A single amino acid in γ -aminobutyric acid ρ 1 receptors affects competitive and noncompetitive components of picrotoxin inhibition

(Xenopus oocytes/mutagenesis/chloride channel/ligand-gated neuroreceptor)

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ABSTRACT A class of bicuculline-insensitive γ -aminobutyric acid (GABA) receptors, GABA_C, has been identified in retina. Several lines of evidence indicate that GABA_C receptors are formed partially or wholly of GABA ρ subunits. These receptors generate a Cl⁻ current in response to GABA but differ from GABA_A receptors in a number of ways. Picrotoxin, widely accepted as a noncompetitive antagonist of GABA_A receptors, displays competitive and noncompetitive antagonism of GABA_C receptors in perch and bovine retina and GABA ρ 1 receptors expressed in *Xenopus* oocytes. The aim of this study was to identify the molecular basis of the two components of picrotoxin inhibition of GABA ρ 1 receptors. By using a domain-swapping and mutagenesis strategy, a difference in picrotoxin sensitivity between ρ 1 and ρ 2 receptors was localized to a single amino acid in the putative second transmembrane domain. Substitution of this amino acid with residues found in the analogous position in highly picrotoxin-sensitive glycine α and GABA_A subunits increased the sensitivity of ρ 1 mutants 10- to 500-fold. Importantly, the competitive component of picrotoxin inhibition of the ρ 1 mutant receptors was almost eliminated. These findings demonstrate that an amino acid in the putative channel domain of GABA ρ 1 receptors influences picrotoxin sensitivity and mediates agonist binding by an allosteric mechanism.

Picrotoxin, a plant-derived compound, is a noncompetitive antagonist of γ -aminobutyric acid type A (GABA_A) receptors and certain glycine receptors (1). Picrotoxin may exert its effect on these receptors by blockade of the ion conduction pore (1-4). However, in crustacean muscle, picrotoxin has been observed to inhibit GABAA receptors in a competitive manner, suggesting an allosteric mechanism of action (5, 6). Binding studies of picrotoxin analogues and single-channel studies of picrotoxin action on rat sympathetic neurons are consistent with this concept (1, 7). Recently, a class of picrotoxin-inhibitable GABA receptors has been characterized in the retina of several species (5, 8-11). These receptors, variously termed bicuculline-insensitive, ρ -like, and GABA_C, generate a Cl⁻ conduction in response to GABA that is insensitive to bicuculline, benzodiazepines, and barbiturates (5, 12, 13). These receptors are referred to as GABA_C, although their precise definition remains a matter of debate. GABA ρ subunits cloned from retina share many of these pharmacologic properties, suggesting that GABA_C receptors are composed partially or entirely of ρ subunits (9, 13–18).

The picrotoxin sensitivity of $GABA_C$ receptors is of interest because it differs from $GABA_A$ receptors in several ways. First, $GABA_C$ receptors in mammalian retina are less sensitive to picrotoxin than $GABA_A$ receptors located in retina or in cortex (9, 12). Second, the inhibitory effects of picrotoxin have

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a weak use-dependent component compared to $GABA_A$ receptors (12). Third, picrotoxin inhibition of $GABA_C$ receptors in perch and bovine retina depends on agonist concentration, suggesting a complex type of antagonism (12, 19). Together, these data suggest that the site and mechanism of picrotoxin action on $GABA_C$ receptors may differ from $GABA_A$ receptors.

The goal of the present study was to identify amino acids underlying the picrotoxin inhibition of GABA receptors formed of $\rho 1$ subunits. Like GABA_C receptors, GABA $\rho 1$ and ρ^2 receptors expressed in *Xenopus* oocytes are considerably less sensitive to picrotoxin than homooligomeric or heterooligomeric receptors formed of GABAA subunits, and GABA ρ 1 receptors display both noncompetitive and competitive components of picrotoxin inhibition (18-20). Differences in picrotoxin sensitivity of $\rho 1$ and $\rho 2$ homooligometic GABA receptors facilitated identification of a single position in the $\rho 1$ subunit critical for picrotoxin sensitivity. Mutating this residue to amino acids found in the analogous position of GABAA or glycine subunits significantly increased picrotoxin sensitivity over wild-type $\rho 1$ and virtually abolished the agonist-dependent elements of picrotoxin inhibition. Therefore, both components of picrotoxin of inhibition of GABA ρ 1 receptors are determined by the same amino acid.

