

Acute Functional Tolerance to Ethanol Mediated by Protein Kinase C ϵ

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A low level of response to ethanol is associated with increased risk of alcoholism. A major determinant of the level of response is the capacity to develop acute functional tolerance (AFT) to ethanol during a single drinking session. Mice lacking protein kinase C epsilon (PKC ϵ) show increased signs of ethanol intoxication and reduced ethanol self-administration. Here, we report that AFT to the motor-impairing effects of ethanol is reduced in PKC ϵ (–/–) mice when compared with wild-type littermates. In wild-type mice, *in vivo* ethanol exposure produced AFT that was accompanied by increased phosphorylation of PKC ϵ and resistance of GABA $_A$ receptors to ethanol. In contrast, in PKC ϵ (–/–) mice, GABA $_A$ receptor sensitivity to ethanol was unaltered by acute *in vivo* ethanol exposure. Both PKC ϵ (–/–) and PKC ϵ (+/+) mice developed robust chronic tolerance to ethanol, but the presence of chronic tolerance did not change ethanol preference drinking. These findings suggest that ethanol activates a PKC ϵ signaling pathway that contributes to GABA $_A$ receptor resistance to ethanol and to AFT. AFT can be genetically dissociated from chronic tolerance, which is not regulated by PKC ϵ and does not alter PKC ϵ modulation of ethanol preference.

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INTRODUCTION

Genetic factors account for 50–60% of the risk of developing an alcohol-use disorder (Prescott and Kendler, 1999). One behavioral characteristic related to this risk is a low level of response to an acute alcohol challenge (Schuckit, 1998), a trait that carries a similar estimated heritability of 40–60% (Schuckit and Smith, 1996; Heath *et al.*, 1999). Linkage studies have identified regions on chromosomes 10, 11, 13, and 20 with LOD scores ≥ 3.0 that may contain genes that contribute to the level of response to alcohol (Schuckit *et al.*, 2004). Few genes have been identified, although recently the long allele of the SLC6A4 gene, which encodes the serotonin transporter, and the Ser 385 variant of the GABRA6 gene, which encodes the $\alpha 6$ subunit of GABA $_A$ receptors, have been associated with a low level of response to alcohol in humans (Hu *et al.*, 2005). The Collaborative Study on the Genetics of Alcoholism has identified the GABRA2 gene

located on chromosome 4 encoding the $\alpha 2$ -subunit of the GABA $_A$ receptor, the GABRB1 gene on chromosome 4 coding for the $\beta 1$ subunit of the GABA $_A$ receptor, and the CHRM2 gene on chromosome 7, which encodes the type 2 muscarinic receptor as susceptibility genes for alcoholism (Dick *et al.*, 2005), although it is not known if they contribute to the level of response to alcohol. In addition, animal studies have identified several other genes that alter acute responses to alcohol in rodents as candidates for the regulation of the level of response in humans (Schuckit *et al.*, 2004).

The level of response to alcohol involves two components: initial sensitivity and acute functional (behavioral) tolerance (AFT). Initial sensitivity is measured as the level of intoxication that appears within minutes after ethanol administration, when blood ethanol levels are rising. AFT is characterized by a greater level of intoxication during the rising phase of the blood ethanol curve than at equivalent concentrations on the falling phase. AFT develops during a single session of ethanol exposure and is not owing to changes in ethanol clearance. It is distinguished from rapid tolerance, which is measured 8–72 h after an episode of ethanol exposure when ethanol is completely metabolized, and from chronic tolerance, which is the progressive resistance to ethanol intoxication following repeated sessions of ethanol exposure (Kalant *et al.*, 1971; Kalant,

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1996). As the most consistent differences in response to an alcohol challenge in human subjects are detected 60 min or more after ethanol administration when blood alcohol levels are declining, it is likely that they reflect differences in AFT (Schuckit, 1984a,b; Schuckit *et al*, 1987, 1988; Newlin and Thomson, 1990).

We previously found that mice deficient in protein kinase C epsilon (PKC ϵ) show an increased duration of the ethanol-induced loss of the righting reflex (LORR), a measure of hypnotic response, when compared with wild-type littermates (Hodge *et al*, 1999). This is associated with a normal response to the direct GABA $_A$ agonist muscimol, but an increased enhancement of GABA $_A$ receptor function by ethanol (Hodge *et al*, 1999; Proctor *et al*, 2003). As GABA $_A$ receptor modulation and behavioral responses to benzodiazepines (Hodge *et al*, 1999) and neurosteroids (Hodge *et al*, 2001), which also act at GABA $_A$ receptors, are greater in PKC ϵ ($-/-$) mice, increased modulation of GABA $_A$ receptor function by ethanol is likely to explain the enhanced level of response in these mice.

