

Chicken GABA_A receptor β 4 subunits form robust homomeric GABA-gated channels in *Xenopus* oocytes

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Abstract

Chicken GABA_A receptor β 4L and β 4S subunits were expressed in *Xenopus* oocytes by cRNA injection. Oocytes expressing either β 4 subunit alone or in combination with the chicken α 1 subunit were studied using the two-electrode voltage-clamp technique. Both the β 4L and β 4S subunits form homomeric GABA-gated Cl[−] channels with similar efficiencies. In comparison, oocytes expressing either the chicken α 1 or β 2S polypeptide show no or barely detectable GABA responses, as reported by others for most single-subunit vertebrate GABA_A receptors. The GABA-gated currents due to the β 4L-subunit homomer were not affected by the presence of actinomycin D during cRNA expression, indicating that nascent oocyte polypeptides are not required for channel formation. The homomeric β 4L-subunit receptors show high affinity for GABA with an EC₅₀ value of 4.3 ± 0.4 μ M and a Hill coefficient of 1.1 ± 0.1 ($n = 6$). In response to GABA application at the EC₂₅ value, currents elicited from the β 4L-subunit receptor are enhanced by 50 μ M pentobarbital ($110 \pm 10\%$, $n = 3$) and 10 μ M loreclezole ($60 \pm 3\%$, $n = 3$), inhibited by 10 μ M picrotoxinin ($93 \pm 3\%$, $n = 3$), but not affected by 1 μ M diazepam. These properties are similar to those found for oocytes expressing heteromeric chicken α 1 β 4L and α 1 β 2S receptors. Since the β subunits of GABA_A receptors provide essential determinants for receptor assembly and subcellular localization, homomeric β 4-subunit receptors are a useful model system for further study of the structure and function of GABA_A receptors. © 1998 Elsevier Science B.V. All rights reserved.

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1. Introduction

GABA is the predominant inhibitory neurotransmitter in the vertebrate brain. Most of the rapid actions of GABA occur, through GABA_A receptors on the postsynaptic membrane, by the gated opening of a Cl[−] channel integral to the receptor complex. The resulting increase in Cl[−] permeability of the postsynaptic membrane produces neuronal inhibition. GABA_A receptors are composed of subunits from six families (α , β , γ , δ , ϵ and π). There are

multiple isoforms within three of these subunit families (α 1-6, β 1-4 and γ 1-4) and additional diversity is produced by variations in the splicing of certain RNAs (reviewed by Darlison and Albrecht, 1995; Rabow et al., 1995). The function of GABA_A receptors is allosterically modulated by a number of therapeutically important agents including benzodiazepines, barbiturates, and neurosteroids (Macdonald and Olsen, 1994). The contribution of different subunits to some of the modulatory sites on GABA_A receptors has been elucidated. The benzodiazepine-binding site of the GABA_A receptor is determined by the type of α and γ subunit present, with the γ polypeptide being essential for benzodiazepine binding (Pritchett et al., 1989; Pritchett and Seeburg, 1990). Loreclezole, a broad-spectrum anticonvulsant compound, shows selective affinity for recombinant GABA_A receptors containing certain β subunits (Wingrove et al., 1994). More recently, it has been shown that reduced sensitivity of recombinant GABA_A

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receptors to general anaesthetic agents and pregnanolone is conferred by the ϵ and π subunits, respectively (Davies et al., 1997a; Hedblom and Kirkness, 1997).