MATERIALS AND METHODS

Construction of Chimeric cDNAs. Chimeric DNA fragments coding for the indicated regions of $\rho 1$ and $\rho 2$ as shown in Fig. 1 were generated by two-step PCR (21) on $\rho 1$ and $\rho 2$ cDNA templates by using three oligonucleotide primers. The primers used for constructing the $\rho 1\rho 2$ -A chimeric cDNA were as follows (boldface indicates $\rho 1$ cDNA sequence):

5'- ρ 1 primer: 5'-ATA <u>GGATC</u> ATGTTGGCTGTCCCAAAT -3'
BamHI

 $\rho 1 \rho 2$ -A hybrid primer:

5'-**CAGACGGTTGTACCAGCCA**GTGCTGCTGTAGAAAGCCAGT-3' 3'-ρ2 primer: 5'-TATCTGCAGCCCCCTC<u>GATGTC</u>TA-3' PstI

The first step of PCR amplified an 850-bp fragment using 0.1 μ M 5'- ρ 1 primer and 0.01 μ M ρ 1 ρ 2-A hybrid primer and ρ 1 cDNA template. Since the hybrid primer contained 20 bp of ρ 2 cDNA sequences (boldface), the resulting 850-bp PCR product contained this sequence at the 3' end and served as the 5' primer for a second PCR using 0.1 μ M 3'- ρ 2 primer and 1 ng of ρ 2 cDNA as template. The presence of the 5'- ρ 1 primer in the reaction mixture enabled the amplification of a 1568-bp chimeric cDNA (Fig. 1). The ρ 1 ρ 2-A chimeric cDNA was

Abbreviations: GABA, γ -aminobutyric acid.

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produced by the same procedure using the following PCR primers (boldface indicates $\rho 1$ cDNA sequence):

5'-p2 primer: 5'-CGGC<u>GGATCC</u>ACAGGCTACTGGAAAGCAGC-3' BamHI

 $\rho 1 \rho 2$ -A hybrid primer:

5'-CTGGCCTTCTACAGCAGCAGCAGC**AGGCTGGTACAACCGTCTC**-3' 3'-p1 primer: 5'-**CTGCAGCTTTCCAAAGCT**-3'

After PCR amplification, both $\rho 1\rho 2$ -A and $\rho 2\rho 1$ -A chimeras were subcloned into the PCRII vector (Invitrogen) and completely sequenced. PCR errors were corrected by site-directed mutagenesis techniques described below.

In Vitro Mutagenesis. Oligonucleotide-mediated sitedirected mutagenesis was performed on uracil-containing single-stranded DNA by the method of Kunkel (22). Mutated clones were identified by hybridization with the radiolabeled oligonucleotides. All positive mutants were verified by dideoxynucleotide sequencing of the mutated region. Five mutagenic oligonucleotides were used, with nucleotides producing the mutation underlined:

P309A 5'-GATACCTAAGGCGACTCTGGC-3' P309G 5'-GTGATACCTAAGCCGACTCTGGCAGG-3' P309S 5'-GAGATACCTAAGGAGACTCTG-3' P309V 5'-TGTGATACCTAAGACGACTCTGGCAGG-3' S319T 5'-GATGGTGGTCATGGTTAGCAC-3'