In this study, we examined PKC ϵ ($-/-$) mice to determine if their increased level of response to ethanol is owing to increased initial sensitivity or decreased AFT and is related to altered regulation of GABA $_A$ receptors by ethanol. Our findings provide evidence for ethanol-induced activation of a PKC ϵ -dependent signaling pathway that contributes to acute tolerance of GABA $_A$ receptors to ethanol and to AFT but not to chronic tolerance.

MATERIALS AND METHODS

Animal Care

A null mutation was generated in the mouse *Prkce* gene by homologous recombination in embryonic stem cells, as described (Khasar *et al*, 1999). Male and female PKC ϵ ($+/-$) mice maintained on inbred 129S4 and backcrossed (>10 generations) C57BL6/J backgrounds were mated to produce PKC ϵ ($+/-$) C57BL6/J \times 129S4 F1 hybrid mice for breeding. These mice were intercrossed to generate F2 hybrid PKC ϵ ($+/+$) and PKC ϵ ($-/-$) littermates for experiments. Mice were genotyped by PCR of tail biopsies (5–10 mm fragment) obtained from nonanesthetized, 10-day-old pups. All animals were male and between 2 and 6 months old at the time of experimentation. We observed no age-related differences in any of the behavioral tests performed. All procedures were conducted in accordance with institutional IACUC policies and the Society for Neuroscience's Policy on the Use of Animals in Neuroscience Research. For each behavioral test, a different group of mice was used.

LORR

Initial sensitivity to ethanol-induced loss of righting was measured using the 'up and down' method (Dixon, 1965; Findlay *et al*, 2002) to calculate an ED $_{50}$ for ethanol-induced LORR. A mouse was given an intraperitoneal (i.p.) injection of 10% (v/v) ethanol in saline, and 5 min later, tested for LORR greater than 1 min. If this first mouse lost the righting reflex, then the next mouse was given a lower ethanol dose. If the first mouse did not lose the righting reflex within

5 min after injection, then the next mouse was given a higher dose. The log dose interval was 0.0138, corresponding to approximately a 0.1 g/kg difference between doses. The ED $_{50}$ value was determined as described (Findlay *et al*, 2002), with 95% confidence intervals (CIs) calculated by the following equation: 95% CI = dosing increment \times square root of $(2/n) \times 1.96$, where n is the the last six trials and 1.96 reflects the 0.05 α level (Dixon, 1965). When 95% CIs do not overlap, the ED $_{50}$ values for each group of animals are considered significantly different.

LORR duration was measured by administering ethanol (20% (v/v) in 0.9% saline) i.p. in different volumes to obtain two test doses (3.2 or 4.0 g/kg). In mice, these concentrations yield LORR durations ranging from 30 to 120 min, and cover the range of doses between the minimum necessary to produce the response and the maximum that will not cause death. After injection, mice were placed on their backs. LORR was defined as the animal being unable to right itself three times within a 30-s period. The mouse was considered to have recovered the righting reflex when it could right itself three times within a 1 min time frame. A blood sample was taken from the tail vein at the time of recovery for the determination of blood ethanol concentration (BEC) in mice that had received 4.0 g/kg ethanol. Mice were tested at both concentrations of ethanol using a random order, balanced design, with at least 1 week elapsing between test sessions.

Rotarod Ataxia

On the day preceding testing, ethanol-naïve mice were trained to remain on a rotarod (model 7650; Ugo Basile, Italy) rotating at 14 r.p.m. for a 3-min period. The following day, mice were tested for their ability to stay on the rotarod for 3 min. Mice were then injected with 2.0 g/kg ethanol i.p. and placed back on the apparatus at 15, 30, 45, 60, 75, and 90 min postinjection. The time to fall from the rotarod was recorded for each trial. For each mouse, blood samples were taken 5 min postinjection and at recovery from ataxia, and were analyzed for BEC.

To measure initial sensitivity to ethanol-induced ataxia, the 'up and down' method was used to calculate the estimated ED $_{50}$ for ethanol. An i.p. injection of 10% ethanol (v/v) in 0.9% saline was administered and the mouse was immediately placed on the rotarod for 5 min. If the mouse fell off the rotarod twice during the test, it was judged to be ataxic. If the first mouse showed ataxia, then the next mouse was given a lower ethanol dose. If the first mouse did not show ataxia within 5 min after injection, then the next mouse was given a higher dose. The log dose interval was 0.0276, corresponding to approximately a 0.2 g/kg difference between doses. ED $_{50}$ values and CIs were calculated as described above for ethanol-induced LORR.