As model systems, homomeric GABA_A receptors with properties that are comparable to native, heteromeric GABA_A receptors could be useful in the further investigation of the pharmacology, assembly, subcellular localization, and structure of this ion channel. Homomeric GABA_A receptors showing a limited number of native receptor properties have been reported after heterologous expression of certain subunits in *Xenopus* oocytes (Blair et al., 1988; Sigel et al., 1989, 1990; Sanna et al., 1995; Cestari et al., 1996; Krishek et al., 1996), human embryonic kidney 293 (HEK) cells (Pritchett et al., 1988; Po et al., 1990; Slany et al., 1995; Krishek et al., 1996; Davies et al., 1997b), and insect cell lines (Atkinson et al., 1992; Joyce et al., 1993). The GABA-gated currents elicited by homomeric receptors are typically much smaller than those produced by heteromeric receptors. Most of the functional homomeric GABA_A receptors reported are formed from β subunits, which have been shown to be a key component for robust expression of recombinant receptors (Sigel et al., 1990; Verdoorn et al., 1990). Homomeric β 1-subunit GABA_A receptors exhibit species-dependent differences in functional properties. Human and bovine β 1-subunit receptors yielded small GABA-gated Cl[−] currents (10–50 nA oocyte whole-cell currents elicited by 100 μ M agonist) that were potentiated by pentobarbital and inhibited by picrotoxin (Blair et al., 1988; Pritchett et al., 1988; Joyce et al., 1993; Sanna et al., 1995; Krishek et al., 1996), whereas murine and rat β 1-subunit receptors were not gated by GABA but showed spontaneous openings that were blocked by picrotoxin (Sigel et al., 1989, 1990; Krishek et al., 1996). On the other hand, rat β 3-subunit GABA_A receptors expressed in HEK cells showed neither spontaneous activity nor sensitivity to GABA, but they could be activated by pentobarbital (Davies et al., 1997b). There is no report of a functional homomeric β 2-subunit GABA_A receptor, in accord with the observation that individual murine α 1, β 2 and γ 2L subunits were not introduced to the surface membrane, but were retained within the endoplasmic reticulum of transfected HEK cells (Connolly et al., 1996a,b). However, Cestari et al. (1996) recorded pentobarbital-induced currents from murine β 2- and β 3-subunit homomers, which were unresponsive to GABA. Some studies demonstrated that GABA_A receptor β subunits are able to form receptors without incorporating endogenous polypeptides of either oocytes or HEK cells (Slany et al., 1995; Krishek et al., 1996), but other experiments show possible involvement of endogenous HEK polypeptides, including the GABA_A receptor β 3 subunit which may normally be present in HEK cells (Kirkness and Fraser, 1993; Fuchs et al., 1995; Davies et al., 1996; Ueno et al., 1996).

The GABA_A receptor β 4 subunit has, to date, been identified only in chick brain. There are two alternatively

spliced variants (β 4L and β 4S), with the β 4L subunit containing a 4 amino acid insert of unknown function (Bateson et al., 1991b). It is intriguing that neither a mammalian homologue of the β 4 subunit nor a chicken homologue of the β 1 polypeptide has been reported. Baumgartner et al. (1994) have shown that the β 4-subunit mRNA accumulates more rapidly than those encoding the α 1 and β 2 subunits during maturation of cultured neurons from the chick embryonic cerebral cortex. Immunoprecipitation and immunoblotting experiments (Tehrani et al., 1995) suggest that the β 4 subunit is a major component of GABA_A receptors in the chick cerebral cortex. However, the GABA_A receptor β 4 polypeptide has not been functionally expressed, either alone or in combination with other subunits. To investigate the properties of the β 4 subunit, *Xenopus* oocytes expressing homomeric and heteromeric GABA_A receptors were examined by the two-electrode voltage-clamp technique.