Oocyte Expression and Electrophysiology. Oocyte injection and a two-electrode voltage-clamping technique were performed as described (18). RNA injection was used for $\rho 1$, $\rho 1$ mutants, and the $\rho 1 \rho 2$ -A chimera. The amount of RNA injected in a 50-nl vol was as follows: ρ 1, 5 ng; ρ 1 ρ 2-A, 5 ng; P309G, 5 ng; P309S, 10 ng; P309A, 25 ng. Nuclear injection of the ρ^2 cDNA and $\rho^2\rho^{1-A}$ chimeric cDNA was performed to achieve efficient expression of ρ^2 and $\rho^2\rho^{1-A}$. The 3-kb ρ^2 cDNA fragment was subcloned into the Sca I site of the nuclear injection vector pMT3 and the $\rho 2\rho 1$ -A chimera was subcloned into the EcoRI site of the pMT3 vector (23). For studies of picrotoxin sensitivity, the amount of complementary RNA or pMT3 cDNA injected was varied to obtain similar whole-cell currents (150-300 nA). For nuclear injection, 1.5 ng of pMT3-p2 or 1.5 ng of pMT3-p2p1-A was injected. Two to 4 days after RNA or cDNA injection, oocytes were recorded in ND96 solution under voltage-clamp measurements with an Axoclamp-2A amplifier (Axon Instrument, Burlingame, CA). ND96 contains 96 mM NaCl, 2 mM KCl, 2 mM CaCl₂, 1 mM MgCl₂, and 5 mM Hepes·NaOH (pH 7.4). GABA and picrotoxin were purchased from Sigma. GABA was dissolved in bathing solution (ND96). Picrotoxin was dissolved in dimethyl sulfoxide (DMSO) as 5 M stock solution and was diluted in ND96 into test concentrations. The final concentration of DMSO was kept below 0.1%.

Curve Fit and Statistics. The dose-dependent inhibition data were best-fitted by inverse Hill equation using the N-fit program (University of Texas, Galveston). The equation was $I(x) = I_{\text{max}}/1 + (x/K_d)^n$, where x represents the drug concentration, K_d is the drug concentration for eliciting half-maximal response, and n is the Hill coefficient. Student's t test was used to determine the statistical significant difference between two sample groups. In this study, independent sampling and comparison were used and P < 0.01 was set as a significant level.

RESULTS

The Difference in Picrotoxin Sensitivity Between $\rho 1$ and $\rho 2$ Receptors Is Conferred by a Single Residue in the Second Membrane-Spanning Domain (M2). GABA $\rho 2$ receptors were consistently more sensitive to picrotoxin than GABA $\rho 1$ receptors in oocytes expressing either $\rho 1$ or $\rho 2$ receptors with similar whole-cell GABA-gated currents (range, 150–200 nA) (Fig. 1.A). Picrotoxin did not alter the linearity of the wholecell current, indicating that the blockage is voltage independent (data not shown). The IC₅₀ (concentration of picrotoxin that inhibits 50% of maximal currents) derived from these curves differed significantly between $\rho 1$ and $\rho 2$ receptors (P < 0.01) (Table 1). The minimal picrotoxin concentration that inhibited $\rho 1$ and $\rho 2$ currents was 1 μ M. Total inhibition of $\rho 2$ currents was observed when picrotoxin concentration reached 30 μ M, whereas 250 μ M was required for complete inhibition of $\rho 1$ receptors. The Hill coefficients of picrotoxin inhibition for $\rho 2$ and $\rho 1$ receptors were also significantly different (P < 0.01) (Table 1).

