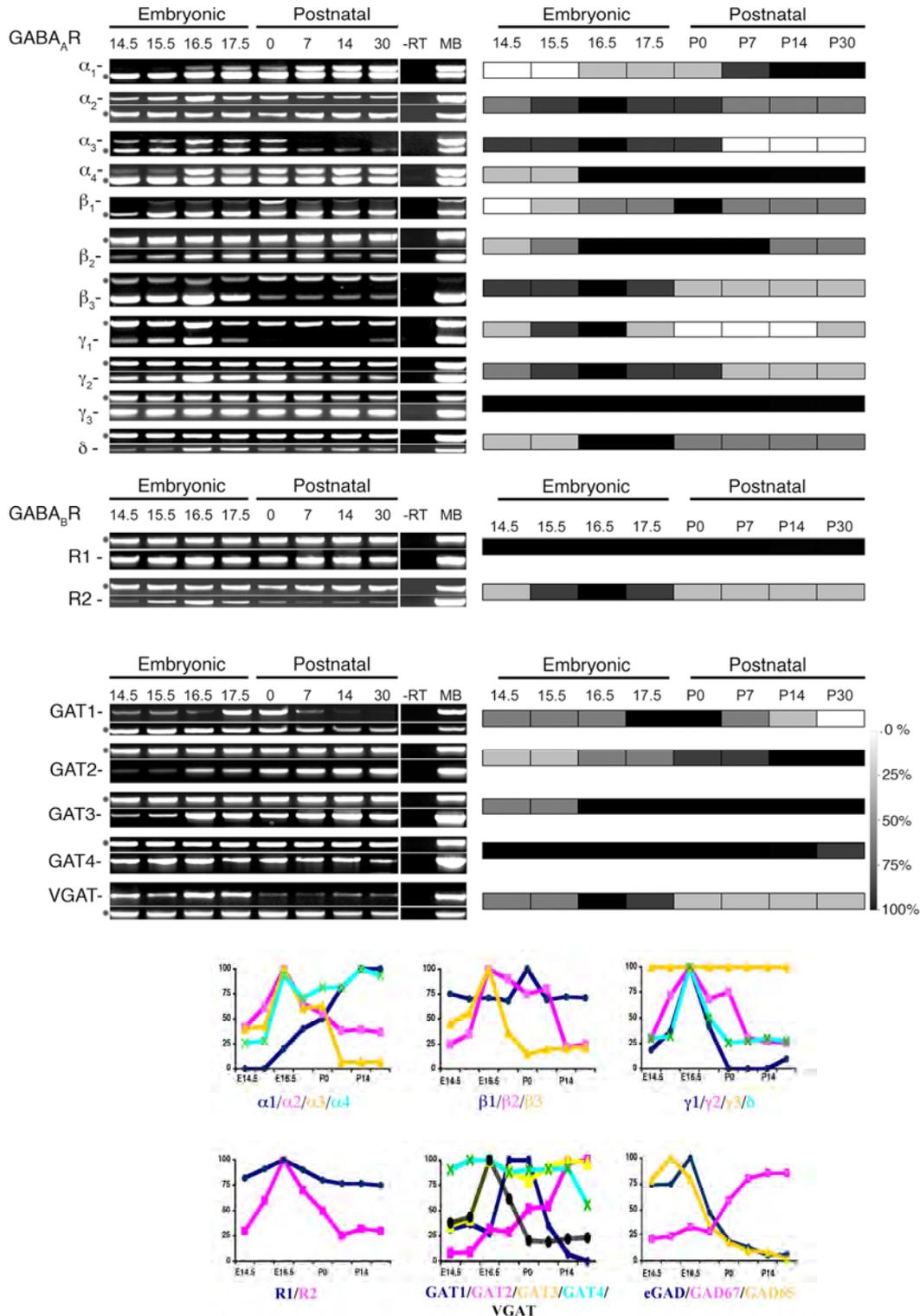


Fig.28. Temporal expression profile of GABA signaling components in the developing mouse lens. A: Expression of GABA_A and GABA_B receptor subunits and GABA transporters as revealed by semi-quantitative RT-PCR. **Left panel:** Representative gel images showing the PCR products amplified from reverse-transcribed total RNA that was isolated from embryonic (E14.5-17.5) and postnatal (0, 7, 14 and 30 days) lenses and adult brain (MB). Specific primer pairs derived from different exons and product size for each cDNA are listed in Table 1. β-actin (*) was co-amplified with the target genes. **Right panel:** Relative expression levels of GABA receptor subunits and GABA transporters, as assayed by the fluorescence intensity of the PCR products (left panel). Densitometry analysis was performed using IMAGE (NIH, v. 1. 34.). Expression levels were normalized to β-actin, and represented on a grey scale as % of maximum level. **B:** Charts represent comparison of developmental expression profiles of different GABA signaling components. Expression levels were calculated from three independent experiments and normalized to β-actin and were expressed as a % of the maximum expression in each group.

3.3. Spearman correlation identifies similar expression profiles of different GABA signaling components during lens development

The striking similarities between expression patterns of different GABAergic components prompted their grouping according to developmental stage (embryonic, fetal, neonatal and postnatal; **Fig.29A**). During embryonic stages mostly GAD65 and the embryonic GAD transcripts are expressed and this is paralleled by predominant expression of VGAT and the membrane GAT4. The prevalent configuration of the GABA_AR at this stage is $\alpha_{2,3,4}/\beta_3/\gamma_{2,3}$. This receptor type predominates in the spinal cord at early embryonic stages (Ma et al., 1993) which points to the existence of common features of development of the GABA signaling pathway. In the fetal lens most of the GABA signaling components are greatly up-regulated in correlation with the increased speed of generation of lens fibers during this period (Kaufman, 1993). GAD65 and EGAD continued to predominate, paralleled at the beginning by VGAT and GAT4, and later by GAT3 and GAT1 (**Fig.29A**). The predicted GABA_AR pentamer during this period is $\alpha_{2,3,4}/\beta_{2/3}/\gamma_{1,2,3}$ with all these subunits highly expressed.

Around birth (neonatal stage), GABA_AR β_1 and GAT1 were selectively upregulated and the embryonic/fetal components were suppressed.

The postnatal stage is clearly characterized by fewer and highly expressed components. Remarkably, these stages are predominated solely by GAD67, while GAD65 is last detected, at extremely low levels, at P14. The GAD65-GAD67 switch is paralleled by an almost complete down-regulation of VGAT, GABA_AR α_2 , α_3 , β_1 , β_3 , γ_1 , γ_2 , δ , GABA_BR2 subunits, up-regulation of GAT1,2,3 and the GABA_AR α_1 subunit, which is predominantly synaptic in brain.

These findings gained further support from the Spearman correlation analysis (Kotlyar et al., 2002; **Fig.29: B1-B4**), which clearly showed high expression

correlation between the predominantly prenatal GAD65 and EGAD ($r_s=0.94$), GAD65-GABA_AR α_2 ($r_s=0.80$), GABA_AR α_2 -GABA_AR γ_2 ($r_s=0.86$) and GAD65-VGAT ($r_s=0.67$). GAD67 expression was highly correlated with that of GAT2 ($r_s=0.99$) and GABA_AR α_1 ($r_s=0.96$; Fig. 29B5, B6), all expressed postnatally.

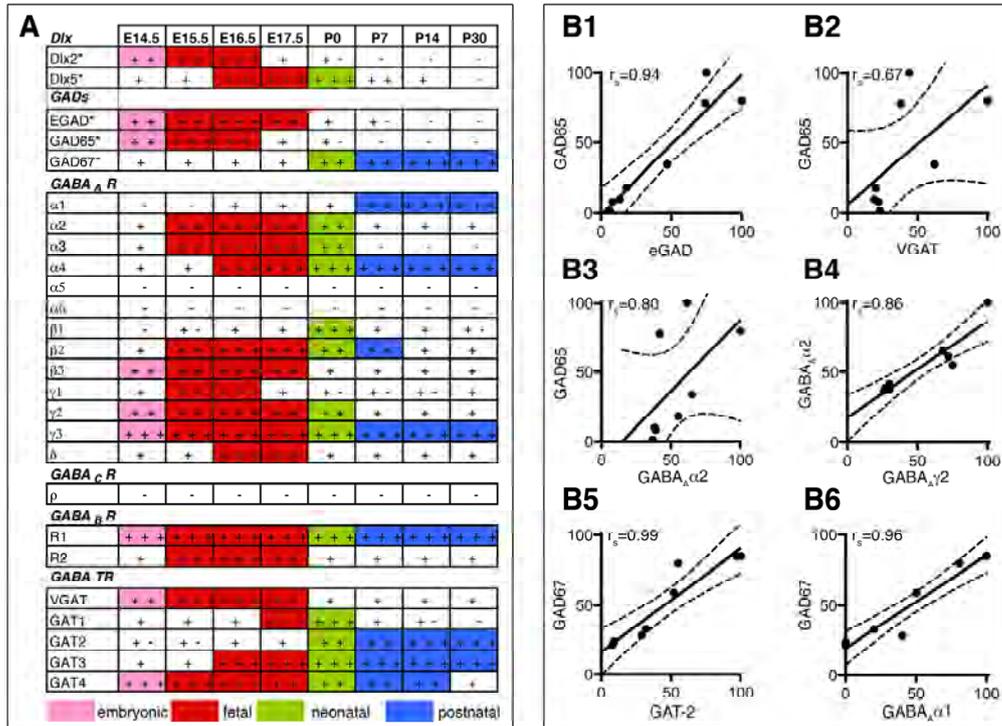


Fig.29. Developmental stage-specific expression profiles and coordinated expression of genes of GABA signaling components in the developing mouse lens. **A:** Schematic presentation of temporal expression profiles based on RT-PCR data in Fig.L1A and L2A Relative expression level of each gene at different developmental stages (E14.5-P30) indicated as follows: +++: high; ++: medium; +: low; -: no expression. Developmental stages: embryonic \leq E14.5; fetal: E15.5-E17.5; P0: neonatal; postnatal: P7-P30. **B:** Coordinated expression of GAD, GABA receptor subunits and transporters: eGAD vs. GAD65 (**B1**), VGAT vs. GAD65 (**B2**), GABA α_2 vs. GAD65 (**B3**), GABA α_2 vs. GABA α_2 (**B4**), GAT-2 vs. GAD67 (**B5**) and GABA α_1 vs. GAD67 (**B6**). Pair-wise correlation analysis of relative gene expression levels at different developmental stages was performed using Prism software. Spearman's rank correlation coefficient (r_s) for each gene pair is indicated on the scatter plots. The best fit (solid line) is shown with 95% confidence intervals (dotted lines). Numbers on x- and y-axes represent % of relative gene expression levels.

3.4. Temporal expression of GAD proteins and GABA in the developing lens

We used Western blot and HPLC analysis to study the synthesis of GAD proteins and GABA content through E15.5 embryonic-P30 postnatal stages. The protein homogenates from E15.5-P30 lenses were run on SDS-PAGE gels, blotted at conditions optimized for each isoform and stained with a pan-GAD rabbit serum #6799 (Katarova et al., 1990; Szabo et al., 1994). β -actin was used to monitor the amount of protein applied on the gel. P30 retina homogenate expressing high levels of

all GAD isoforms was run in parallel (**Fig.30A: P30-R**). In agreement with the RT-PCR results, GAD65 was more abundant at earlier stages (E15.5-E17.5) peaking at E17.5, thereafter its level quickly dropped and was not detected at P30. In contrast, GAD67 appeared first as a faint band at P0, slightly increasing until P30 (**Fig.30A**). EGAD (GAD25 and GAD44) showed a transient temporal expression pattern similar to that characteristic for the embryonic nervous system (Szabo et al., 1994): it was low at the earliest stages (E15.5), peaked at E17.5, gradually decreased until P14 and was undetectable beyond this stage (**Fig.30A**).

The GABA content of the lens at different developmental stages showed a clear correlation with the temporal profile of GADs: it was low at earlier stages, increased considerably and peaked at E17-P0, then leveled to 30% of P0 value at the postnatal stages (**Fig.30B**).