2. Materials and methods

2.1. Generation of full-length complementary DNAs (cDNAs)

A full-length β 2S-subunit cDNA was constructed by ligating part of the previously-reported chicken GABA_A receptor β 2L-subunit partial cDNA (~1.5 kb; Harvey et al., 1994), which lacks sequences that encode the carboxy-terminal part of the large intracellular loop and the fourth membrane-spanning domain (M4), to a polymerase chain reaction (PCR)-derived cDNA fragment. This latter fragment was generated using primers 5'-AGTAAGCTTTG G T C A A C T A C A T C T T C T T T G -3' and 5'-T A A C T C G A G T T C T T G C T T C C T G T G T G G C T T -3'; the former recognizes the sequence that specifies the amino acids Glu-Tyr-Ala-Leu-Val-Asn-Tyr-Ile-Phe-Phe which span the end of the third membrane-spanning domain (M3), while the latter is complementary to part of the 3'-untranslated region. Amplification of random-nonamer primed embryonic day-18 chick brain first-strand cDNA was for 35 cycles of 94°C for 1 min, 65°C for 1 min and 72°C for 1 min. The resultant products (497 bp; β 2S-subunit cDNA, and 548 bp; β 2L-subunit cDNA) were digested with *Hinc*II and *Xho*I, at unique restriction endonuclease recognition sites (underlined) that were incorporated into the PCR primers, and cloned into *Hinc*II- and *Xho*I-restricted pBluescript SK+ yielding plasmids pcGR β 2S.3'end and pcGR β 2L.3'end, respectively. Finally, the pBluescript II SK+ plasmid containing the ~1.5 kb β 2L-subunit partial cDNA (Harvey et al., 1994) was digested with *Eco*RI (which cuts at the 5' end of the insert) and *Hinc*II (which cuts at a sequence that encodes the end of M3), and the released cDNA fragment was ligated into *Eco*RI- and *Hinc*II-restricted pcGR β 2S.3'end. The resultant construct was completely sequenced.

A full-length cDNA encoding the $\beta 4L$ subunit (previously called the $\beta 4'$ subunit) was constructed by ligating a previously-reported chicken GABA_A receptor $\beta 4L$ -subunit partial cDNA (Bateson et al., 1991b), which lacks sequences that encode the carboxy-terminal part of the large intracellular loop and M4, to a PCR-derived cDNA fragment. This latter fragment was generated using primers 5'-GGCTCAGGAATTCAGTTCCGCAAGCCACTG-3' and 5'-CTATCAAGCTTGGAGGCTGCAGGCATCA-3'; the former recognizes the sequence that specifies the amino acids Gly-Ser-Gly-Ile-Gln-Phe-Arg-Lys-Pro-Leu in the large intracellular loop, while the latter is complementary to part of the 3'-untranslated region. Amplification of random-nonamer primed 1-day-old chick brain first-strand cDNA was for 30 cycles of 94°C for 1 min, 65°C for 1 min and 72°C for 1 min. The resultant product (289 bp) was digested with *EcoRI* and *HindIII*, at unique restriction endonuclease recognition sites (underlined) that were incorporated into the PCR primers. Next, the pBluescript plasmid containing the $\beta 4L$ -subunit partial cDNA (Bateson et al., 1991b) was completely digested with *HindIII* (which recognizes a unique site within the vector polylinker, 3' of the cDNA sequence), and then partially digested with *EcoRI*. The resultant mixture of products was ligated together with the *EcoRI*- and *HindIII*-digested PCR-generated fragment, and a construct containing a full-length $\beta 4L$ -subunit cDNA (pcGR $\beta 4L$) was selected by restriction mapping and fully sequenced.

A full-length cDNA encoding the $\beta 4S$ subunit (previously called the $\beta 4$ subunit; Bateson et al., 1991b), which lacks the amino acid sequence Val-Arg-Glu-Gln in the large intracellular loop, was generated by in vitro mutagenesis of plasmid pcGR $\beta 4L$ using the method of Kunkel (Sambrook et al., 1989). Single-stranded DNA was mutagenized using the antisense oligonucleotide 5'-ACCGTAAGGGTTCGACCTCTTTTCTTCATA-3', which is complementary to the sequences that flank the 12-bp insertion in the $\beta 4L$ -subunit cDNA. Note that this primer contains a single mismatch to the published sequence (Bateson et al., 1991b) which generates a *SalI* site (underlined); this permitted the rapid detection of mutated cDNAs but does not change the encoded amino-acid sequence. Fourteen mutated plasmids were identified. One of these was fully sequenced to confirm that the 12 nucleotides had been deleted and to check the integrity of the complete $\beta 4S$ -subunit coding sequence.

The chicken GABA_A receptor full-length $\alpha 1$ -subunit cDNA used in these studies was that described by Bateson et al. (1991a).