Two chimeras were generated to determine regions conferring the different picrotoxin sensitivity to ρ subunits. The ρ 1- ρ 2-A chimera included the cDNA sequence of the N-terminal extracellular domain of the ρ 1 subunit fused with the Cterminal half of ρ^2 containing putative membrane-spanning domains M1–M4 (Fig. 1B). A reciprocal chimera, $\rho 2\rho 1$ -A, was also created that contains the N-terminal domain of the ρ^2 subunit fused with the C-terminal half of the ρ 1 subunit. The $\rho 1 \rho 2$ -A chimera produced robust currents (200 nA to 1 μ A; n = 9) when expressed in oocytes similar to those observed with the ρ 1 subunit. However, receptors formed from this chimera were 10-fold more sensitive to picrotoxin than wild-type $\rho 1$ receptors with an IC₅₀ value and Hill coefficient close to that obtained from ρ^2 receptors (Fig. 1B). Nuclear expression of the $\rho 2\rho 1$ -A chimeric cDNA produced current levels (150–200 nA; n = 6) similar to those observed in ρ^2 cDNA-injected oocytes. However, unlike ρ^2 receptors, the $\rho^2\rho^{1-A}$ chimeric receptors were more resistant to picrotoxin blockage with an IC₅₀ value and Hill coefficient similar to that of $\rho 1$ receptors (Fig. 1B). These data indicate that the C-terminal half of ρ subunits containing M1-M4 determine picrotoxin sensitivity in ρ subunits.

Since prior studies had suggested that picrotoxin may block the ion conduction pore, we compared the amino acid sequence of the putative channel-lining M2 domains of $\rho 1$ and ρ^2 subunits. Two differences were noted: a proline at codon 309 of ρ 1, whereas ρ 2 has a serine in the analogous position, and a serine at codon 319 of ρ 1, while ρ 2 has a threenine. To determine whether either residue altered picrotoxin sensitivity, two mutant ρ 1 subunits were created. In one mutant, the proline residue at position 309 of ρ 1 was changed to serine (P309S) and, in the other, the serine residue at position 319 was substituted with threonine (S319T). Both mutants were expressed readily in oocytes with whole cell currents ranging from 200 nA to 1 μ A. Picrotoxin inhibition of currents generated by the S319T mutant was indistinguishable from $\rho 1$ (Table 1). However, the IC_{50} of P309S to picrotoxin is significantly different from $\rho 1$ (P < 0.01) but very similar to $\rho 2$ receptors (Table 1). Furthermore, the P309S and ρ 2 receptors have Hill coefficients for picrotoxin around 2, which is 2-fold larger than that of $\rho 1$ receptors (Table 1). Therefore, mutation of this single amino acid to that found in the ρ^2 subunit altered the picrotoxin interaction of $\rho 1$ to match that of $\rho 2$.

Picrotoxin Sensitivity of $\rho 1$ Receptors Is Increased by Substituting the Residue at Position 309 with Those Found in Picrotoxin-Sensitive Glycine α and GABA_A β and *Rdl* Receptor Subunits. To assess the relationship between the nature of the amino acid at position 309 of $\rho 1$ and picrotoxin sensitivity, mutants P309G, P309V, and P309A were constructed to correspond with residues found in the analogous position in glycine and GABA_A α , β , and wild-type *Rdl* subunits, respectively. (P309S corresponds with the serine residue found in GABA_A γ and δ subunits.) Two mutants, P309A and P309G, expressed GABA-gated currents that differed markedly in picrotoxin sensitivity (Table 1 and Fig. 2). For example, 5 μ M picrotoxin did not inhibit $\rho 1$ currents, whereas the same concentration of picrotoxin decreased P309G currents by $\approx 30\%$ and almost totally abolished currents generated by