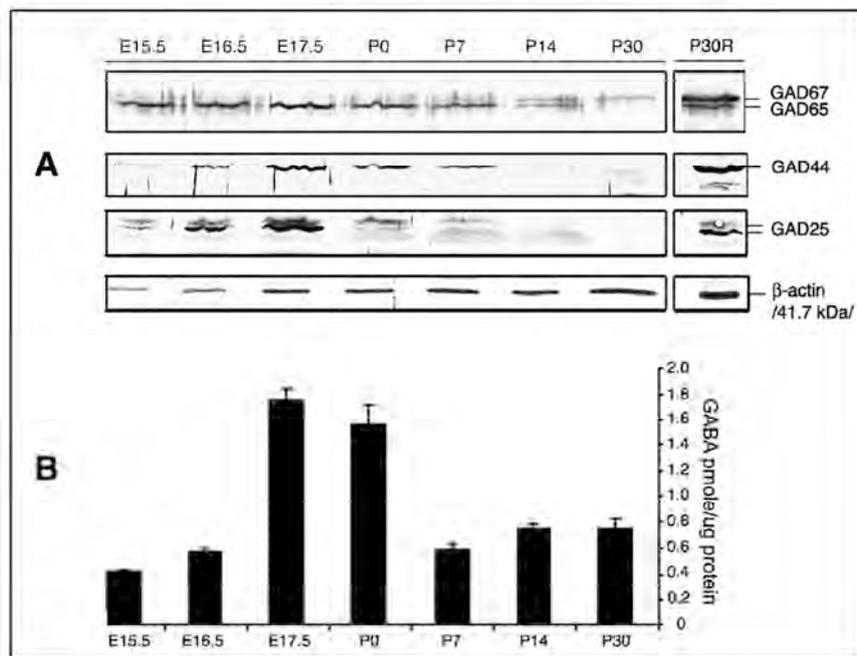


Fig.30. GAD proteins and GABA content in the developing mouse lens at stages E15-P30. **A:** Western blot analysis of GAD forms in the eye lens: protein homogenates from lenses isolated from the different stage embryonic lenses next to P30 retina (P30R) were run on 10% SDS-PAGE, blotted and stained with an anti-GAD rabbit serum #6799. β -actin was stained as control. **B:** Analysis of GABA content in the developing mouse lens: GABA concentration was measured in E15.5-P30 lens homogenates by HPLC and expressed in pmole/ μ g protein. Bars represent means of three measurements \pm SEM. GABA measurements by HPLC were done in collaboration with Mária Baranyi.

3.5. GAD protein expression in the developing lens as revealed by immunohistochemistry: spatial segregation of different GAD forms

The spatial pattern of GAD65 expression in embryonic lenses was revealed by immunostaining with the GAD65-specific N-65 antibody of E9.5-E17.5 eyes (**Fig.31: a,b,c,d,e**). Strong GAD65 staining was first visible in the thickened surface ectoderm of the lens placode (**Fig.31a**) and persisted through the lens pit (E10.5; **Fig.31b**) and lens vesicle (not shown). At E12.5, strong labeling was detected in both, prospective lens epithelium and elongating cells of posterior wall - prospective primary lens fibers (**Fig.31c**). The core lens fibers were still prominently stained at E14.5 (**Fig.31d**), thereafter the intensity was reduced (E17.5; **Fig.31e**). Faint GAD65 expression was still detected in P0 and P7 lens nucleus, but disappeared at later stages (not shown).

To detect the GAD67 gene-encoded GAD isoforms we used an anti-GAD antibody (serum #6799), which was raised in our laboratory against recombinant mouse GAD67, but also recognizes the embryonic GAD25 and GAD44. This antibody recognizes to a lesser extent GAD65 on Western blots, but not on tissue sections under the conditions used in this study. Based on our RT-PCR and Western blot results we could conclude that the pattern obtained with serum #6799 reflects the expression of EGAD (GAD25 and GAD44) mRNA in embryonic and predominantly that of GAD67 in the postnatal lens. As shown on **Fig.31a'**, GAD67 staining was already detected in the lens placode, an area of a thickened surface ectoderm and the underlying optic vesicle at E9.5. At E10.5, uniform immunolabeling was observed in the lens cup (**Fig.31b'**). At E12.5, strong staining was observed in both the posterior wall, containing elongating primary fibers, and epithelial cells at the anterior pole, which remain undifferentiated and retain their proliferation capacity (**Fig.31c'**).

From E14.5 on, the GAD67-related proteins were localized in both primary and secondary fiber cells with strongest labeling at E16.5-E17.5 (**Fig.31e'**). This pattern

was preserved until birth. In postnatal lenses, a gradual change could be followed from moderately strong intensity to complete absence of staining (P30) of the lens core concomitant with increase in immunolabeling of elongating secondary fibers at the equatorial region (EQ in P14, P30).

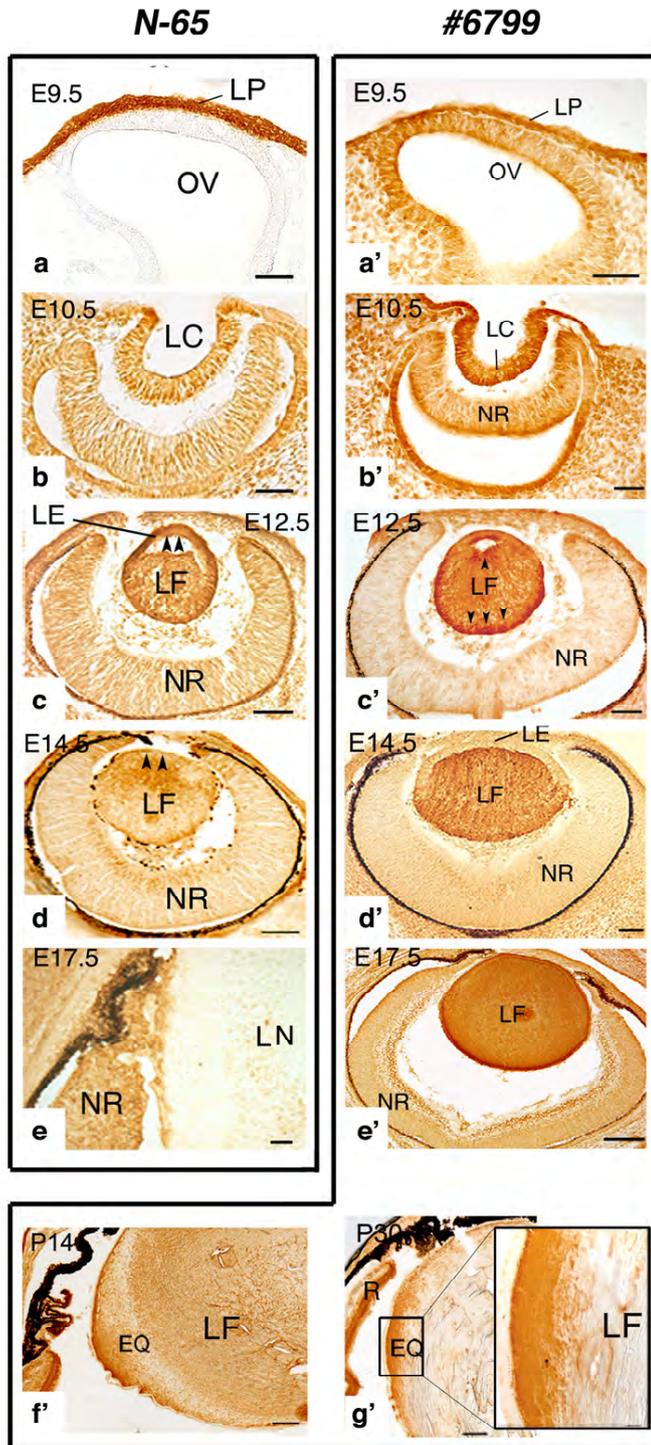


Fig. 31. Expression pattern of GAD proteins in the developing mouse lens.

a-e: coronal sections from different stage embryos stained with anti-GAD65 antibody N-65. GAD65 protein was localized initially to the lens placode, lens vesicle and later to lens primary fiber cells. Arrowheads point to lens epithelium **a'-g'**; Coronal sections from E9.5-P30 eyes were stained with anti-GAD serum #6799. Not that immunoreactivity during embryonic stages is due to GAD25 and GAD44; only adult GAD67 is expressed beyond P7. GAD67-related immunoreactivity was detected from the lens placode stage, thereafter was localized to the lens pit, primary fibers and central epithelium. **LC-** lens cup; **LF-** lens fibers; **LP-** lens placode; **NR-** neural retina; **OV-** optic vesicle. **EQ-**equator; Arrowheads in point to the high accumulation of immunostaining product at the anterior/posterior lens poles. Scale bars: a-e, a'-e', 100µm; f'-g'. 50 µm.

3.6. GABA receptors and transporters exhibit spatially restricted localization in the newborn mouse lens

The cellular distribution of vesicular and membrane GABA transporters and the two main GABA receptor subtypes was studied by immunohistochemistry using specific antibodies on tissue sections from eyes at different developmental stages. The ionotropic GABA_AR and metabotropic GABA_BR were detected with antibodies specific for GABA_Aβ3 and GABA_BR2, the subunits indispensable for functional assembly of the receptors (see Introduction Chapter). Here we describe the staining patterns at neonatal stage (P0) (**Fig.32**). Staining patterns were quite similar between E16-P0. In neonatal lens labeling with all antibodies was strong in the equatorial secondary fibers and overlaying lens epithelium and extremely weak in the lens nucleus, composed of primary lens fibers. GAT1 spanned a larger expression domain of the lens cortex (**Fig.32b**), in accordance with the elevated expression of GAT1 mRNA at this stage. VGAT and the GABAR showed a strongly segregated expression in the apical/basal membranes of epithelial cells and the closely positioned ciliary body implying the existence of intensive GABA_{A(B)}- mediated signaling between these domains (**Fig.32: a,d,e,e'**).

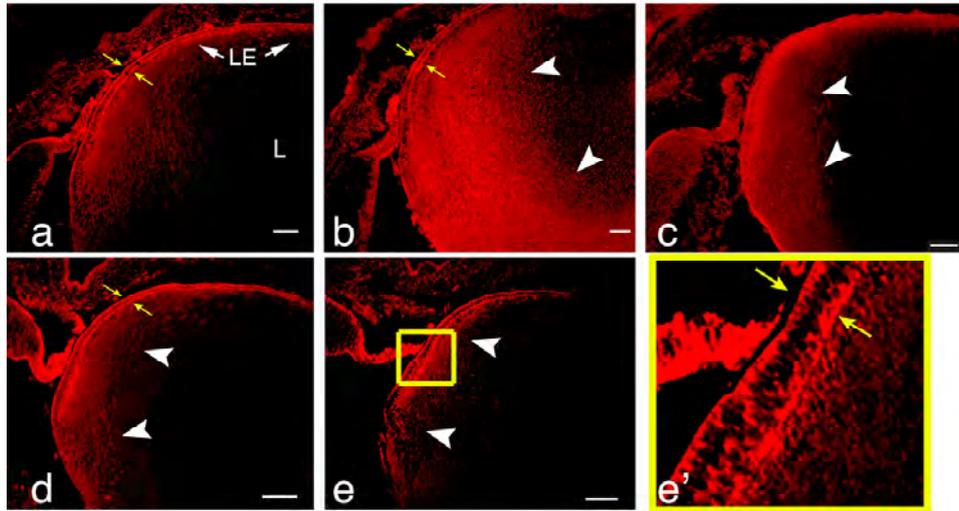


Fig. 32. Cellular localization of the GABA transporters and receptors in the newborn mouse lens: images of coronal cryosections of P0 mouse lens stained with antibodies specific for VGAT (a), GAT1 (b), GAT3 (c), GABA_Aβ3 (d) and GABA_BR2 (e,e'). Immunoreactivity for all antibodies is predominantly localized to the apical/basal membranes of lens epithelial cells (yellow arrows on a-d,e') and to the equatorial fibers. Note that GAT1 staining shows more expanded expression in the cortical region containing secondary fibers compared with VGAT, GAT3, GABA_Aβ3 and GABA_BR2 (arrowheads). Abbreviations: L-lens, LE-lens epithelial cells. Scale bars: 50 μm .

3.7. GABA is produced in the epithelial and fiber cells and induces rise of $[Ca^{2+}]_i$ in equatorial cells of intact newborn lenses

In the newborn (P0) mouse lens GABA is predominantly localized in the lens epithelium and fiber cells with the strongest labeling found at the equatorial region and the forming sutures (**Fig.33I: a,c,e**), a pattern entirely consistent with the expression of the two GAD isoforms and the GABA signaling components (see Figs. 31,32). An inward gradient of GABA staining in the lens cortex and nucleus follows the progressive fiber maturation and denucleation: the signal declined towards the border of secondary/primary fibers as secondary fibers loose their nuclei, but became strong again in external, still nucleated primary fibers (**Fig.33I: a-c**). The nuclei followed a concomitant change in shape and chromatin compaction followed by nuclear membrane disintegration (**Fig.33I: e-e'**). Our results show that GAD and GABA synthesis is turned off shortly before primary/secondary fiber denucleation. To test the functionality of the GABA signaling in intact lenses we performed Ca^{2+} imaging using laser-scanning microscopy and a specially designed perfusion system

(Fig.33II: A-D). In the intact lens only epithelial and equatorial fiber cells were loaded with fluo-4, since they have sufficient esterase activity (Bassnett et al., 1994). Applied GABA evoked an increase in fluorescence intensity mostly in cells of the equatorial region that are undergoing transition from epithelial to fiber cells (Fig.33II: A-C). Usually, several equatorial cells reacted simultaneously with a moderate increase in fluorescence amplitude. Application of KCl (at 20 mM) after GABA washout always induced fluorescence increase, even when the response to GABA was low (data not shown) demonstrating the viability of the responding cells. However, a drawback of this method was that only a small fraction of GABA positive cells in the P0 lens could be labeled with Ca^{2+} sensing dyes and borders of the reacting cells were poorly discernible.