2.2. cRNA synthesis

cRNAs were synthesized using the T3 and T7 mMES-SAGE mMACHINE kits (Ambion) following the protocol recommended by the manufacturer. cRNAs were ethanol precipitated, resuspended in 0.1 M KCl at a final concentration of 1 μ g/ μ l, and stored at -80°C until use.

2.3. Oocyte preparation and microinjection

Xenopus oocytes were prepared and injected following routine protocols (Parent and Gopalakrishnan, 1995). De-folliculated oocytes were injected with 50 nl of solution containing a single GABA_A receptor subunit cRNA, or combinations of $\alpha 1$ - and β -subunit cRNAs each at a concentration of approximately 600 ng/ μ l. In experiments differentiating chicken GABA_A receptor subunit expression from that of the host cell, oocytes were incubated in the presence of actinomycin D (50 μ g/ml) following cRNA injection to inactivate transcription of *Xenopus* genes.

2.4. Electrophysiological recording

Whole-cell inward currents in response to GABA were measured by two-electrode voltage-clamp (Oocyte clamp OC-725B, Warner Instrument) 4–8 days after cRNA injection. During electrophysiological recording, oocytes were voltage-clamped between -60 and -70 mV and perfused with modified Barth's medium (in mM: 88 NaCl, 1 KCl, 2.4 NaHCO₃, 10 HEPES, 0.82 MgSO₄, 0.33 Ca(NO₃)₂, 0.91 CaCl₂, pH 7.4) at 6 ml/min throughout the experiment. GABA_A receptor modulators were applied in the perfusate for 30 s before GABA addition. GABA was applied for 30–45 s or until the peak of the response was observed. Intervals of 10–15 min were allowed between applications to ensure full recovery from desensitization. Each batch of oocytes expressing GABA_A receptors was tested for their ability to produce consistent GABA-induced currents by repeatedly applying 100 μ M GABA during a period of at least 90 min. GABA_A receptor modulators were dissolved in dimethylsulphoxide (DMSO) prior to dilution into the perfusate. The resulting levels of DMSO ($\leq 0.1\%$) had no effect on either non-injected, sham-injected or GABA_A receptor subunit cRNA-injected oocytes. Loreclezole was a gift from Janssen Laboratories, diazepam was donated by Hoffmann-La Roche, and picrotoxinin and pentobarbital were purchased from Sigma.

2.5. Data analysis

The computer software CLAMPX 6.0 from pCLAMP (Axon Instruments) was used for on-line data acquisition and analysis. Membrane currents were digitally sampled at 5 kHz (200 μ s/data point) and simultaneously traced by chart recorder. To study the current-voltage relationship of the GABA-induced current, discontinuous voltage pulses (-70 to -10 mV) of 200 ms duration were applied from a holding potential of -70 mV before and after application of 100 μ M GABA. The peak amplitude of the GABA-evoked currents at various potentials was normalized by assigning 100% to the value at -70 mV. All measurements for the GABA dose-response curves were standardized by assigning 100% to the current amplitude

elicited by 1 mM GABA. All measurements of the effects of GABA_A receptor modulators were standardized by assigning 100% to the current amplitude evoked by an approximately EC₂₅ concentration of GABA (predetermined for each individual oocyte based on the maximum current response induced by 2 mM GABA). Dose–response curves were calculated using a nonlinear regression fit (Marquardt–Levenberg algorithm, SigmaPlot v. 2.0, Jandel Scientific) of the equation: $I_C/I_{\max} = 1/(1 + (EC_{50}/C)^n)$, where C is the GABA concentration, I_C is the current elicited by GABA at concentration C , I_{\max} is the current elicited by 1 mM GABA, and n is the Hill coefficient.

3. Results

Co-expression of the chicken GABA_A receptor $\alpha 1$ and $\beta 4L$ subunits in *Xenopus* oocytes resulted in a GABA-gated inward current (Fig. 1A). The peak response to 100 μ M GABA (1.9 ± 0.3 μ A, mean \pm S.E.; $n = 4$ oocytes) was comparable to that of oocytes co-expressing the chicken $\alpha 1$ and $\beta 2S$ subunits (2.5 ± 0.3 μ A, $n = 4$).

