The α 1 Subunit in Sickness and in Health: Properties of the Most Predominant GABA_A Receptor Subunit and Implications of Its Dysfunction in Human Epilepsy

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The aim of this review is to highlight the role of the a1 subunit of the GABA_A receptor under normal conditions and to examine the consequences of its dysfunction in the context of epilepsy. First, background information relevant to GABA_A receptors is discussed, followed by a summary of the biophysical properties conferred by the a1 subunit and its developmental expression patterns. Next, key findings from a1 subunit knockout mice are reviewed. Lastly, the important role of the a1 subunit in regulating inhibitory tone in the CNS is highlighted by examining consequences of mutations in the a1 subunit implicated in generalized human epilepsy syndromes.

Keywords *GABA*, *GABA*, *receptor*, *heterogeneity*, *alpha-1 subunit*, *epilepsy*

Introduction

GABA_A receptors (GABA_ARs) are a family of chloride-selective, ligand-gated ion channels that mediate the majority of fast inhibition in the adult central nervous system¹. GABA_ARs belong to a larger superfamily of ligand-gated ion channels called Cys-loop receptors, which also includes nicotinic acetylcholine receptors, glycine receptors, and serotonin type III receptors^{2,3}. The GABA_AR gene family is comprised of at least 19 different subunits which are classified by sequence homology into 8 subtypes. These subunits can assemble in various combinations to produce functional GABA _ARs: α 1-6, β 1-3, γ 1-3, δ , ϵ , π , θ , and ρ 1-3⁴. Based on the large number of different GABA_AR subunits, there are seemingly myriad different subunit combinations possible, although only a subset of these theoretical combinations has been identified in vivo5. The cache of possible subunit combinations coupled with differing spatial and temporal expression patterns provides considerable structural and functional heterogeneity to GABA_ARs. Distinct subunit combinations produce distinct receptor isoforms which display highly variable properties throughout development and in adulthood⁶⁻⁹. This topic will be discussed in further detail below.

The vast majority of GABA_ARs exist as a combination of two α subunits, two β subunits, and either a γ or $\delta^{5,10}$ subunit arranged as shown in **Figure 1A-B**. The ε , π , and ρ 1-3 subunits are far less common and are generally positioned

in place of the γ or δ subunit, while the θ subunit can assume the position of the β subunit. The morphology of each GABA_AR subunit includes several characteristic features. A large extracellular domain at the N-terminus contains the characteristic disulfide bridge between two cysteine residues that creates the "Cys-loop" for which the receptor family is named. There are four helical transmembrane domains termed M1-M4, with the M2 of each subunit lining the ion pore. conntecting M3 and M4, there is a large intracellular domain, and beyond M4 is a very small extracellular C-terminal domain (**Figure 1C**)^{1,2,11}.

Full activation of GABA_ARs requires the binding of two molecules of the neurotransmitter γ -amino butyric acid (GABA)—one at each α/β subunit interface^{12,13}. Upon activation of the receptor, the channel opens and chloride (Cl⁻) flows down its electrochemical gradient through the pore. In the mature brain, this leads to an influx of Cl⁻, which causes the membrane potential of the cell to hyperpolarize and thus inhibits the generation of action potentials.

As previously mentioned, the wide variety of GABA_AR isoforms supports extensive functional heterogeneity, perhaps best demonstrated by the existence of two distinct forms of GABA_AR-mediated inhibition: tonic and phasic¹⁴. Tonic inhibition is mediated by extrasynaptically localized GA-BA_ARs largely comprised of a δ subunit with an α 4 and/ or an α 6 subunit¹⁵, though some extrasynaptic GABA_ARs containing α 5 and lacking a δ subunit are known to func-



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Figure 1: *Structure of GABA_A receptors.* A: Top down view of typical subunit arrangement with GABA binding site indicated by red ovals. B: Side view of receptor depicting chloride flowing through the pore. C: Morphology of an individual subunit.

tion extrasynaptically as well¹⁶. Extrasynaptic GABA_ARs are continuously activated by low levels (1M) of ambient GABA typically overflowing from the synaptic cleft. The properties of these receptors are conducive to persistent extrasynaptic activation, as they are highly sensitive to GABA, activate relatively slowly, and desensitize minimally¹⁷. The functional role of tonic inhibition seems to be the regulation of neuronal excitability. Conversely, phasic inhibition is mediated by synaptic GABA_ARs, which most often contain a γ subunit with $\alpha 1$, $\alpha 2$, and/or $\alpha 3$ subunits. Unlike their extrasynaptic counterparts, synaptic GABA_ARs are transiently activated by much higher (1mM) concentrations of GABA released from the presynaptic neuron into the synaptic cleft. In contrast to extrasynaptic GABA_ARs, synaptic GABA_ARs mediate tonic inhibition to allow for fast transmission of a fleeting signal and are thus less sensitive to GABA, activate rapidly, and desensitize extensively¹⁸. The short-lived activation of these receptors produces a transient inhibitory post-synaptic current (IPSC) whichtranslates presynaptic GABA release into a post-synaptic signal^{19,20}. The remainder of this review will focus on the most predominant subtype of synaptic receptors: those containing the a1 subunit. The distinct receptor properties and known roles of the $\alpha 1$ subunit, as well as important findings from a1 subunit knockout mice and recently discovered epilepsy-associated mutations in the $\alpha 1$ subunit are discussed below.