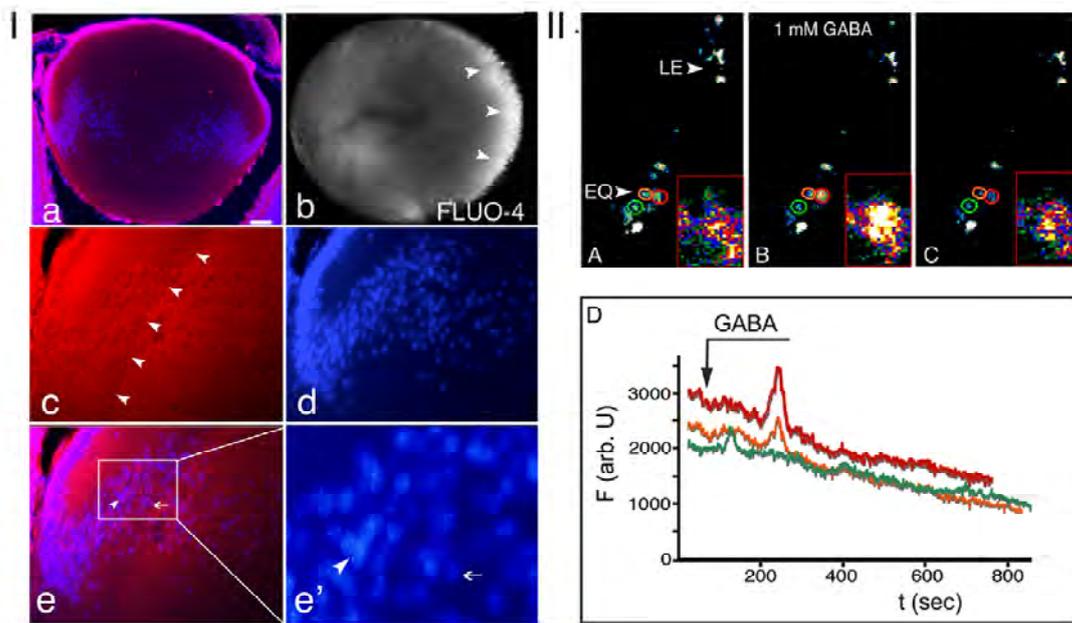


Fig. 33. GABA expression and function in modulation of $[Ca^{2+}]_i$ in the neonatal lens. I: Staining with GABA antibody of the neonatal mouse lens (a, c, e). Nuclei were visualized by DAPI staining (a, d, e, e'). GABA immunoreactivity is visible predominantly in equatorial secondary fibers possessing intact nuclei (arrowheads in e, e') and in primary fibers with nuclei in the process of disintegration (arrows in e, e'). The border between primary and secondary fibers is well demarcated by strong GABA staining as is the equatorial lens region containing the newly generated secondary fibers (arrowheads in c). A P0 lens loaded with Fluo-4/AM (b). **II:** Images from a representative recording of the fluorescence emission in three cells (shown in D in three different colors) of the equatorial region of a Fluo-4/AM loaded newborn mouse lens in buffer alone (A), at the addition of 1 mM GABA (B) and after GABA washout (C). The cells responding to GABA are circled in A, B and C in colors matching the respective recordings shown in D. arb.U-arbitrary units; EQ-equator; F-fluorescence; LE-lens epithelium; t-time. Scale bars: 100 μ m in a-b; 50 μ m in c-e.

To overcome these difficulties, we performed Ca^{2+} imaging in primary LEC culture, an *in vitro* system successfully used before for similar studies (Churchill et al., 2001). To this end first we established and characterized the primary LEC culture system, then we tested whether the LEC cells express *in vitro* the molecular components of GABA signaling.

3.8. Characterization of LEC cultures

Primary LEC cultures were derived from newborn mouse lenses according to a modified protocol and were used for further studies (Fig.28). P0 was chosen as a stage with high expression levels of GABA signaling components as well as ample supply of easy-to-dissect lens epithelial and young fiber cells. As shown in **Fig.34**, after two weeks primary LEC cultures were comprised of both undifferentiated epithelial cells that can be spread or form epithelial monolayers (**Fig.34: e,f**) and randomly distributed lentoids- multilayered cell aggregates that appear to bulge from the surrounding epithelial cells (**Fig.34: d,e**). On the basis of morphological criteria and expression profiling they are considered an equivalent of elongating newly-formed fiber cells (Menko et al., 1984; TenBroek et al., 1994; Frenzel and Johnson, 1996; Wagner and Takemoto., 2001).

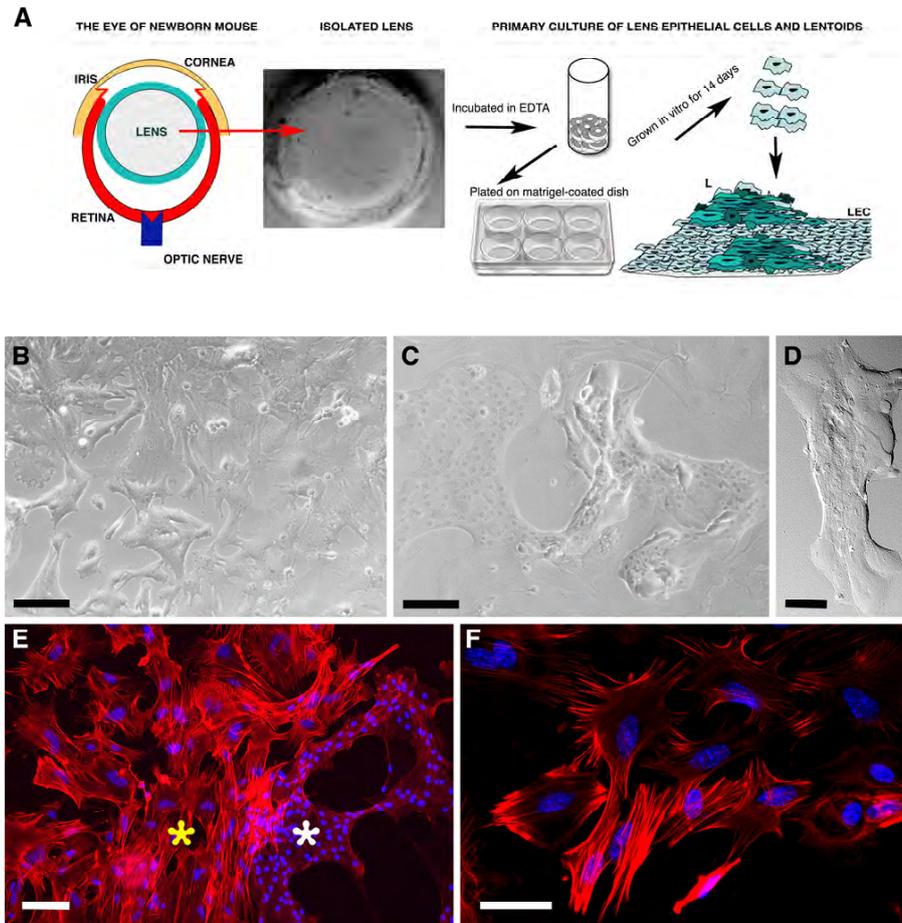


Fig.34. LEC culture established from newborn mouse eye. **A:** Scheme for lens cell isolation procedure and establishing of LEC culture. Following removal of retina and vitreous body from eyes of newborn (P0) mice, lenses were incubated in EDTA solution, centrifuged and plated in M199 medium on matrigel-coated coverslips. **B-F.** Images of LEC after seven (B) or fourteen (C-F) days in vitro by which time culture comprised of epithelial monolayers (E,F) and spontaneously differentiated fiber cells in lentoids (C,D,E). Cells were visualized by phase contrast (B, C), DIC (D) or by immunofluorescence phalloidin staining for filamentous actin (E, F). Stars in E indicate the area of epithelial cell monolayer (yellow) and lentoids (white). The cell nuclei were counterstained with DAPI (blue). Scale bars: 100µm (B, C, E), 50µm (D, F).

First, we examined the cultures for the expression of cellular markers and lens fiber specific proteins. The molecular chaperone α A-crystallin, one of the most abundant proteins of the lens epithelium and fibers (Piatigorsky et al., 1989) and the small α B-crystallin, more abundant in fiber cells were both detected in lentoids and epithelial cells (**Fig.35: A,B,D**). Lentoids, but not single epithelial cells stained strongly for N-cadherin (**Fig.35C**), while E-cadherin only stained the membranes of epithelial monolayers, but not lentoids (**Fig.35: E,G**).

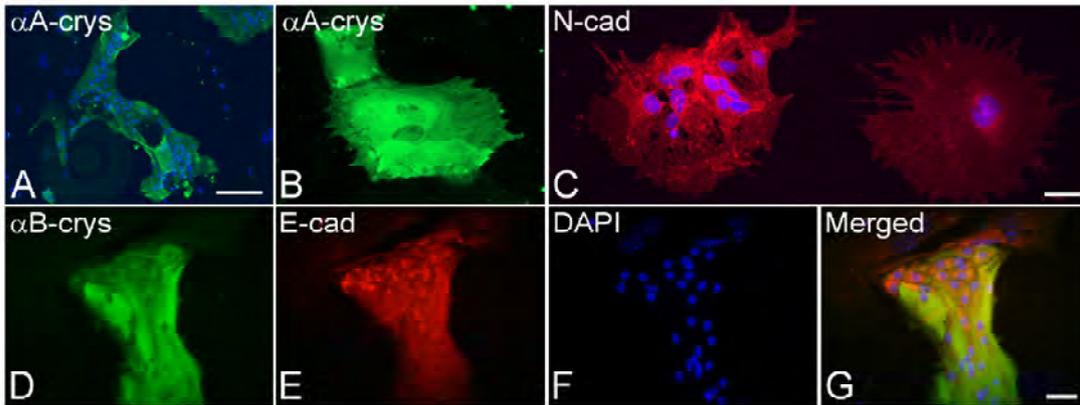


Fig.35. Characterization of mouse lens epithelial primary cultures Immunofluorescent staining of primary LECs with antibodies to lens specific α A- and α B-crystallin, E- and N-cadherin. *A, B, D*. As the cells began to differentiate and form lentoids and as the lentoids began to become more highly differentiated, express not only α A- but also α B- crystallin. *C-G*. The polyclonal N-cadherin-specific antibody stains specifically the long (lateral) sides of fibers in lentoids, while epithelial cells and monolayers are not stained. E-cadherin stained only the membranes of epithelial monolayers, but not lentoids. Scale bars: 100 μ m in *A*, 50 μ m in *B-C*, 20 μ m in *D-G*.

3.9. Lens epithelial cells and lentoids grown *in vitro* express key GABA signaling components and contain GABA

Similar to intact lenses, cells in primary lens epithelial cultures prepared from P0 lenses expressed GAD65 and GAD67 forms that co-localized with the vesicular and membrane GABA transporters, respectively (**Fig.36**). In general, lentoids showed the strongest expression, while epithelial monolayers were less prominently stained. Some GAT1⁺ cells in LEC cultures did not express GAD (**Fig.36: E,G,H**), which suggests that GAT1 in these cells operates as GABA uptake only, similar to CNS glia involved in the GABA clearing from the synaptic cleft.

Moreover, strong labeling of GABA_A β 3 and GABA_BR2 was detected over the cell borders of the lens epithelial monolayers and lentoids (**Fig.37**). The staining had a punctate appearance and was greatly enriched near cellular membranes (**Fig.37: A' B'**). Single cells showed little or no staining (data not shown). In conclusion, epithelial sheets and lentoids expressed all GABA signaling components required for building up a functional pathway.