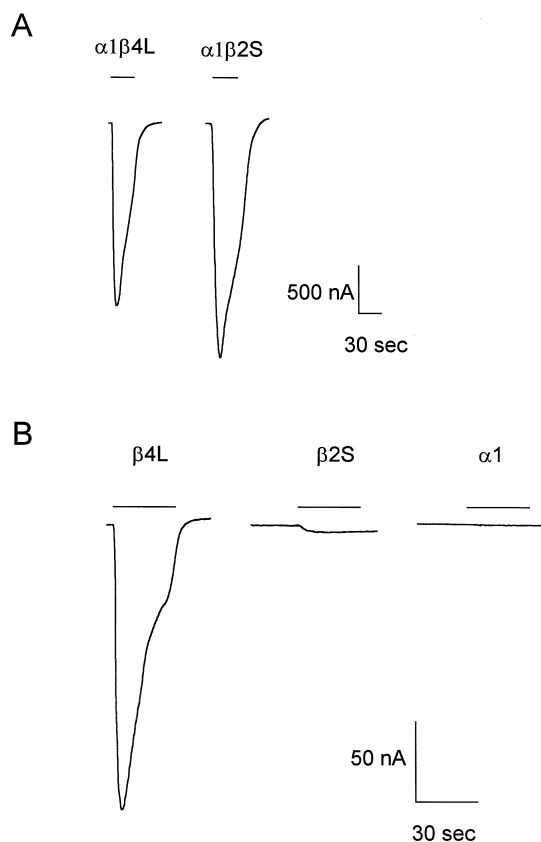


Fig. 1. Membrane currents in response to the application of GABA. (A) Oocytes expressing either $\alpha 1\beta 4L$ - or $\alpha 1\beta 2S$ -subunit receptors. (B) Oocytes expressing either the $\beta 4L$, $\beta 2S$ or $\alpha 1$ subunit alone. GABA (100 μ M) was applied for the times indicated by the horizontal bars above the current traces.

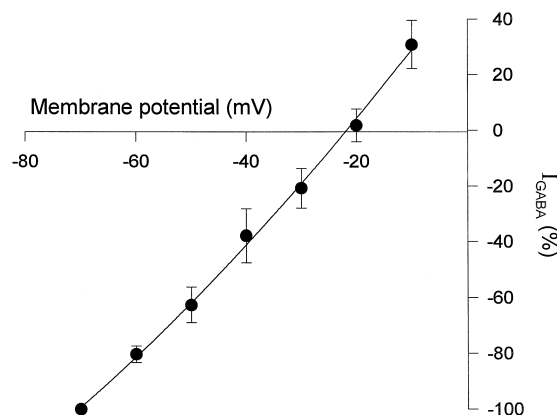


Fig. 2. Current–voltage relationship of the GABA-induced current from oocytes expressing homomeric $\beta 4L$ -subunit GABA_A receptors. Before application of 100 μ M GABA and during the peak current, discontinuous voltage pulses (–70 to –10 mV) of 200 ms duration were applied from a holding potential of –70 mV. The amplitudes of the GABA-induced currents at various potentials were normalized by assigning 100% to the value obtained at –70 mV and are expressed as the mean \pm S.E. ($n = 3$ oocytes).

Injection of the $\beta 4L$ -subunit cRNA alone (Fig. 1B) produced receptors that also yielded a GABA-gated inward current with a peak response of 110 ± 10 nA ($n = 8$). Similar currents were obtained with $\beta 4S$ -subunit channels (not shown). In paired experiments similar to those shown in Fig. 1B, the peak currents evoked by 100 μ M GABA from $\beta 4S$ -subunit receptors were not significantly different from those of $\beta 4L$ -subunit channels when analyzed by t -test ($t = 0.498$, $df = 3$, P -value = 0.653). However, expression of the chicken GABA_A receptor $\alpha 1$ subunit or $\beta 2S$ subunit alone resulted in no or a barely detectable response to the application of 100 μ M GABA (Fig. 1B). GABA had no effect on either non-injected or sham-injected oocytes (not shown).