Biophysical Properties of the α 1 Subunit Relative to Other Synaptic α Subunits

The α 1 subunit is the most predominant α subunit in the adult brain and is most often assembled into the $\alpha 1\beta 2\gamma 2$ GABA_AR isoform comprising 60% of all GABA_ARs⁸. Further heterogeneity exists among synaptic receptors. The specific α subtype influences biophysical properties of GA-BA_ARs including GABA sensitivity and the rates of activation, deactivation, and desensitization²¹⁻²⁵. The primary strategy employed to determine the contribution of each α subunit to a particular property is to express different α subunits with the same β and γ subunit partners in heterologous cells and explore the properties of interest. One such property, GABA sensitivity, is defined by the concentration of GABA that is required to produce a given response. A common measure of GABA sensitivity is the concentration of GABA required to elicit a half-maximal response in a given receptor subtype, known as the EC50. A low EC50 indicates higher sensitivity and vice versa. The activation rate of a particular GABA, R is the rate at which receptor current increases from 10% to 90% of the maximal or peak current, and the deactivation rate is the rate at which current amplitude decreases after GABA is removed.

The desensitized state of the receptor is a high-affinity state in which GABA is bound but the ion channel is closed. The desensitization rate is the rate at which the response diminishes in continued presence of GABA. Each of the biophysical properties defined above influence the shape and time course of GABA_AR-mediated IPSCs. Thus, the identity of the α subunit within the receptor in large part

| Table 1: Relative Biophysical Properties Conferred by Synaptic α Subunits | |
|--|----------------------------------|
| GABA Sensitivity | $\alpha 1 > \alpha 2 > \alpha 3$ |
| Activation Rate | $\alpha 2 > \alpha 1 > \alpha 3$ |
| Deactivation Rate | $\alpha 1 > \alpha 2 > \alpha 3$ |
| Desensitization Rate | $\alpha 1 = \alpha 2 > \alpha 3$ |

Table 1: *Biophysical properties by synaptic* α *subunits.*

dictates the properties of inhibitory currents. The biophysical properties of $\alpha 1$ containing GABA_ARs relative to other synaptic α subunits are summarized in **Table 1**²¹⁻²⁴.

Interestingly, the properties of GABA, R mediated IPSCs are known to change throughout development^{26,27}—namely the decay kinetics which are heavily influenced by the deactivation and desensitization rates of the GAB-A_ARs^{28,29}. In fetal and early postnatal development, GABA_AR-mediated IPSCs decay relatively slowly; later in development, the IPSCs decay much more rapidly. The timing of this change coincides with the timing of a wellestablished developmental change in GABA_AR α subunit expression^{6,30,31}. Early in development, the $\alpha 2$ and $\alpha 3$ subunits predominate, but soon after birth their expression begins to wane while the expression of the $\alpha 1$ subunit steadily increases to become the most abundant α subunit by postnatal day 12 in mice⁶. A comparison of juvenile and mature GABA Rmediated IPSCs is shown in Figure 2. Given that the identity of the α subunit impacts IPSC properties, it is feasible that the developmental changes in α subunit expression and IPSC decay kinetics are causally linked. Indeed, it has been shown that in mice lacking the $\alpha 1$ subunit juvenile IPSC kinetics persist into adulthood³²⁻³⁴. The functional role of this developmental switch in α subunit expression and the concomitant change in IPSC kinetics is currently not well understood.

Findings From α1 Subunit Knockout Mice

In 2001, transgenic mice lacking the α 1 subunit of the GABA_AR exhibited a 50-60% decrease in the total number of GABA_ARs in the brain^{33,35}. Consistent with this finding, the expression of the β 2/3 and γ 2 subunits—the most common binding partners of the α 1 subunit—is also decreased in al knockout mice^{36,37}. Given that the α 1 subunit is the most abundant subunit and its loss results in a loss of the majority of GABA_ARs in the brain, it is very surprising that these animals are viable and lack any obvious phenotypic abnormalities aside from a slight handling-induced tremor. The fact that the mice are overtly normal could suggest that changes occur within the GABA R system to compensate for the loss of al. Indeed, posttranscriptional increases in the expression of the other α subunits have been observed^{36,37}, but the nature and extent of the compensation seems to vary among brain regions and has not been systematically quantified in the entire brain.