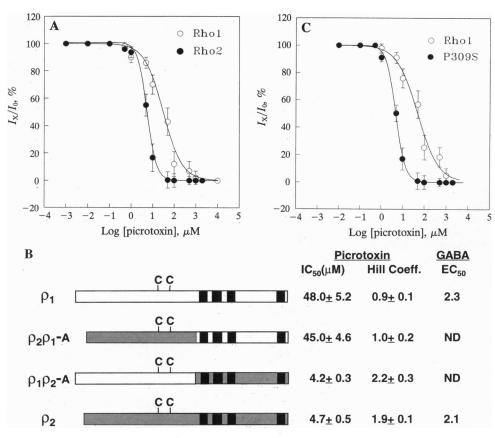


FIG. 1. Inhibitory effects of picrotoxin on $\rho 1$ and $\rho 2$ receptors. (A) Picrotoxin dose-dependent inhibition curve of $\rho 1$ and $\rho 2$ receptors. \bigcirc , $\rho 1$ (n = 3); \bullet , $\rho 2$ (n = 3). Data are expressed as I_X/I_0 vs. concentration of picrotoxin. I_0 , current elicited by 20 μ M GABA; I_X , current evoked by different concentrations of picrotoxin coapplied with 20 μ M GABA. Data are means \pm SE. (B) Comparison of picrotoxin sensitivity between $\rho 1 \rho 2$ -A chimera and $\rho 2\rho 1$ -A chimera. Structures of $\rho 1$ (open rectangle) and $\rho 2$ (shaded rectangle) and chimeric channels composed of $\rho 1$ and $\rho 2$ sequences are shown on the left, with the four putative transmembrane segments aligned. The responses of each construct to picrotoxin (IC₅₀ values and Hill coefficients) are shown as means \pm SE (n = 3). (C) Dose-dependent picrotoxin and GABA. Data are expressed as normalized picrotoxin response $(I_X/I_0) \pm$ SE vs. log scale of picrotoxin concentration. \bigcirc , $\rho 1$ (n = 3); \bullet , P309S mutant (n = 3). I_0 and I_X are the same as in A.

P309A, the most sensitive mutant. Unlike the P309S mutant, however, the Hill coefficients for picrotoxin inhibition of the P309G and P309A mutants were near unity, similar to wildtype $\rho 1$ (Table 1). Oocytes injected with the P309V mutant failed to express GABA-gated channels (n = 5). To determine whether a stable P309V mutant subunit was translated in oocytes, complementary RNAs synthesized from P309V and from $\rho 1$ were coinjected at ratios of 1:1 and 4:1. At the higher ratio, the expressed currents were 4-fold more sensitive to picrotoxin than $\rho 1$ alone (n = 5), suggesting that the P309V

Table 1. Comparisons of picrotoxin and GABA responses in $\rho 1$, $\rho 2$, and $\rho 1$ mutants

	Picrotoxin		
		Hill coefficient	GABA
	IC ₅₀ , μM		C ₅₀ , µM
GABA ρ1	48.0 ± 5.0	0.9 ± 0.1	2.3 ± 0.5
GABA p2	4.7 ± 0.5	1.9 ± 0.1	2.1 ± 0.3
P309S	4.8 ± 0.8	1.8 ± 0.2	6.7 ± 1.1
P309G	5.8 ± 1.2	0.9 ± 0.2	6.2 ± 1.5
P309A	0.1 ± 0.4	0.9 ± 0.3	7.2 ± 1.2
P309V	, <u> </u>		
S319T	47.2 ± 6.1	1.0 ± 0.1	
GABA p1			
+ P309V*	11.3 ± 4.0	0.8 ± 0.1	34.5 ± 5.4

Data are expressed as means \pm SE (n = 3). —, No current.

*Recordings after coinjection of 5 ng of ρ 1 and 20 ng of P309V per oocyte.

mutant, although nonfunctional as a homooligomer, could interact with $\rho 1$ subunits and form heterooligomeric receptors with increased picrotoxin sensitivity (Table 1).

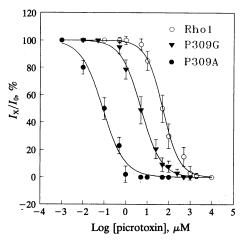


FIG. 2. Dose-dependent inhibition curves of picrotoxin generated by $\rho 1$ receptors and mutant $\rho 1$ receptors P309G and P309A. Responses to a wide concentration range of picrotoxin were collected from oocytes expressing homooligomeric channels generated from mutants P309G (\mathbf{v}) (n = 5), P309A (\mathbf{o}) (n = 5), and $\rho 1$ (\bigcirc) (n = 5). The current elicited by 20 μ M GABA without picrotoxin is the maximal current response (I_0). The current response after picrotoxin application is represented as I_X . Normalized current (I_X/I_0) vs. log scale of picrotoxin concentration is plotted \pm SE.