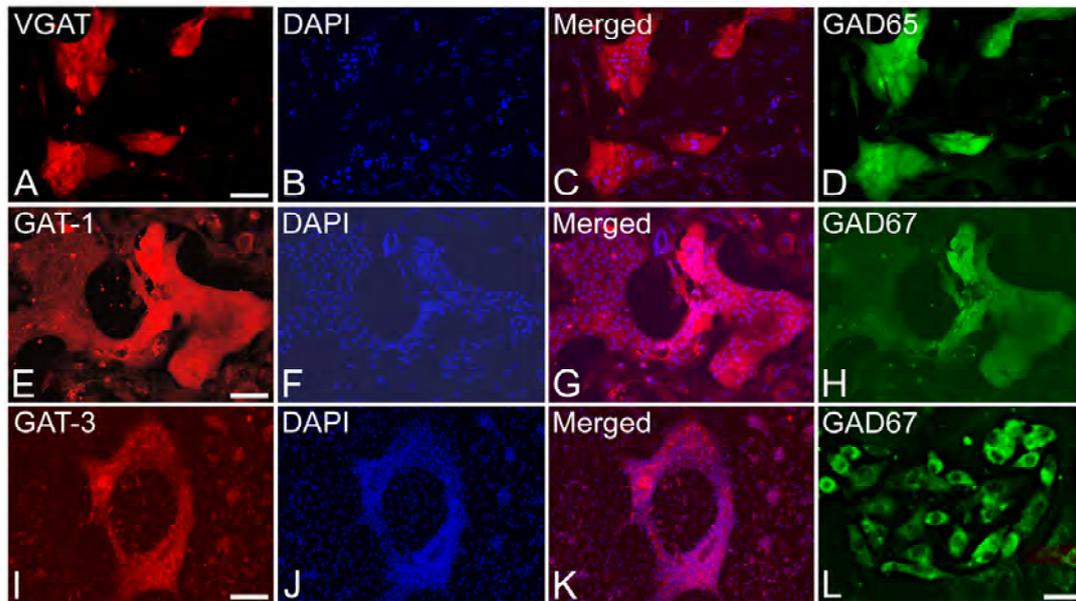


Fig.36. Expression of GAD and GABA transporters in primary cultures of mouse lens epithelial cells (LEC). A-D. VGAT entirely co-localizes with GAD65; staining of the same preparation with the polyclonal anti-VGAT serum and monoclonal anti-GAD65 antibody GAD-6. E-H. Co-localization of GAT1 and GAD67 The membrane localization of GAT1 is more evident in the less prominently stained epithelial monolayer than the strongly positive lentoid I-K. Staining with anti-GAT3 antibody. L. GAD67 is also found in small cells outside lentoids and monolayers that are positive for α A-crystallin (not shown). Scale bar: 100 μ m in A-K, 50 μ m in L.

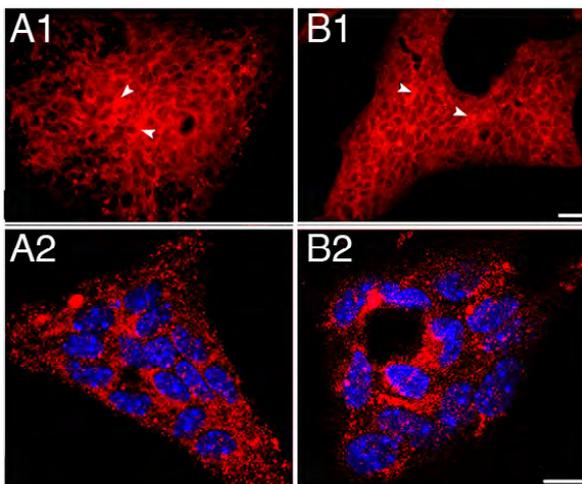


Fig. 37. Expression GABA transporters receptors and GABA in primary cultures of mouse lens epithelial cells (LEC). GABA_A and GABA_B receptors are enriched in multicellular lentoids. LEC cultures were stained with rabbit anti-GABA_A β 3 (A) or anti-GABA_BR2 (B) antibodies and visualized by epifluorescence (A1,B1) or laser scanning microscopy (A2,B2). Nuclei were revealed by DAPI staining (A2,B2). The staining is greatly enriched near cell borders and at the surface of lentoids (arrowheads in A, B). Scale bars, 20 μ m in A, B, 10 μ m in A1,B1.

Finally, GABA synthesis and accumulation in primary LEC was verified by staining with a specific antibody (**Fig.38**). Lentoid structures showed much higher accumulation of GABA compared to adjacent LEC monolayers (**Fig.38: a, a'**), while single cells remained unstained (**Fig.38: c, c'**). This expression gradient was typically found in the vicinity of all lentoids and was accompanied by a gradual change in

nuclear shape and chromatin condensation reminiscent of the inwardly directed progressive nuclear loss in maturing fibers of intact lenses (**Fig.38: b, b'**; see also Fig. 33). Thus, primary LEC cultures provide a useful and matching model to study lens fiber differentiation.

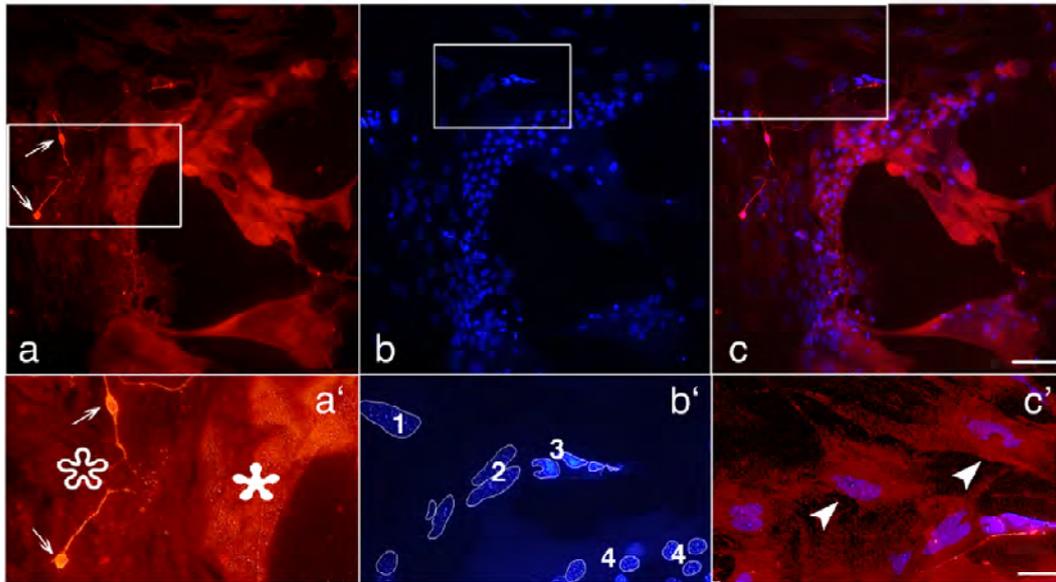


Fig.38. GABA is produced in LEC cultures. Primary LEC cultures derived from P0 mouse lenses were stained with a rabbit antibody against GABA (**a, a'**; **c, c'**). Nuclei were stained with DAPI (**b, b'**). Boxed regions in **a, b,** and **c** are enlarged in **a', b'** and **c'**, respectively. A large multicellular lentoid-like structure (**a-c**) is highly positive for GABA in bulging parts (solid asterisk in **a'**) and less pronouncedly stained at the flatter, presumably monolayer regions (empty asterisk in **a'**), while cells outside the multicellular structure were unstained (boxed area in **c** devoid of staining was enhanced in **c'** to reveal the presence of cells in this area). Rare GABA⁺ neurons can be seen (arrow in **a**) in the preparations that display similar intensity of staining as lentoids (solid asterisk in **a'**). **b'**-circled nuclei of different shapes and level of chromatin compaction (numbered 1-4) correlating with stages of fiber cell differentiation proceeding from less differentiated (nuclei #1, 2) to more differentiated (nuclei#3 and 4) stages (91). Scale bars: 50 μ m in **a-c,** 20 μ m in **a'-c'**.

3.10. GABAR-mediated modulation of $[Ca^{2+}]_i$ levels in primary LEC and lentoids

The functional assembly and activity of GABA_A and GABA_B receptors expressed by primary LEC was verified by measuring $[Ca^{2+}]_i$ changes in cultures upon bath application of GABA and selective agonists/antagonists. Most of the medium size lentoids and single cells were successfully loaded, while some larger lentoids, likely to represent a later stage of lens fiber cell differentiation remained unlabeled (data not shown).

GABA (1mM) and GABA_AR agonist muscimol (100 μ M) induced a simultaneous increase of fluorescence intensity in epithelial cells or cell groups belonging to

smaller lentoids (not shown). Cell-to-cell propagation of the signal, sometimes reaching out to adjacent lentoids was clearly observed and it was rarely followed by oscillations. The muscimol-evoked $[Ca^{2+}]_i$ rise was almost reversibly blocked in the presence of the GABA_AR antagonist bicuculline (0.5 ± 0.34 , $n=6$; amplitude $19.52 \pm 5.84\%$) demonstrating the specific involvement of the receptor in the response (**Fig.39**). Lower number of cells responded to GABA (2.45 ± 0.42 , $n=20$) than to muscimol (4.2 ± 1.01 , $n=10$), but the amplitude of their responses was almost identical ($48.06 \pm 9.26\%$ and $46.5 \pm 5.49\%$, respectively; **Fig.39: C1,C2**). However, this difference may be incorrectly estimated due to difficulties in defining the contours (and ultimately the exact numbers) of reacting cells within the lentoids.

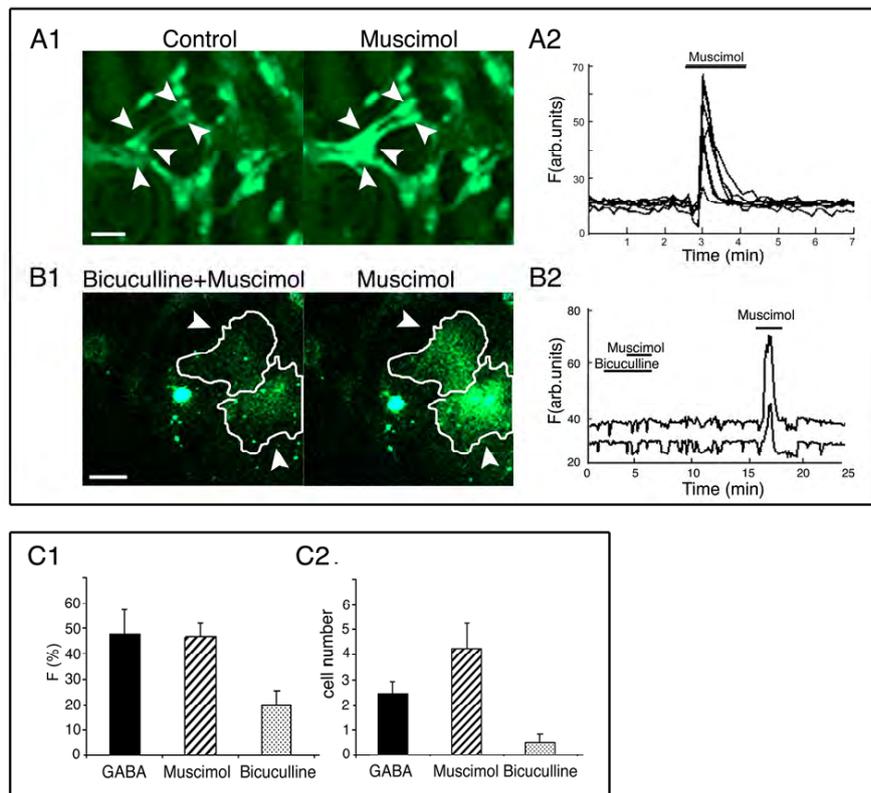


Fig.39. GABA_A receptor activation induces rise in $[Ca^{2+}]_i$ in lens epithelial cultures. *A1, A2, B1, B2.* Representative recordings of GABA_A receptor-evoked changes in fluorescence of Fluo4/AM-loaded primary, lens cells (arrows in A1, B1). Cultures were initially perfused with AHH buffer to obtain a steady baseline (Control) followed by 100 μM muscimol (Muscimol; A1, A2). Alternatively, following 20 μM bicuculline, 100 μM muscimol and 20 μM bicuculline were co-administered (Bicuculline+Muscimol) and then 100 μM muscimol (Muscimol) alone (B1, B2). A1, B1 confocal images were taken at different time points of the recording shown in panels A2, B2. Note that the muscimol-evoked fluorescence rise was completely blocked by bicuculline (B1, B2). *C1, C2.* Bar charts illustrating the average number of cells (C1) and the amplitude change (C2) responding to GABA, muscimol alone and muscimol in the presence of bicuculline. F% represents the change in fluorescence intensity relative to the basal fluorescence. Data represent the mean ± S.D. ($p < 0.05$). Scale bar: 50 μm.

GABA_BR agonist baclofen (20 μ M) induced calcium transients in LEC, which differ in amplitude and kinetics between different cells and sometimes within a single cell after repeated drug applications. The baclofen-evoked transients in single cells had the smallest amplitude, which can be explained by the low number of receptors, their low sensitivity to baclofen or the presence of baclofen-evoked hyperpolarizing responses (not shown). Application of baclofen sometimes induced a secondary wave of signal propagation spreading further into the lentoid cells and adjacent epithelial monolayer parts and followed by persistent oscillations, which sustained even after complete washout of drugs (**Fig.40: C1,C2**). The response to baclofen was typically completely blocked in the presence of the GABA_BR antagonist CGP55845. These observations were highly reminiscent of responses involving mobilization of intracellular Ca²⁺ in lens cultures mediated by ATP-receptors and strongly suggests that it may be propagated through intercellular gap junctions and/or purinergic receptors (Duncan et al., 1993; Churchill et al., 1996; Williams et al., 2001). In addition, initial application of CGP sometimes induced small-amplitude and short-term oscillations in single cells (not shown) which we could not explain at the moment. Furthermore, preliminary data indicates that baclofen-induced calcium transients are reduced, but not abolished, in the absence of extracellular calcium, suggesting that they might partially depend on influx of calcium from the extracellular space.