One study suggests that neurons upregulate the subunits they normally express rather than expanding their subunit repertoire³⁶, which is consistent with a post-transcriptional mode of upregulation. The consequences of these compensatory changes are not completely understood, but al knockout mice fail to develop mature IPSC kinetics32-34 and exhibit a lower threshold for pharmacologically induced seizures³⁸. This phenotype could indicate a decrease in inhibitory tone, although it was reported that a1 knockout mice did not experience spontaneous seizures³⁵. However, it is important to note that these conclusions were drawn by visual inspection only rather than EEG analysis. Certain types of seizures, such as absence seizures, result in very subtle alterations in behavior that are difficult to detect even in a human, much less in a mouse. Thus it is possible that these animals did have seizures but did not display any easily detectable seizure behavior.

The α1 Subunit and Epilepsy

Because GABA_ARs are the primary source

Electroencephalogram (EEG):

Method to measure electrical activity of the brain using electrodes on the surface of the scalp.

Missense Mutation:

Non-synonymous point mutation in which the identity of a single nucleotide is changed resulting in a codon that codes for an amino acid that differes from that of the WT protein at that particular location.



Figure 2: Developmental change in α subunit expression and IPSC kinetics. A: Juvenile α 3 containing receptor with much slower IPSC delay kinetics. B: Mature α 1 containing receptor with fast IPSC decay kinetics.

Endoplasmic Reticulum Associated Degradation:

A cellular process that marks misfolded proteins within the ER for ubiquitination followed by degradation via the proteasome.

Nonsense Mediated Decay:

Cellular mechanism to detect nonsense mutations (premature stop codons) and prevent translation of truncated proteins by degrading the mutant mRNA. of inhibition in the central nervous system, it is not surprising that several mutations in various GABA, R subunits have been identified in patients with idiopathic generalized epilepsy (IGE) syndromes such as childhood absence epilepsy and juvenile myoclonic epilepsy³⁹⁻⁴¹. Epilepsy is a large collection of syndromes diagnosed upon the occurrence of two or more unprovoked seizures. Epilepsy may be classified as IGE if the cause is thought to be genetic and the seizures appear to involve the entire brain simultaneously with no obvious focal origin. Most IGEs are thought to be multigenic, which renders developing animal models fairly difficult. However, the identification of monogenic forms of IGE has permitted the generation of animal models based on human disease-associated mutations. To date, there have been four mutations identified specifically in the $\alpha 1$ subunit in human patients suffering from idiopathic generalized epilepsy (IGE.) Two of these mutations, K353delins18X and D219N42, were only recently reported and have not been fully characterized. The other two however, A322D43 and S326fs328X44, have been extensively studied.

a1(K353delins18X) Mutation

The α 1(K353delins18X) mutation was identified in 1 unaffected and 3 affected individuals with IGE exhibiting generalized tonic-clonic seizures. These seizures manifest as a sudden tensing of skeletal muscles followed by rapid contractions and relaxations resulting in characteristic convulsions⁴⁵. This mutation involves the insertion of 25 base pairs into intron 10 which interrupts splicing and causes the retention of intron 10 in the transcript. The inclusion of intron 10 leads to an 18-amino acid insertion into the protein as well as the truncation of the fourth transmembrane domain due to a premature stop codon. Work in heterologous expression systems revealed that the protein is localized to the ER with complete loss of cell surface expression. In agreement with these findings, GABA-mediated currents were absent in these cells⁴². The fate of the mutant protein and the mechanism by which the $\alpha 1(K353delins18X)$ mutation leads to epilepsy is yet to be determined.

a1(D219N) Missense Mutation

The D219N missense mutation was identified in 4 of 5 affected individuals with IGE in a French-Canadian family exhibiting IGE or febrile seizures(FS)—so aptly named due to their coincidence with fever⁴⁵. Two of the four individuals with FS also reported a single generalized tonic clonic seizure. Studies conducted in heterologous expression systems indicate that surface expression of the mutant subunit is reduced as compared to WT a1 subunit, consistent with observations of decreased GABAevoked peak current amplitudes. Additionally, a1(D219N) subunit-containing receptors were shown to desensitize more rapidly than WT $\alpha 1$ subunit-containing receptors⁴². Further studies characterizing the effects of this mutation both in vitro and in vivo will be required to elucidate the mechanism by which it promotes the development of epilepsy.