Sigel et al. (1990) have suggested that currents associated with homomeric $\beta 1$ -subunit GABA_A receptors may be due to incorporation of endogenous oocyte polypeptides. In order to investigate this possibility, oocytes injected with the $\beta 4L$ -subunit cRNA were incubated with actinomycin D (50 μ g/ml) to inactivate *Xenopus* gene transcription. Similar responses to 100 μ M GABA were observed after incubation either with (130 ± 14 nA, $n = 3$) or without (110 ± 16 nA, $n = 3$) actinomycin D. To characterize the GABA-gated current attributed to the $\beta 4$ -subunit homomer, the current–voltage relationship of the inward current induced by 100 μ M GABA was examined (Fig. 2). The reversal potential was –20 mV, consistent with the equilibrium potential for Cl^- in oocytes (Barish, 1983).

The relationship of the peak current amplitudes to the concentration of applied GABA is shown in Fig. 3. The curves represent the best fit by Marquardt–Levenberg algorithm for the least-square solution of the parameters (EC₅₀ value and Hill coefficient). It is apparent that the

homomeric $\beta 4L$ -subunit receptor shows a high affinity for GABA. The arithmetic mean of individual EC_{50} values from 6 oocytes injected with $\beta 4L$ -subunit cRNA was $4.3 \pm 0.4 \mu M$ and the Hill coefficient was 1.1 ± 0.1 . When the GABA_A receptor $\beta 4L$ subunit was co-expressed in oocytes with the $\alpha 1$ subunit, a lower affinity for GABA was observed ($EC_{50} = 16 \pm 3 \mu M$, Hill coefficient = 1.3 ± 0.2 , $n = 3$). A similar dose–response curve (Fig. 3) was obtained for heteromeric $\alpha 1\beta 2S$ receptors ($EC_{50} = 14 \pm 2 \mu M$, Hill coefficient = 1.1 ± 0.2 , $n = 3$). At the lowest GABA concentration tested ($1 \mu M$), the data for both heteromeric receptors show a small deviation from the fitted curves. However, this is unlikely to significantly affect the resulting EC_{50} values.

The allosteric modulation of GABA-elicited currents from the $\beta 4L$ -subunit GABA_A receptor was also investigated (Fig. 4). At GABA concentrations representing the EC_{25} value, the peak currents were enhanced by $50 \mu M$ pentobarbital ($110 \pm 10\%$, $n = 3$) and by $10 \mu M$ loreclezole ($60 \pm 3\%$, $n = 3$). In the presence of $10 \mu M$ picrotoxinin, the responses were inhibited by $93 \pm 3\%$ ($n = 3$), while $1 \mu M$ diazepam had no significant effect. Similar modulatory effects were also observed for heteromeric $\alpha 1\beta 4L$ and $\alpha 1\beta 2S$ receptors (Fig. 4) with $50 \mu M$ pentobarbital enhancing ($110 \pm 17\%$ and $150 \pm 23\%$, respectively), $10 \mu M$ loreclezole enhancing ($65 \pm 7\%$ and $87 \pm 2\%$, respectively), and $10 \mu M$ picrotoxinin inhibiting ($87 \pm 2\%$ and $89 \pm 3\%$, respectively) the GABA-gated currents. Higher concentrations of pentobarbital, which presumably could directly gate channels in the absence of GABA, were not tested. Under conditions similar to those