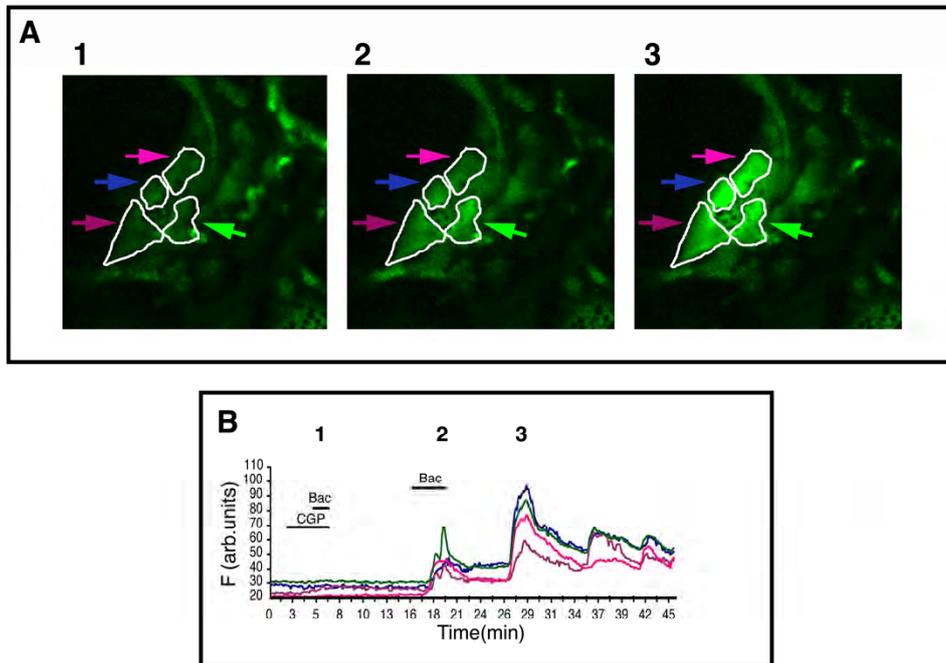


Fig.40. GABA_B receptor activation induces rise in [Ca²⁺]_i in lens epithelial cultures. Cells were initially washed for 2 min with AHH buffer containing 20 μM CGP55845 (CGP) followed by 20 μM CGP+20 μM baclofen (CGP+Baclofen) and 2 min of baclofen alone (Baclofen). In some cells, mostly belonging to small lentoids, single application of baclofen induced repetitive and prolonged rises in intracellular calcium concentration.. **A.** Images (1,2,3) were taken at time point of the recordings shown in panel **B** (1,2,3, respectively). Scale bar, 50 μm.

VII DISCUSSION

1. MOUSE ES CELLS EXPRESS FUNCTIONAL GABA SIGNALING

1.1. Structural components of GABA signaling are expressed in the mouse ES cells of line R1

The expression of molecular components of GABA signaling in the undifferentiated ES cell was examined by RT-PCR and immunocytochemistry. We showed for the first time that the mouse ES cells, developmentally closely related to the ICM of the mouse blastocyst express all structural components needed for assembly of functional GABA signaling: GABA, GABA synthesizing enzymes (GAD65 and EGADs), membrane and vesicular GABA transporters (GAT1 and VGAT), as well as multiple GABA_A ($\alpha 1, \alpha 3, \alpha 4, \alpha 5, \beta 3, \gamma 2S, \gamma 3, \delta, \epsilon$) and GABA_B (R1, R2) receptor subunits. Furthermore, the enrichment of GABA_A $\beta 3$ and GABA_BR2 at sites of putative contacts between adjacent cellular membranes, revealed by immunofluorescent labeling, suggests that both types of GABA receptor are engaged in intensive cell-cell communications within the ES colony.

Remarkably, 11 out of 19 known to date GABA_AR subunits belonging to all major subunit classes are moderately or abundantly expressed in undifferentiated mES cells providing the structural requirements for the formation of functional GABA_A receptors. As previously shown, despite the high number of subunits for each subclass, there is only limited number of pentameric combinations actually exist in the CNS (Barnard et al., et al., 1998; Costa, E., 1998; Mohler, 2006). Interestingly, we detected all but two α ($\alpha 2$ and $\alpha 6$), all γ , in addition to δ and ϵ subunits, but only a single type of the GABA-binding β subunit ($\beta 3$) in undifferentiated ES cells. The $\beta 3$ gene is part of the $\alpha 5$ - $\beta 3$ - $\gamma 3$ gene cluster on mouse Chr. 7 (human CHR 15) implicated in Angelman's syndrome (Hommanics et al., 1997; DeLorey and Olsen,

1999) the only one that is fully turned on in ES cells, the remaining four clusters (Barnard et al., 1998) are only partially transcribed. This expression pattern may be important consequences for the GABA_A receptor expression regulation and receptor properties at later developmental stages. First, it may be involved in defining a future expression paradigm for the co-ordinate regulation of the genes within a cluster, second it should allow for only a limited number of pentamers to be formed in the pluripotent ES cells. An obvious consequence of the absence of $\beta 2$ subunit would be the lack of the major brain synaptic GABA_A receptor type composed of $2\alpha 1-1\beta 2-2\gamma 2$ (Mohler, 2006; Mody, 2008) and the preferential assembly of predominantly extrasynaptic receptors due to abundant expression of subunits found mostly ($\alpha 5$, Collinson et al., 2002) or exclusively ($\alpha 4$, δ and ϵ ; Mohler, 2006; Mody, 2008) extrasynaptically. $\alpha 1$, which is mostly synaptic in adult brain can also localize to extrasynaptic sites when assembled with δ (Glykys et al., 2007). Notably, of the two $\gamma 2$ splice variants only $\gamma 2S$ (short) is present, while the $\gamma 2L$ (long) form, which incorporates preferentially at postsynaptic sites (Meier and Grantyn, 2004) is altogether missing. The $\gamma 2S$ can also physically interact with GABA_BR1 and anchor it to the membrane (Balasubramanian et al., 2004).

However, a synaptic type GABA_A receptor can still be assembled between $\alpha 1-3$, $\beta 3$ and $\gamma 2S/\gamma 3$. Interestingly, $\gamma 3$ is very restricted in adult brain (Sieghart and Sperk, 2002), but is the most abundant of all γ subunits in mouse ES cells. This observation holds true also for the developing lens (see next Chapter), and may indicate common features in the developmental programs of these systems.

In brain synaptic GABA_AR mediate fast (phasic) neurotransmission, while extrasynaptic GABA_AR are responsible for the slow and long-lasting “tonic” inhibition (reviewed in Mody, 2008). Extrasynaptic receptors containing δ and ϵ subunits are potentiated by allosteric modulators like alcohol, anaesthetics or

neurosteroids (Mohler, 2006; Mody, 2008;). Furthermore, GABA_A receptor containing the ϵ subunit, both splice variants of which we found highly expressed in ES cells, can replace not only γ , but also α or β subunits and assemble into variable and often constitutively active receptors (Neelands et al., 1999. Bollan et al., 2008) that are readily potentiated by steroids in the absence of GABA (Bollan et al., 2008).

The metabotropic GABA_B receptor was represented in ES cells by the R1a and R1b variants of GABA_BR1 and the GABA_BR2 subunit and found at the protein level in a specific vesicular pattern previously found in neural (Ritter et al., 2005) and peripheral Schwann cells (Magnaghi et al., 2004).

The enrichment of GABA receptors and GABA transporters at sites of putative contacts between adjacent cellular membranes suggests that they are engaged in intensive cell-cell communications within the ES colony.

1.2. GABA receptor of unique features act synergistically to modulate $[Ca^{2+}]_i$ in ES cells

Herein we show for the first time that in ES cells multiple GABA_AR and/or GABA_BR subunits assemble into functional receptors, and their activation with specific ligands triggers Ca^{2+} transients. The GABA_AR-mediated $[Ca^{2+}]_i$ rise in response to muscimol is clearly of dual nature due to Ca^{2+} influx through Cd^{+} -sensitive VGCC and mobilization of CPA-sensitive internal stores. Accordingly, the cellular machinery required for these responses is present in the ES and displays several specific features. First, the low level of L-type (long-lasting, high voltage-activated) channel and moderate to high expression of the N (neural), R (residual) and especially T (transient), moderate-to-low voltage activated type Ca^{2+} channels (Han and Lee, 2005; Lee et al., 2006) suggests that Ca^{2+} rise may occur at low excitation levels, but long-lasting and possibly detrimental $[Ca^{2+}]_i$ rises are mostly avoided. Second, the

abundant expression of two out of three IP₃R (Yanagida et al., 2004,) and detectable RyR1 and RyR2 receptors suggests a greater role of the intracellular Ca²⁺ stores (see below).

The [Ca²⁺]_i increase induced by non-synaptic GABA_A receptor activation in the embryonic nervous system (see Introduction) or non-neuronal cells (Kuroda et al., 1999; Xie et al., 2004) is well documented and ES cells are of no exception from the classical examples. Quite unexpectedly, however GABA_B receptor activation by the highly specific agonist baclofen also induced Ca²⁺ transients, which were mostly, if not exclusively due to Ca²⁺ release from the intracellular Ca²⁺ stores (Verkhatsky, 2005) as they were insensitive to externally added Ca²⁺ or the presence of VGCC inhibitors, but could be blocked almost entirely by the ER-SR Ca-ATPase blocker CPA. This response is the opposite of the well-documented inhibitory effect of GABA_B receptor on [Ca²⁺]_i rise in both mature and immature neurons and non-neuronal cells (Watanabe et al., 2002; Nicoll, 2004) and is very similar to the recently described baclofen-evoked and entirely intracellular store-dependent [Ca²⁺]_i rise in primary astroglia cultures (Meier et al., 2008).

Previous data showed that mouse ES cells or ICM express exclusively IP₃R and there is a developmental switch from IP₃R to RyR-mediated Ca²⁺ signalling (Rosemblyt et al., 1999; Yanagida et al., 2004.). However, we have detected low-to-moderate expression of RyR and both RyR₁ and RyR₂ have been found in human ES cells (Goh et al., 2005).

Interestingly the GABA_B receptor antagonist CGP35348 in the same experimental paradigm was similarly incapable of suppressing the Ca²⁺ rise following KCl-baclofen application pointing again at the specific features of the receptor. Thus, further experiments will be needed to determine the exact sequence of downstream

events and components involving intracellular Ca^2 stores following GABA_B receptor activation in mouse ES cells.

GABA_B receptors can also facilitate Ca^{2+} influx via store-operated channels (SOC, New et al., 2006) or potentiate the L-type VGCC (Carter and Mynlieff, 2004; New et al., 2006) in an external Ca^2 - dependent way, although these comprise a smaller component in the overall response.

Another notable feature of the GABA_B R-mediated Ca^2 signalling in ES cells is the simultaneous reaction of neighbouring cells often followed by persistent oscillations that may share similarities with IP_3 R-mediated response in mesenchymal stem cells (Sun et al., 2007 and ref therein).

Intercellular Ca^2 waves become important at the blastocyst stage with the increase of cell number and demand for co-ordinate cellular activity (Webb and Miller, 2003). The Ca^2 transients observed in small clones or groups of neighbouring cells on the surface of bigger clones upon activation of the GABA_B R result clearly from a coordinate response most probably involving intercellular spread of Ca^{2+} and/or IP_3 through gap junctions formed by Cx43 abundantly expressed in both ES cells and ICM (Verkhatsky et al., 1998; Webb and Miller, 2003).

Surprisingly, single transients could be blocked by CGP55485 but repeated long-lasting transients were insensitive to two different GABA_B R blockers (FENS, Schwirtlich et al., 2006) and may result from a constitutive activation of GABA_B receptors.

It should be stressed that CGP alone could induce a moderate rise in the basal Ca^2 level indicating the existence of a classical inhibitory component of the GABA_B R response involving VGCC inhibition and/or GIRK channel activation. This Ca^2 rise however is also potentiated by baclofen in ES cells. Hence, our results suggest that ES

cells express pharmacologically different GABA_BR types that exert distinct functions through modulation of the $[Ca^{2+}]_i$ level.