Mutation of Position 309 in p1 Influences the Agonist-Dependent (Competitive) Component of Picrotoxin Inhibition. The channel properties and responses of the residue 309 mutants to GABA were determined to assess whether the mutations had any effect on other receptor properties. Mutants P309S, P309G, and P309A have linear I-V relationships with a reversal potential close to the Cl⁻ equilibrium potential (data not shown). The three mutants had the same anion selectivity series as $\rho 1$ receptors; $Cl^- > Br^- > I^-$ when measuring the slopes of the cord conductance, and $Cl^- > Br^-$ > I⁻ when measuring shifts in reversal potential (n = 3 for each). The discrepancies in selectivity series measured by these two different methods are similar to previous reports of GABA-gated Cl⁻ currents from isolated neurons (24). These data suggest that intrinsic biophysical properties of the Clchannel formed by the $\rho 1$ mutants were not significantly altered. However, the agonist-dependent activation differs between $\rho 1$ and the residue 309 mutants. Regardless of the substituted amino acid, all mutants have significantly higher EC₅₀ values for GABA activation than $\rho 1$ (P < 0.01) (Table 1). These results prompted an analysis of the sensitivity of $\rho 1$ mutants to picrotoxin at submaximal concentrations of GABA. Picrotoxin inhibition curves for P309S and P309G at three different GABA concentrations are compared with $\rho 1$ in Fig. 3. ρ 1 receptors showed a 30-fold increase in the picrotoxin IC_{50} between 1 and 5 μ M GABA applications (18). In contrast, P309S and P309A had minor changes in picrotoxin IC₅₀ values between these two concentrations of GABA. For P309S, the IC₅₀ value for picrotoxin was 1.0 \pm 0.6 μ M at 1 μ M GABA (n = 5) and 3.1 \pm 0.8 μ M (n = 5) at 5 μ M GABA, and it reached $4.8 \pm 0.7 \ \mu M \ (n = 5)$ at a saturating level of GABA (20 μM). Similar minor changes in the IC₅₀ of picrotoxin between 1 μ M $[1.0 \pm 0.8 \ \mu M \ (n = 3)], 5 \ \mu M \ [2.5 \pm 0.7 \ \mu M \ (n = 3)], and 20$ $\mu M [5.4 \pm 0.5 \mu M (n = 3)]$ GABA concentrations were observed for the P309A mutant.

DISCUSSION

This report identifies a molecular basis for the picrotoxin sensitivity of GABA ρ 1 receptors. Substitution of proline-309 of $\rho 1$ with a serine (P309S), the corresponding residue in $\rho 2$, decreased the IC₅₀ and increased the Hill coefficient of this mutant to precisely match ρ^2 receptors. This suggested that the difference between $\rho 1$ and $\rho 2$ subunits in picrotoxin sensitivity is primarily determined by a single amino acid in the M2 domain. Two lines of evidence indicate that this single mutation is specific for picrotoxin interaction. First, several important properties of the mutant receptor (current-voltage relationship, Cl⁻-reversal potential, anion selectivity) were unaltered. Second, this mutation produced a gain in function (i.e., increased sensitivity to picrotoxin) rather than a loss of function. The distinction is critical because increased function provides evidence that the structural integrity of the domain that interacts with the compound has not only been preserved but has been altered specifically to enhance its interaction with the compound.