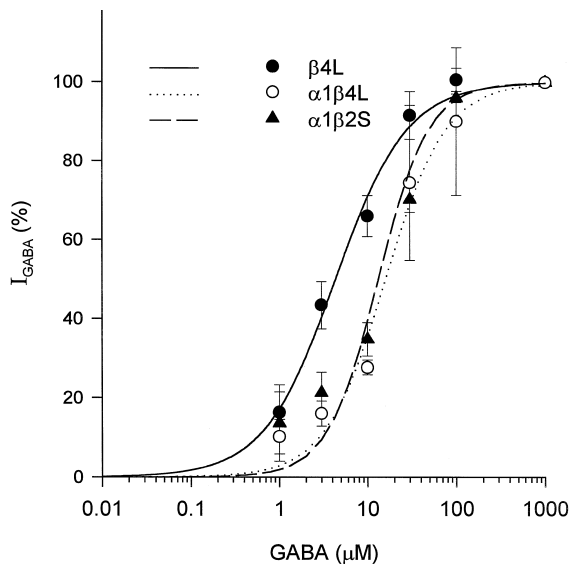


Fig. 3. GABA dose–response curves for homomeric $\beta 4L$ -subunit and heteromeric $\alpha 1\beta 4L$ - and $\alpha 1\beta 2S$ -subunit GABA_A receptors. The peak currents were normalized to the values obtained with 1 mM GABA and represent the mean \pm S.E. The values for n (number of oocytes), the Hill coefficients, and the EC_{50} values are given in the text. Individual dose–response curves were fitted as described in Section 2.5.

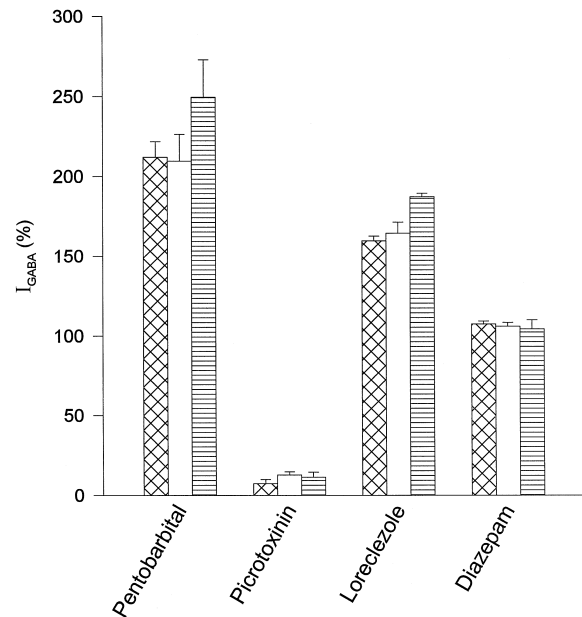


Fig. 4. Effect of $50 \mu M$ pentobarbital, $10 \mu M$ picrotoxinin, $10 \mu M$ loreclezole and $1 \mu M$ diazepam on homomeric $\beta 4L$ -subunit and heteromeric $\alpha 1\beta 4L$ - and $\alpha 1\beta 2S$ -subunit GABA_A receptors. After application of the indicated modulator for 30 s, GABA at the appropriate EC_{25} value, determined from Fig. 3, was introduced into the perfusate. Values for the peak GABA-gated currents are expressed as a percentage of controls (100%) in which the modulator was omitted and represent the mean \pm S.E. ($n = 3$).

used to generate the data shown in Fig. 4, $\beta 4L$ -subunit receptors did not elicit detectable currents during application of either pentobarbital, loreclezole or picrotoxinin in the absence of GABA (deflections $< 3 \text{ nA}$; $n = 4$; not shown). Likewise, no response to the application of any of these modulators ($n \geq 4$) was observed with oocytes injected with either the $\alpha 1$ - or $\beta 2S$ -subunit cRNA alone or with non-injected or sham-injected oocytes.

4. Discussion

These experiments provide the first description of GABA_A receptor $\beta 4L$ -subunit function. A combination of chicken GABA_A receptor $\beta 4L$ and $\alpha 1$ subunits co-expressed in *Xenopus* oocytes yielded GABA-gated currents of a magnitude similar to those recorded from an $\alpha 1\beta 2S$ -subunit combination. Oocytes injected with the $\beta 4L$ -subunit cRNA alone also showed GABA-evoked currents which were approximately 10% of those obtained with the $\alpha 1\beta 4L$ -subunit receptor. Since both the $\beta 4S$ and the $\beta 4L$ homomer gave similar sized GABA-gated currents, it seems unlikely that the additional sequence of 4 amino acids in the predicted large intracellular loop region of the $\beta 4L$ subunit has a substantial effect on receptor assembly. The failure of actinomycin D to affect the expression of homomeric $\beta 4$ -subunit currents appears to rule out potential

contributions of nascent oocyte polypeptides in the assembly of these channels. For the formation of GABA_A receptor β 1-subunit channels, such endogenous polypeptides have been suggested to be required (Sigel et al., 1990).