GABAR as modulators of cell proliferation in ES cells

GABA has long been known as a modulator of cellular proliferation of neuronal progenitors (Barker et al., 1998; Haydar et al., 2000; Varju et al., 2001; Owens and Kriegstein, 2002b; Ben-Ari et al., 2007). Pioneer experiments showed that either GABA or GABA_AR agonists inhibit cell proliferation in neocortical ventricular zone (VZ) cells (LoTurco et al., 1995). Similar results were obtained for adult neural stem cells (Liu et al., 2005) or cancer cells (Watanabe et al., 2006), while it stimulates proliferation of VZ cortical (Haydar et al., 2000) or cerebellar (Fiszman et al., 1999) progenitors, growth plate chondrocytes (Tamayama et al., 2005), Schwann cell (Magnaghi et al., 2004), oligodendrocytes (Luyt et al., 2007) and cultured embryonic neural progenitor cells (Fukui et al., 2008).

Our results show, that co-activation of both GABA receptors in mES cells has little effect on the proliferation due to the opposite action of the two receptor types. Notably, the persistent activation of GABA_AR has a differentiate effect based on the alkaline phosphatase staining experiments and induces expression of neuronal-specific markers. Thus, the action of GABA and different GABA ligands on proliferation and cell cycle are modulatory and not regulatory. Therefore, our results support the data obtained from GABAR knock-out experiments, which do not exert an overt phenotype in the preimplantation embryo (Burt, 2003)

1.4. GABA SIGNALLING DURING IN VITRO NEURONAL DIFFERENTIATION OF MOUSE ES CELL RECAPITULATION OF EMBRYONIC DEVELOPMENT

We found that *in vitro* model of neuronal differentiation of mouse ES cells (4-/4+) recapitulated many aspects of the normal development program that directs *in vivo* GABAergic differentiation. Our RT-PCR results revealed that all molecular components necessary for the assembly of functional GABA signalling follow a specific pattern of temporal regulation during different stages of neuronal differentiation. The embryonic and adult GAD forms were sequentially induced in the order GAD25, GAD44, GAD67 and GAD65 following the overlapping and transient expression of the upstream regulators, *Dlx2* and *Dlx5*, respectively.

We detected 11 GABA_A receptor subunits in our cultures: the ones whose genes are located on chromosome 4 ($\alpha 2/4$, $\beta 1$ and $\gamma 1$) and 15 ($\alpha 5$, $\beta 3$ and $\gamma 3$) were induced and upregulated during early stages (*dai2*, *dai4*) of neuronal differentiation and those located on chromosome 5 ($\alpha 1$, $\beta 2$ and $\gamma 2L$) were induced at a later differentiation period which is in agreement with data obtained in CNS. GABA_B receptor subunits R1 and R2 were highly expressed in the first days after RA induction. Finally, the expression of both membrane (mostly GAT1) and vesicular GABA transporters (VGAT) appeared at early stages and increased in parallel with neuronal maturation, thus both vesicular and GAT-mediated GABA release may occur in the absence of synaptic specializations. As revealed by GABA immunocytochemistry, the earliest stages of differentiation were characterized by initial strong expression of GABA over all cells of the embryoid bodies (EBs), followed by gradual appearance first in bipolar cells, followed by a massive outmigration of GABAergic progenitors and sprouting of neuritis concomitant with the disappearance of staining from the EB remnant. These highly reproducible stages match the expression of a selective

repertoire of genes and may indicate the existence of a self-autonomous GABAergic differentiation programme.

2. DYNAMIC REGULATION OF MOLECULAR COMPONENTS AND FUNCTIONALITY OF THE GABA SIGNALING DURING MOUSE LENS DEVELOPMENT

The vertebrate lens develops from a series of interactions between the surface ectoderm, the optic vesicle, and the surrounding mesoderm, involving successive steps of bias, competence, specification, and differentiation (McAvoy, 1999; Hirsch and Grainger, 2000). In addition to that, lens is attractive as a model system due to its regional compartmentalization. Proliferation occurs in the anterior lens epithelium, migration in the equatorial zone, and differentiation proceeds with cell elongation at the anterior/posterior poles.

Due to the above reasons, lens is a frequently used model to study the role of different signaling molecules in basic developmental processes.

Expression of different GAD forms recapitulates different phases of lens development

We found expression of all four known GAD forms during lens development : GAD65 and the three highly related proteins encoded by the GAD67 gene (mostly embryonic - GAD25, GAD44 and “adult” GAD67). During earlier embryonic stages (E14.5-E16.5) predominantly GAD65 and the embryonic forms GAD25 and GAD44 were expressed, adult GAD67 was only detected at a low level by RT-PCR, but not by Western blot, indicating that it may be present at extremely low levels. Our

immunohistochemistry results also showed that during E9.5-E16.5, the GAD65 and GAD67-related (mostly attributable to EGAD) immunolabeling was essentially overlapping, localized in both lens epithelium and lens fiber cells.

The highly coordinated expression of GAD65 and EGADs implies that these two forms have a synergistic role in the primary fiber differentiation (involving the processes of cell migration, elongation, and withdrawal from the cell cycle).

In accordance with their expression pattern, GAD44 along with GAD65 may provide GABA for the early stages of epithelial cell proliferation and primary fiber cell differentiation (**Fig.40**). Due to preferential accumulation of the “early” GADs at the apical and basal tips of the primary fibers, GABA synthesized by these forms may play a role in remodeling of the membrane skeleton during fiber elongation. Furthermore, the “early” GAD related GABA signaling may be involved in formation of transient cell-adhesion complexes between elongating fiber cells and overlying lens epithelium anteriorly and the lens capsule at the posterior lens wall. Similarly, the extremely high levels of “early” GADs at the anterior and posterior lens fiber tips and in the central lens epithelium during E13.5-E16.5 suggests that GABA synthesized by GAD44/GAD65 is involved in cell-cell adhesion mechanisms between fiber tips of opposite sides during formation of lens sutures (Fig.4, Kwakowski et al., 2007; Beebe et al., 2001). The highest level of expression of both EGADs and GAD65 was observed in the primary lens fibers at stages 16.5-E17.5 characterized by extensive fiber elongation and beginning of nuclear disintegration, indicating a role in this process as well. GAD65/EGAD segregation of function may be linked to different mechanisms of GABA release, VGAT or GAT, respectively (**Fig.40**; Szabo et al., 1994).

The function of GAD25, the enzymatically inactive truncated form of glutamate decarboxylase, is more intriguing. Similar to the nervous system, GAD25 is

transiently expressed during early development, but it is also found in adult brain (Szabo et al., 1994, 2000).

From E17.5 on, GAD65, EGAD and GAD67 expression patterns change in accordance with the induction of primary fiber terminal differentiation and more enhanced secondary fiber generation, which continues into adult (**Fig.40**). GAD65 and EGADs gradually disappear from the lens core, concomitant with up-regulation of GAD67 at the equatorial region containing elongating secondary fibers. Therefore, it can be speculated that GAD67 is involved mostly in the differentiation of secondary fibers after birth.

Thus, there is a clear spatio-temporal segregation of “early” (GAD65 and EGAD) and “late” (GAD67) GAD forms. The predominant expression of GAD67 in the elongating nucleated fibers of the postnatal lens, where the lens maturation occurs much faster, in a narrow, 1-2 cell-wide region may be explained by the need for a quick recruitment of active GAD, which is provided by GAD67 existing in a PLP-bound holo-GAD, but not by GAD65, which is found mostly as an inactive apo-GAD (Martin et al., 2000; Varju et al., 2001) that is regulated by the PLP/ATP level in the cells.

Finally, the previously reported expression of GAD67 and embryonic forms in the developing rat lens (Li et al., 1995) shows remarkable similarities to that in the mouse with some notable differences: in E17.5 rat lens (corresponding roughly to E16.5 in the mouse), when the highest level of GAD is reached, mostly GAD67, but not EGAD is synthesized, furthermore, GAD67 was not detected before E15 and shortly after birth. Among these, the absence of GAD67 staining from the postnatal cortical fibers may be due to a much lower level of GAD67 in the rat compared to mouse lens and/or a lower affinity of the GAD67-specific antibody used in this study. This may also help explain the discrepancy between lack of GAD67 immunoreactivity and the

presence of GABA in the postnatal rat lens (Li et al., 1995 and references therein). GABA has been detected at variable levels also in the lenses of other vertebrates (discussed in Li et al., 1995).

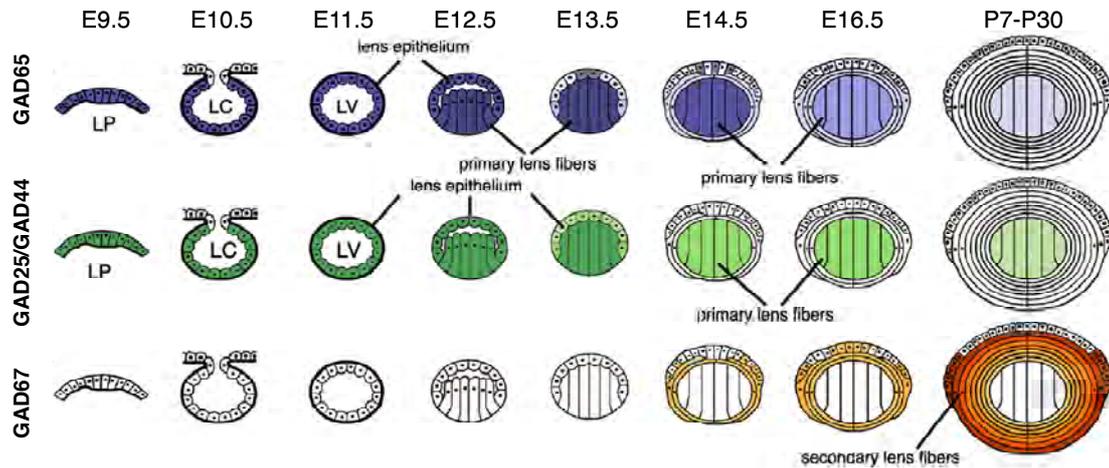


Fig. 40. Schematic presentation of GAD65 and GAD67 expression during lens development. Starting from the lens placode stage to E16.5, GAD65 (Blue) and GAD67-related (GAD25 and/or GAD44; Green) expression entirely overlaps. GAD65 expression overlaps with EGAD in lens epithelium and the primary fiber cells, where it is down-regulated after P14. At P30 and thereafter, only GAD67 (Orange) is expressed in the elongating secondary fiber cells and some cells of the epithelium, but not in lens nucleus.

2.2. Transcriptional regulation of GAD in the lens - involvement of *Dlx2* and *Dlx5*

The earliest GAD (GAD65, EGAD) and GABA expression was observed in lens placode where multiple transcription factors like Pax6, Six3, Eya, Dach1, Mafk, Sox2 and 3 and Prox1 acting as key regulators in lens induction and fiber cell differentiation are present (see Introduction chapter, Fig. Kawauchi et al., 1999; Kim et al., 1999; Ogino and Yasuda, 2000; Ring et al., 2000; Purcell et al., 2005). Among these, Pax6 and Sox have binding sites on conserved regions of GAD65 and GAD67 promoters and are potential cis-regulators of GAD. Pax6, Six3 and Dach1 expression patterns (Purcell et al., 2005) largely overlap with GAD at early developmental stages not only in lens, but also in retina and ciliary marginal zone. Whether any of these

transcriptional factors directly regulate the transcription of GAD genes in the lens is a matter of speculation at present.

The Distal-less (Dlx) homeobox containing transcription factors also share overlapping domains of expression with GADs in the midgestation embryo (Katarova et al., 2000; Maddox and Condie, 2001), although it is not clear at present whether they can bind to and directly regulate GAD transcription. In the developing brain, GAD65 and GAD67 expression domains grossly overlap with that of the Dlx1-2 and Dlx5-6 implicated as key regulators of differentiation and migration of a subset of GABAergic inhibitory neurons (Anderson et al., 1997a,b; 1999; 2001; Cobos et al., 2005; Marin and Rubenstein, 2001; Katarova et al., 2000); Panganiban and Rubenstein, 2002; Pleasure et al., 2000). Dlx2 and Dlx5 may also regulate GAD in neurons outside the brain, like the migratory LHRH neurons, or in non-neuronal structures such as teeth, vibrissae, palate, where they show overlapping expression patterns (Porteus et al., 1994; Thomas et al., 1997; Katarova et al., 2000; Maddox and Condie, 2001; Depew et al., 2005; Givens et al., 2005; Tamayama et al., 2005; Levi et al., 2006).

The presence and possible function of Dlx2 and Dlx5 during lens development has not been described previously, although detailed expression studies in the mouse embryo have been published (reviewed in Panganiban and Rubenstein, 2002; Zerucha and Ekker, 2000). We find by semi-quantitative RT-PCR that the two genes are clearly expressed in an overlapping and sequential manner. Dlx2, which prevails at early stages, shows a temporal profile indistinguishable from that of EGAD and very similar to GAD65 may be needed for induction and/or maintenance of the early GAD forms (EGAD and GAD65). Dlx5 expression correlates only partially with that of GAD67: its initial up-regulation precedes that of GAD67 (E17.5 vs P0) and is not expressed beyond P14 in contrast to GAD67, which shows elevated expression until

P30. These findings strongly suggest that *Dlx5* may be needed for induction, but not for maintenance of expression of *GAD67*. Further experiments will be needed to verify this hypothesis.