The nature of the amino acid at position 309 accounts for the differences in picrotoxin inhibition not only between $\rho 1$ and $\rho 2$ receptors but also among GABA $\rho 1$ receptors and GABA_A receptors. Alignment of the amino acid sequences of the M2 domain of mammalian GABA_A receptor subunits and ρ subunits reveals that the amino acid residue at this position is highly diverse. Our data demonstrate that substituting the proline residue of $\rho 1$ with valine, alanine, serine, or glycine, the amino acids found in GABA_A subunits and the glycine α subunit, increased the sensitivity toward picrotoxin. They also suggest a starting point to assess the differences in the picrotoxin blocking effect between mammalian retinal GABA_C receptors and those characterized in perch, salamander, and catfish retina (9, 10, 12, 19, 20, 25). Identification of the residues occupying position 309 in ρ subunits from these species helps

determine whether variation at this location accounts for the increased sensitivity of nonmammalian GABA_C receptors.

Residue 309 of $\rho 1$ corresponds to one of the amino acids in glycine receptors shown to affect picrotoxin sensitivity. Mutation of 13 residues in the putative channel-lining M2 domain

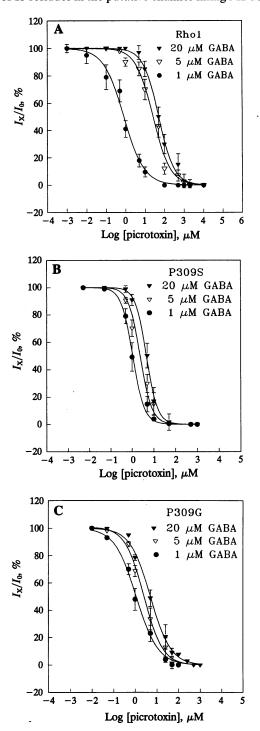


FIG. 3. Agonist-dependent picrotoxin inhibition curves of $\rho 1$ and $\rho 1$ mutants P309S and P309G. (A) Picrotoxin inhibition responses of the $\rho 1$ receptor observed at three GABA concentrations: 1, 5, and 20 μ M (n = 3 for each). Data are expressed as I_X/I_0 vs. concentration of picrotoxin. I_0 , current response to 1, 5, or 20 μ M GABA; I_X , current response to 1, 5, or 20 μ M GABA coapplied with a certain concentration of picrotoxin. (B) Picrotoxin inhibition curves for the P309S mutant at the same three concentrations of GABA as in A (n = 5 for each). (C) Picrotoxin inhibition curves for the P309G mutant at the same three concentrations of GABA as in A (n = 3 for each). Data are means \pm SE.

of the β glycine receptor subunits to those found in the α_3 subunit increased picrotoxin sensitivity of α_3/β receptors to levels observed in α_3 homooligomeric receptors (3). It also is the same residue that is altered in a GABA_A receptor subunit of *Drosophila Rdl*. This naturally occurring point mutation confers insecticide resistance to the *Rdl* (resistance to dieldrin) strain (4, 26). Since picrotoxin and the cyclodiene insecticides compete for the same binding site (27), this amino acid substitution identified a biologically important location for picrotoxin interaction with invertebrate GABA_A receptors. These observations are consistent with the concept that picrotoxin noncompetitively inhibits GABA_A, glycine, and GABA ρ 1 receptors by interacting with residues forming the ion pore.

One of the aims of this study was to identify residues facilitating competitive picrotoxin antagonism of ρ 1 receptors. Our results suggest that the nature of the amino acid at position 309 plays a critical role in this process. The presence of a proline residue confers a competitive component to the GABA $\rho 1$ receptor. Substitution with residues found in the analogous location in GABAA receptors eliminates this component. The residue 309 location is distinct from the putative agonist binding region of $\rho 1$ receptors (28). Thus, picrotoxin may inhibit GABA activation of $\rho 1$ receptors by an allosteric mechanism in addition to blockade of the open channel. This could explain why picrotoxin acts noncompetitively on $GABA_A$ receptors but displays mixed antagonism of GABA ρ and GABA_C receptors. Characterization of the molecular composition of retinal GABA_C receptors will help confirm this hypothesis.

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