The β 4L-subunit GABA_A receptor has a much higher apparent affinity for GABA than other homomeric GABA_A receptors reported previously (cf. Blair et al., 1988). Interestingly, the GABA EC₅₀ value for the homomeric β 4L-subunit channel is also lower than that of the heteromeric α 1 β 4L- and α 1 β 2S-subunit receptors (Fig. 3). It has been suggested that the GABA binding site in GABA_A receptors is formed by the interface between α and β subunits (Smith and Olsen, 1995). The proposed locus of GABA binding includes Phe⁶⁴ and flanking residues of the α 1 subunit (Sigel et al., 1992; Smith and Olsen, 1994) and two motifs in the β 2 subunit, one surrounding Tyr¹⁵⁷ and the other at Tyr²⁰⁵ (Amin and Weiss, 1993). The GABA_A receptor β 4-subunit sequence (Bateson et al., 1991b) shows conservation of these β 2-subunit motifs, but lacks that in the α 1 subunit. Nevertheless, our data imply that the interface between two β 4 subunits is capable of forming a site whose affinity for GABA gating is similar to that of native GABA_A receptors.

At GABA concentrations near the EC₂₅ value, pentobarbital and picrotoxinin give similar potentiation and inhibition, respectively, with homomeric β 4L- and heteromeric α 1 β 4L- and α 1 β 2S-subunit GABA_A receptors. In the absence of GABA, application of either pentobarbital or picrotoxinin has been reported to modulate the currents from misassembled GABA_A receptors (Krishek et al., 1996). However, in our studies when applied alone, neither compound had an effect on either non-injected oocytes or sham-injected oocytes, or on oocytes expressing chicken GABA_A receptor subunits. Therefore, the GABA-gated currents from homomeric β 4L-subunit channels are not artifacts of improperly assembled receptors.

The affinity for loreclezole is more than 300-fold higher with GABA_A receptors containing either a β 2 or a β 3 subunit than those having a β 1 subunit. This selectivity is determined by a single amino-acid residue at the distal end of M2, Asn²⁸⁹ and Asn²⁹⁰ in human GABA_A receptor β 2 and β 3 subunits, respectively, and Ser²⁹⁰ in the β 1 subunit (numbering from the amino-terminal methionine residue; Wingrove et al., 1994). The predicted amino acid sequences of the chicken β 4 and β 2 subunits (Bateson et al., 1991b; Harvey et al., 1994) show that both contain Asn at this position (Asn²⁸⁹). Co-application of 10 μ M loreclezole together with a GABA concentration at the EC₂₅ value enhanced the currents from homomeric β 4L- and heteromeric α 1 β 4L- and α 1 β 2S-subunit receptors to a similar degree (60–87%). Since 10 μ M loreclezole produced much smaller enhancements at GABA_A receptors containing a β 1 subunit (Wingrove et al., 1994), it is apparent that the β 4 subunit functionally resembles the β 2 and β 3 subunits. This is consistent with the conservation of Asn, in M2, in the β 2, β 3, and β 4 polypeptides.

In the putative intracellular loop region between M3 and M4, the chicken GABA_A receptor β 4 subunit shows substantial divergence from the mammalian β 1- and chicken β 2- and β 3-subunit sequences. This points to possible novel properties for the β 4 polypeptide. One of these is the ability to form robust homomeric agonist-gated channels, a property not found among other vertebrate GABA_A receptor β subunits. Furthermore, investigation reveals that homomeric β 4-subunit receptors possess a very high affinity site for GABA. Although such homomeric receptors are unlikely to occur in vivo, we believe that the β 4-subunit homomer will prove useful for examining receptor assembly and trafficking as well as the relationship between molecular structure and receptor function.

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