The sequential and overlapping expression of *Dlx2* and *Dlx5* in the lens is reminiscent to that in the brain (Liu et al., 1997; Eisenstat et al., 1999; Panganiban and Rubenstein, 2002) and may be related to the sequential induction of *EGAD* and *GAD67* in the differentiating GABAergic progenitors (Katarova et al., 2000; Result Chapter 2.1). No lens phenotype has been described for *Dlx1/2*, *Dlx5/6* double knock-out mutants, which exhibit embryonic or neonatal lethality (Zerucha and Ekker, 2000; Panganiban and Rubenstein, 2002; deMelo et al., 2005). This would suggest that *Dlx* genes have a subtle effect on lens morphogenesis during embryonic development, which can be better clarified in conditional knock-out experiments.

2.3. Different molecular components of the GABA signaling pathway exhibit similar expression profiles during lens development- evidence for co-regulation

We presented for the first evidence on the presence and dynamic regulation of all component of GABA signaling machinery involved in GABA synthesis (*GAD*), release (*VGAT*, *GAT*), binding (*GABA_A* and *GABA_B* receptors) and uptake (*GAT*) in the developing mouse lens.

We showed that selected *GABA_A*R subunits ($\alpha_{1,2,3,4}$; $\beta_{1,2,3}$; $\gamma_{1,2,3}$ and δ), both *GABA_B*R subunits, the vesicular and membrane GABA transporters are expressed in a strict spatio-temporal manner correlating with different stages of lens development.

Our RT-PCR data indicate a clear switch from embryonic to postnatal expression of different components. Thus, the subunit composition of *GABA_A*R changes from the predominant embryonic $\alpha_{3}/\beta_{3}/\gamma_{1,2,3}/\delta$ to the predominant postnatal $\alpha_{1,4}/\beta_{2}/\gamma_{3}$ that

could result in changes in the receptor kinetics. For instance, an ontogenetic replacement of α_2 with α_1 has been shown to result in a change from a slow decay to fast decay receptor kinetics (Okada et al., 2000). Similarly, a reduction of $\alpha_{2,3}$ subunits with concomitant retainment of α_1 -containing, faster decay kinetics receptors in $GABA_A\beta_3^{-/-}$ mutant mice (Ramadan et al., 2003) or the compensatory up-regulation of α_2/α_3 (and down-regulation of $\beta_{2/3}$ and γ_2) subunits in $\alpha_1^{-/-}$ mutants (Kralic et al., 2002) implies that expression of $\alpha_{2,3}$ - β_3 - γ_2 is coordinately regulated during development. The presence and selective up-regulation of δ , a subunit, which confers high affinity to GABA, slow rate of de-sensitization and sensitivity to neuroactive steroids (Mihalek et al., 1999; Peng et al., 2004) at late gestational stages may indicate a specific function in the maturation of primary fibers that may be modulated by endogenous steroids. Similar role can be attributed to $\alpha_{2,3,4}$, β_3 and γ_1 , which can be differentially regulated by gonadal steroids (Pierson et al., 2005). Although $GABA_A\alpha_5$ gene is found clustered with γ_3 and β_3 in the same locus of mouse Chr 7, it does not seem to interact with γ_3 in brain (Togel et al., 1994). The exceptionally high level of γ_3 expression and the total absence of the closely positioned α_5 indicate that they are oppositely regulated at the transcriptional level throughout lens development. A coordinated transcriptional regulation has recently been demonstrated for the $\tilde{\beta}_2$ - $\tilde{\alpha}_6$ - $\tilde{\alpha}_1$ - $\tilde{\gamma}_2$ subunit gene locus as well (Uusi-Oukari et al., 2000). The lens affords an ideal model to study such phenomena.

In brain and peripheral tissues $GABA_B R1$ subunit is much more abundant than $GABA_B R2$, which is critical for trafficking the receptor to the membrane (Thuault et al., 2004). Similar correlation was observed for $GABA_B R$ of the prenatal lens. However, the extremely low level of $GABA_B R2$ in the postnatal lens may present a limitation for the formation of functional $GABA_B R$ heterodimers, and may suggest the existence of an alternative assembly of $GABA_B R1$ with other protein

partners (Calver et al., 2000).

In the lens, the GABA_AR (and probably GABA_BR) subunit switch strongly correlates with the switch from mainly apo-GAD65 and EGAD to the constitutively active GAD67 and gradual substitution of VGAT with membrane GABA transporters (GATs). In the adult brain GAD65 is greatly enriched in the presynaptic terminals and is thought to contribute mainly to the synaptic GABA pool (Battaglioli et al., 2003).

Our expression correlation data strongly suggest that in the embryonic lens GABA synthesized by GAD65 is released predominantly by VGAT and binds to GABA_AR $\alpha_{2,3}$ subunit-containing receptors, while that produced by GAD67 is released predominantly by the membrane GAT2 and binds to α_1 -containing GABA_AR. GABA produced by the enzymatically active GAD44, which is co-expressed with GAD65 (correlation coefficient 0.94) and is the embryonic counterpart of GAD67 (Szabo et al., 1994) may be released by inversion of one or more membrane GATs, as previously suggested (Szabo et al., 1994; Varju et al., 2001).

The high level of expression of GAT1-4 showing overlapping temporal profiles and especially the robust expression of GAT4 suggests that GAT-mediated GABA uptake/release coupled with Na⁺/Cl⁻ transport performs important function(s) in the lens. Furthermore, different GATs are selectively sorted to the apical or basolateral membranes where they may actively participate in the ion exchange, regulation of the ion homeostasis and cell volume (Schusboe et al., 2004). The developmentally earlier GAT3 is sorted to the apical membranes of polarized (Muth et al., 1998), and lens epithelia and apical/lateral membranes of differentiating lens fibers (Fig.3 and Fig.4, Kwakowski et al., 2008). compared to the postnatal GAT2, which is sorted predominantly to the basolateral membrane (Muth et al., 1998). In comparison, the axonal GAT1, localized preferentially to apical membranes of polarized epithelium (Muth et al., 1998) and apical/basal surfaces of lens epithelium and fibers (Fig.3,

Kwakowski et al., 2007) may carry out GABA transport at the fibers tips at stages of highest GABA production (E17.5-P0). The GABA transport across the lateral membranes is probably carried out by GAT3 during prenatal and GAT2 during postnatal stages. Finally, the predominantly radial glia-specific GAT4 (Jursky and Nelson, 1996) is pharmacologically unique among all transporters since it can operate as a Cl⁻ channel in the absence of Na⁺ (Karakossian et al., 2005). Further experiments will be needed to clarify the possible roles of the abundantly expressed membrane GABA transporters in the lens ion transport and fiber differentiation.

The expression pattern of GAD, GABA, VGAT and GABA receptors highly resembles that of some typical neuronal synaptic vesicular transport-related proteins like synapsin, synaptotagmin, synaptophysin (Frederikse et al., 2004) and Snap-25 (Wride et al., 2003) in the developing lens, which suggests the existence of a GABA signaling system similar to the synaptic GABA transmission in the nervous system. Like synapsin, VGAT, GABA_AR and GABA_BR are present in clusters at the tips of elongating fibers and apical membranes of the lens epithelium reminiscent of focal regions where vesicles interact with the fiber cell surface.

2.4. GABA receptors are functional in the developing lens: activation of GABA receptors triggers rise of intracellular calcium levels

In the adult brain, GABA, binding to GABA receptors exerts an inhibitory action, inducing membrane hyperpolarization and inhibition of Ca²⁺ currents. In immature brain activation of the ionotropic GABA_AR induces membrane depolarization and Ca²⁺ influx through voltage-gated Ca²⁺ channels, while the metabotropic GABA_BR is inhibitory, evoking membrane hyperpolarization and blockade (inhibition) of [Ca²⁺]_i.

GABA receptors are also expressed on non-excitabile cells, where GABA may acts as a trophic factor during the development similar to the CNS (Barker et al., 1998; Owens and Kriegstein, 2002a,2002b; Varju et al., 2001; Represa and Ben-Ari, 2005; Watanabe et al., 2002) but also non-neuronal tissues (pancreas, testis etc; Watanabe et al., 2002; Geigerseder et al., 2003; Franklin and Wollheim, 2004). In these structures GABA induces $[Ca^{2+}]_i$ rise either from internal stores or through voltage gated Ca^{2+} channels.

Using laser scanning confocal microscopy in freshly isolated P0 lenses loaded with Fluo-3/AM we detected simultaneous rise in $[Ca^{2+}]_i$ in epithelial cells and elongating fibers of the equatorial region upon bath application of GABA, indicating the existence of active GABA receptor response in this region.

To characterize the GABA receptor–induced calcium transients we have expanded this approach to an *in vitro* model system: detection of $[Ca^{2+}]_i$ changes in primary LEC cultures using selective GABA receptor agonists/antagonists and GABA. To date, lens cell lines and primary cultures derived from several species have been successfully used to study aspects of fiber cell differentiation and Ca^{2+} signaling pathways (Duncan et al., 1993; Churchill and Louis, 1998). The method, developed by us, was a novel approach by which we could obtain of primary LEC at different differentiation stages (epithelial cells, nucleated and anucleated fibers) found in native lenses and in addition they were far more permeable for Ca^{2+} imaging dyes than the native lens. Furthermore, most of the lens epithelial cells and lentoids, isolated from newborn mice and grown for two weeks in culture, expressed molecular components of GABA signaling: GAD, GABA transporters and GABA receptors found in the lens.

Surprisingly, both muscimol (GABA_AR agonist) and baclofen (GABA_BR agonist) evoked Ca^{2+} transients, predominantly in lentoids and sheets of epithelial cells, but

also in single cells, that could be blocked by the selective antagonists bicuculline and CGP55845, respectively. Whereas the GABA_AR-mediated Ca²⁺ rise in LEC is entirely consistent with the channel properties in the embryonic CNS (Ben-Ari et al., 2007) and/or non-neuronal cells (Watanabe et al., 2002; Franklin and Wollheim, 2004), the GABA_BR- evoked responses were unexpected, as it is widely considered to be inhibitory in both embryo and adult (Owens and Kriegstein, 2002; Watanabe et al., 2002; Lujan et al., 2005). However, more recent reports indicate that activation of GABA_BR can induce Ca²⁺ rise in both neurons (De Erausquin et al., 1992; Guatteo et al., 2004) and astroglia (Meier et al., 2008). In the latter case, the atypical GABA_BR responses displayed a variable and transient character mediated by Ca²⁺ release from IP₃- sensitive intracellular Ca²⁺ stores. Similarly we have found a greater variability in the LEC responses to baclofen compared to muscimol (data not shown), which indicates a dependence on threshold mechanisms. Alternatively, this variability could arise also from the superimposed inhibitory and excitatory effects of GABA_BR existing in different cells. However, the inhibitory component is probably small compared to the excitatory one, since application of the antagonist (CGP55845) alone usually had no effect, but occasionally evoked long-lasting oscillations, the mechanism of which is obscure.

Moreover, we have been able to record two additional specific features of the LEC responses - a synchronous reaction of cell groups, more pronounced at GABA_BR activation, which is often followed by wave-propagation of the signal and persistent synchronous oscillations and delayed Ca²⁺ transients, observed often upon GABA or baclofen, but never by muscimol applications. This wave-propagation of the signal strongly suggests the involvement of gap junctions known to mediate Ca²⁺ waves in primary LEC (Churchill et al., 1996,1998). Interestingly, persistent Ca²⁺ elevation (but not cell-to-cell Ca²⁺ waves) causes gap junction uncoupling and formation of

hemichannels, implying that GABA, acting on GABAR may regulate gap junction formation through $[Ca^{2+}]_i$ and thus ultimately influence the gap-junction-mediated lens transport.

The GABA signaling described here is a novel neurotransmitter signaling system described for the lens in addition to the previously characterized acetylcholine signaling (Duncan and Collison, 2003), which is involved in controlling cellular movements during lens development (Duncan and Collison, 2003 ; Oppitz et al., 2003). Based on previous (Oppitz et al., 2003) and present expression studies, the two systems may operate in the same epithelial and fiber cells to induce rise of $[Ca^{2+}]_i$ via GABAR and/or muscarinic receptors, respectively. Synergistic action of GABA and acetylcholine has been described previously for the chick retinal ventricular zone, where purinergic, muscarinic, GABAergic and glutamatergic stimulation all induce $[Ca^{2+}]_i$ rise (Pearson et al., 2002). Neurotransmitter co-expression and co-release is thought to be an important feature of neuronal progenitors that may provide for compensatory mechanisms during early development (Owens and Kriegstein, 2002). Another important feature of LEC is the GABAR (predominantly GABA_BR)- induced oscillation as it may have a long-term effect on the cell cycle and differentiation similar to the GABAR-mediated oscillations described for the gap junction-coupled embryonic neuroepithelial cells (LoTurco et al., 1995; Owens and Kriegstein, 1998; 2002).