a1(A322D) Missense Mutation

The A322D missense mutation was identified in 8 affected individuals within a large French-Canadian family suffering from a type of IGE called juvenile myoclonic epilepsy (JME). Myoclonic seizures are characterized by sudden, brief, involuntary jerks of the arms or legs⁴⁵. This mutation is autosomal dominant and results in the insertion of a charged aspartate residue in place of a highly conserved alanine within the M3 domain^{39-41,43}. Experiments in heterologous cells indicate that this mutation disrupts the insertion of M3 into the lipid bilayer which results in its retention in the endoplasmic reticulum (ER) and subsequent partial degradation through ER-associated degradation^{39-41,46} (ERAD). However, the mutant subunit is not completely degraded, as total and surface mutant protein is detectable, albeit at significantly lower levels than the WT a1 subunit^{39–41,46}. It has also been postulated that the $\alpha 1(A322D)$ mutation exerts a dominant negative effect by oligomerizing with and trapping WT subunits in the ER which are then subject to ERAD^{39,47}. Consistent with reduced surface expression, the peak GABA-evoked current through receptors containing the $\alpha 1(A322D)$ subunit was reduced 88%⁴⁷. In the mutant receptors, open time of the channel was considerably reduced. Additionally, a1(A322D) subunit-containing receptors exhibited substantially reduced sensitivity to GABA with a nearly 100-fold increase in the GABA EC50^{39,40}. It would be particularly enlightening to study the effects of this mutation in vivo and to that end, our group is in the process of generating an $\alpha 1(A322D)$ knock-in mouse line.

a1(S326fs328X) Frameshift Mutation

The $\alpha 1(S326fs328X)$ mutation is an autosomal dominant de novo mutation identified in a patient with childhood absence epilepsy. In contrast to the previously mentioned seizure types, absence seizures are not associated with any sort of convulsions or motor movements. Rather, they are characterized by sudden brief lapses in consciousness often accompanied by a blank stare⁴⁵. A single base pair deletion leads to a frameshift and premature termination codon (PTC) in the eighth exon which corresponds to the third transmembrane domain of the protein⁴⁴. The PTC has been shown to induce nonsense mediated decay (NMD) of the mutant mRNA, albeit incomplete. Mutant mRNA that escapes NMD is subsequently translated into truncated protein which is retained in the ER and subjected to ERAD^{48,49}. Thus, the $\alpha 1(S326fs328X)$ mutant subunit is not incorporated into the cell membrane and GABA-evoked currents are absent44,48.

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Based on these findings, it is thought that the epilepsy phenotype is a result of haploinsufficiency in the WT α 1 gene. This raises the intriguing possibility that heterozygous α 1 knockout animals could serve as a model for this disease. Indeed, our group recently revealed through EEG analysis that heterozygous α 1 knockout mice do in fact experience seizures, though they are absence seizures rather than convulsive seizures⁵⁰. This may explain why it was previously reported that α 1 knockout mice did not exhibit an epileptic phenotype as mentioned above. The seizures were greatly attenuated by treatment with ethosuximide (ETX), the most commonly prescribed drug for absence seizures in human patients. These findings represent a novel model of absence epilepsy based on a mutation identified in a human epilepsy patient.

Conclusions

GABA_ARs are a heterogeneous population of receptors and their properties are heavily influenced by their α subunit composition. The α 1 subunit is the most predominant subunit in the adult brain and is largely responsible for the maintenance of inhibitory tone in the CNS. As evidenced by the consequences of the dysfunction or loss of the $\alpha 1$ subunit, it also seems to be involved in epilepsy susceptibility. The transgenic $\alpha 1(A322D)$ knock-in and $\alpha 1$ knockout mouse lines that our group focuses on represent the first mouse models of genetic epilepsy based on mutations in the $\alpha 1$ subunit identified in human epilepsy patients. These two mutations were both identified in patients suffering from generalized epilepsy, but their seizure phenotypes were distinct. Based on our preliminary analyses, the epileptic phenotypes of the two mouse lines also differ. By dissecting the similarities and differences in pathogenesis behind these two models, we aim to identify common themes among all generalized epilepsies and also delineate important differences that contribute to distinct disease manifestations. While epilepsy is an exceptionally complicated and heterogeneous condition, the advent of animal models reflecting specific disease-associated mutations in GABA_ARs represents a promising and riveting new avenue for advancing our understanding of the pathogenesis of generalized epilepsy.

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This paper validates heterozygous 0.1 knockout mice as a model for absence epilepsy as it shows that the mice have absence-like seizures that remit with ethosuximide, the most commonly prescribed drug for the treatment of absence seizures.

Further information: http://www.mc.vanderbilt.edu/ neurology/faculty/gallagher.htm