2.4. Lens as a model to study the non-synaptic GABA signaling during early steps of differentiation

Our results show that all molecular components needed to form a functional GABA signaling system are expressed from early stages of lens development in a strict spatio-temporal manner, which parallels different phases of lens cell differentiation

suggesting that GABA may play diverse stage-specific function in this process. Structurally, GABA synthesis (GAD), release (VGAT, GAT) and GABAR binding are co-localized in epithelial and immature lens fiber cells that allow epithelial-to-epithelial epithelial-to-fiber and fiber-to-fiber GABA-mediated signaling (**Fig.41**).

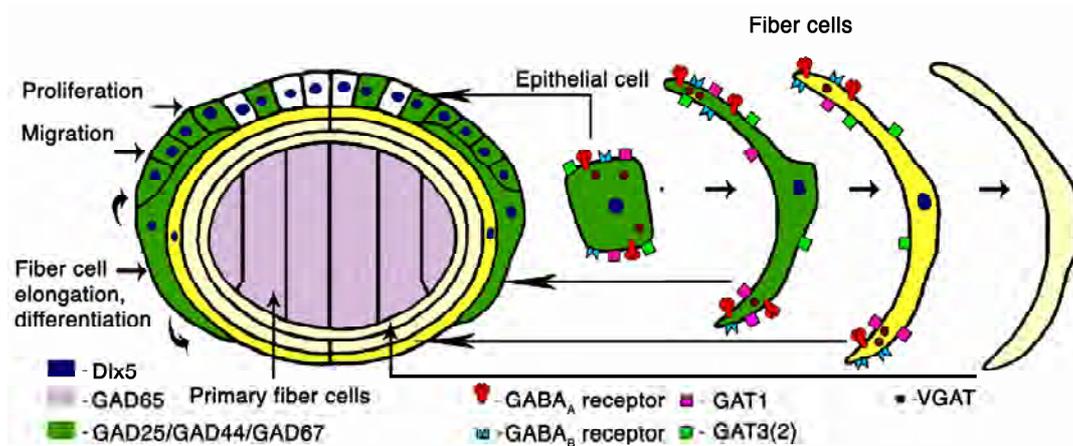


Fig.41. Expression patterns of different GABA signaling components in the neonatal mouse lens: a schematic presentation. GAD65 is localized almost exclusively in the primary lens fibers comprising the lens nucleus, while GAD67 is mostly found in the elongating secondary fibers and lens epithelium cells. Dlx5, an upstream regulator of GAD genes is expressed in the nuclei of the epithelial cells and fibers expressing GAD and GABA. GABA_A and GABA_B receptor subunits, VGAT and GAT1 show preferential expression in the apical/basal membranes (including the lens sutures) of both lens epithelial cells and elongating secondary fibers at the equatorial region. GAT3 is expressed in the apical/basal membranes of epithelial cells and the apical and lateral membranes of fiber cells, but not expressed in the basal tips of elongating secondary fibers. GAT2 may be predominantly localized to lateral membranes.

Our results suggest that GABA, predominantly located in the germinative and migratory zones of the lens epithelium and elongating primary or secondary fibers, act in a paracrine/autocrine fashion to modulate the cell cycle, cell migration and fiber elongation through modulation of the $[Ca^{2+}]_i$ level, by analogy with its similar role in the ventricular (germinative) and subventricular zones of the embryonic nervous system (see Introduction). Both VZ cells and lens epithelial cells are polarized epithelia connected via gap junctions that are permeable for Ca^{2+} signal propagation upon activation of the GABA receptors.

The fiber cell differentiation in the lens is thought to proceed through an unusual, attenuated apoptosis (Ishizaki et al., 1998; Wride, 2000; Bassnett et al., 2002; Zandy et al., 2005; Weber and Menko, 2005), a Ca^{2+} dependent process during which the lens loses all membrane organelles including the nuclei. GABA and GABA signaling components, initially highly expressed in nucleated immature fibers are gradually lost and become undetectable in terminally differentiated enucleated fibers implying that the GABA signaling may be one of the regulators of this process, acting through $[\text{Ca}^{2+}]_i$ changes.

The spatio-temporal segregation of the components downstream of the two kinetically different GAD forms- the predominantly embryonic GAD65 (and EGAD) vs postnatal GAD67 is probably related to the profoundly different mechanisms of denucleation of the primary vs secondary fibers (Wride, 2000).

Furthermore, GABA synthesized by GAD65 and EGAD may be released by VGAT at the apical/basal end or membrane GAT1-3 at both apical and basolateral membranes, while GAD67- produced GABA is released almost entirely by GAT2/3 at apical/basolateral surfaces of epithelial and fiber cells. Therefore, GABA pools synthesized by different GAD enzyme and released by distinct mechanisms may influence differentially the intercellular communications through adherens and gap junctions and/or reorganization of the membrane skeleton, both Ca^{2+} -dependent processes (discussed in the previous chapter). Selective sorting of the GABA transporters to the apical/basal surfaces, an equivalent of the axonal/dendritic neuronal compartments may have an active role in the osmo- and cell-volume regulation (Schousboe et al., 2004). Similar role can be proposed for the lens based on the high level of GAT/GABA expression at stages of most intensive fiber elongation.

In conclusion, the existence of GABA signaling machinery in the developing lens would suggest that it may be a target to a variety of drug treatments acting

predominantly on GABAR like benzodiazepines, hormones and neurosteroids and also in certain diseases affecting the GABA signaling, like IDDM. Furthermore, the lens is a dynamic organ that continues to grow throughout development and calcium signaling plays a key role in it. Several lines of evidence implicate the loss of calcium homeostasis in the lens as key factor in cataract formation (reviewed in Duncan and Wormstone, 1999). The loss of transparency during cataract development severely impairs the ability of the lens to focus on the image on the retina and is the leading cause of blindness worldwide. It would be important to study the effect of such treatments or pathological conditions on the lens transparency in order to develop preventive strategies.

VIII CONCLUSIONS

- Undifferentiated pluripotent embryonic stem (ES) cells express active GABA signalling system assembled from neuron-specific components- GAD, GABAR, GABA transporters and uptake.
- The temporal regulation of the two known GAD genes coding for the GABA-synthesizing enzymes GAD65 and GAD67 and the upstream transcriptional regulators Dlx2/5 follows a highly reproducible paradigm in different in vivo and in vitro differentiation systems.
- Based on our expression analysis, both vesicular and membrane transporter-mediated GABA release may occur in undifferentiated ES cells and during neuronal or lens differentiation, in the absence of inhibitory synapses.
- GABA_A and GABA_B receptors in ES cells and lens epithelial cells display unique composition and properties; both receptor types evoke intracellular Ca²⁺ rise by distinct mechanisms involving both the extracellular Ca (VGCC) and intracellular Ca stores (G-proteins).
- Despite the synergistic effect on the intracellular Ca²⁺ exerted by GABA_A and GABA_B receptors, they exert opposite effects on cellular proliferation and/or differentiation of ES cells. This effect is in sharp contrast to the receptors' properties in immature neurons, but may be similar to these in neural stem cells.
- GABA may serve as a paracrine –autocrine signaling molecule at very first stages of embryonic development affecting the cell proliferation, cell-cell communications and/or differentiation by mechanism(s) similar to that in the embryonic nervous system.
- Due to the spatio-temporal segregation of GABA signalling components along the two GAD proteins, the lens has proven a unique model system affording

the possibility to study the specific roles of GAD65 and GAD67 and components downstream of each gene, especially in view of their correlated regulation during development and in the adult.

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3. **Schwirtlich M**, Zs Emri, K Antal K, Z Máté, Z Katarova and G Szabó Activation of GABA_A and GABA_B receptors of distinct properties adversely affects proliferation of mouse embryonic stem (ES) cells through modulation of intracellular Ca²⁺ signalling. *Manuscript under revision.*

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IX SUMMARY

γ - aminobutyric acid (GABA) is the main inhibitory neurotransmitter in the CNS and as such regulates neuronal excitability. During development GABA acts as a trophic factor modulating processes such as proliferation, differentiation and migration in both neuronal and non-neuronal tissues. The aim of the present study was to characterize GABA signaling components and function during development in two different model systems: mouse embryonic stem (ES) cells and the ocular lens. Our results can be summarized as:

1. Undifferentiated pluripotent embryonic stem (ES) cells and differentiating lens cells express neuron-specific components necessary for an active GABA signaling system including GAD, GABAR and GABA transporters.
2. The temporal regulation of the two known GAD genes coding for the GABA-synthesizing enzymes GAD65 and GAD67 and the upstream transcriptional regulators *Dlx2/5* follows a highly reproducible pattern in the studied in vivo and in vitro differentiation systems.
3. Based on our expression analysis, both vesicular and membrane transporter-mediated GABA release may occur in undifferentiated ES cells and during neuronal or lens differentiation in the absence of inhibitory synapses.
4. GABA_A and GABA_B receptors in ES cells and lens epithelial cells display unique composition and properties; both receptor types evoke intracellular Ca²⁺ rise by distinct mechanisms involving both the extracellular Ca²⁺ and intracellular Ca²⁺ stores.
5. Despite the synergistic effect of GABA_A and GABA_B receptors on the intracellular Ca²⁺-concentration, they exert opposite effects on cellular proliferation and/or differentiation of ES cells. This effect is in sharp contrast to the receptors' properties in immature neurons, but may be similar to these in neural stem cells.

Our findings supports the view that GABA may serve as a paracrine–autocrine signaling molecule at early stages of embryonic development affecting cell proliferation, cell-cell communications and/or differentiation by mechanism(s) similar to that in the embryonic nervous system.

Using these two model systems will allow us to uncover the general mechanism of GABA action in basic developmental processes, which could also apply to the nervous system.

ÖSSZEFOGLALÁS

A gamma-aminovajsav (GABA) a központi idegrendszer legfontosabb gátló neurotranszmittereként kulcsszerepet játszik az idegsejtek ingerelhetőségének szabályozásában.

A GABA-nak az idegrendszer és a nem-idegi szövetek fejlődése során pedig szerepe van a proliferáció, differenciáció és migráció szabályozásában is. A disszertáció alapját képező kísérleteink legfőbb célja az volt, hogy jellemezzük a GABA jelátvitel összetevőit és azok működését az egyedfejlődés során. Ehhez két modellrendszert, az egér differenciálatlan embrionális őssejteket és a fejlődő szemlencsét használtuk. Jelentősebb eredményeink az alábbiakban foglalhatók össze:

1. A GABA jelátviteli folyamat összes komponense: a GABA, GAD, GABA_A, GABA_B receptorok valamint a GABA transzporterek jelen vannak az ES sejtekben és a fejlődő szemlencsében is, mintegy megteremtve a GABA szintézis, kibocsátás, érzékelés és újrafelvétel lehetőségét.
2. A GABA-szintetizáló enzimeket kódoló gének (GAD65 és GAD67) és azok szabályozásában szerepet játszó Dlx2/5 gének időbeli expressziós mintázata nagyon hasonló az általunk vizsgált *in vitro* és *in vivo* differenciációs rendszerekben.
3. Az expressziós vizsgálatunk eredményei szerint a vezikuláris és membrántranszport-mediált GABA-felszabadulás egyaránt előfordulhat a differenciálatlan ES sejtekben és a fejlődő szemlencsében is.
4. A differenciálatlan ES sejtekben és a lencse hámsejtekben a GABA_A és GABA_B receptorok sajátos alegység összetétellel és funkcionális sajátosságokkal rendelkeznek. Mindkét receptortípus aktiválódása sejtenbelüli kalcium-szint növekedést okoz, amelynek a mechanizmusa eltérő, és a Ca²⁺ mind az extracelluláris térből, mind pedig a belső raktárokból is származhat.
5. Annak ellenére, hogy a GABA_A és GABA_B receptorok hasonlóan befolyásolják az intracelluláris Ca²⁺-koncentrációt, a sejtosztódásra és differenciációra gyakorolt hatásuk ellentétes. Ez különbözik az éretlen neuronokban kifejtett hatástól, de hasonló lehet az ideg őssejtekben zajló folyamatokhoz.

Eredményeink alátámasztják azt a feltételezést, hogy a GABA auto- illetve parakrin faktorként befolyásolhatja a sejtosztódást, differenciációt és a sejtek közötti kommunikációt a korai embrionális fejlődés során.

Modellrendszerünk segítségével kideríthetjük a GABA hatás pontos mechanizmusát a fejlődési folyamatokban.

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