Stationary Dowel Test

For this test, we used a modification of the procedure described by Wu *et al* (2001). A horizontal wooden dowel (1.27 cm diameter, 30 cm long) was suspended between two Plexiglas walls 50 cm above a cushioned surface. Mice were trained to balance on the dowel for three 1-min periods at 5-min increments. The mice were considered trained when

they were able to remain on the rod for more than 1 min. For AFT experiments, each mouse was then given an i.p. injection of 1.5 g/kg ethanol (20% (w/v) in saline) and placed on the dowel. After the mouse fell from the dowel, it was retested every 5 min until it was able to remain on the dowel for at least 1 min, at which time a blood sample was taken (BEC1) by tail vein puncture. Following sample collection, the mouse was immediately given a booster injection of 1.5 g/kg ethanol and placed back on the dowel. The time at which the second period of ataxia began was recorded when the animal again fell off the rod. The mouse was retested every 5 min until it was again able to remain on the rod for at least 1 min, at which point a final blood sample (BEC2) was drawn. Some animals were bled 5 min after the second injection to measure BEC, but were not tested further. Blood samples were centrifuged at 10 000g and serum blood alcohol concentrations were determined using an Analox AM-1 analyzer (Analox Instruments, Lunenburg, MA). AFT was calculated as the difference between BEC2 and BEC1. The rate at which AFT developed was calculated as (BEC2–BEC1)/duration of ataxia following the booster injection. The threshold for sensitivity to ethanol-induced ataxia was calculated by extrapolating the line determined by the BEC2 and BEC1 values to zero time. The rationale for this calculation is based on the evidence indicating that AFT develops linearly with the time of ethanol exposure up to a maximum (Radlow, 1994; Wu et al, 2001).

Ethanol Preference

Baseline fluid consumption was monitored by allowing mice access to two bottles containing water for 4 days. After baseline levels of fluid intake were determined, mice were given free access to two drinking bottles, one containing an ethanol solution and the other containing water. Bottles and mice were weighed every 2 days to monitor consumption. The side of the cage in which the ethanol bottle was presented (left vs right) was switched every 2 days. The concentration of ethanol (3, 6, or 10%) was increased every 3–5 days. Consumption of 10% ethanol was assessed for 6 days, after which animals were treated twice daily with i.p. injections of 2.5 g/kg ethanol for 6 days to establish chronic tolerance to ethanol. Starting on day 7, they underwent a second session of two-bottle choice drinking with 10% ethanol for 6 days.

Ethanol Clearance

Ethanol clearance was determined in ethanol-naïve mice by injecting them with 3.6 g/kg of ethanol i.p. and removing 20 μ l blood samples from the tail vein at 30, 60, and 120 min postinjection. The following day, chronic treatment was started with animals injected twice daily with 2.5 g/kg ethanol i.p. On day 7, mice were retested for ethanol clearance.

Preparation of Cerebellar Microsacs

Duplicate pairs of animals of each genotype were injected with 2.0 or 4.0 g/kg of ethanol i.p. or the equivalent volume of normal saline and 45 min later cerebella were dissected

on ice. Cerebella from each pair were pooled to constitute one sample and microsacs were prepared from each sample using a modification of the method described by Harris and Allan (1985). Samples were homogenized by hand (seven to 10 strokes) using a Teflon-glass homogenizer containing 7 ml of ice-cold assay buffer (145 mM NaCl, 5 mM KCl, 1 mM MgCl₂, 10 mM glucose, 1 mM CaCl₂, 10 mM HEPES (pH 7.5), using Tris base) plus a Complete Protease Inhibitor Cocktail Tablet (Roche Applied Science, Indianapolis, IN). The homogenate was centrifuged at 950g for 5 min, using a Marathon 3200R rotor (Fisher Scientific, Fairlawn, NJ). The pellet was resuspended in assay buffer containing 0.2% BSA to yield a suspension of 6 mg/ml protein. It took approximately 15 min to generate microsacs after killing the mice; thus, the ³⁶Cl[–] uptake assay (see below) was started 1 h after *in vivo* injection of ethanol or saline.

³⁶Cl[–] Uptake

Microsacs (200 μ l) were incubated at 37°C for 5 min. ³⁶Cl[–] uptake was started by adding a solution containing ³⁶Cl[–] (0.2 μ Ci/ml of assay buffer), 1 μ M muscimol, or 1 μ M muscimol plus 20 mM ethanol. After 5 s, the reaction was stopped by adding 4 ml of ice-cold assay buffer containing 100 μ M picrotoxin followed by rapid vacuum filtration over 2.5 cm GF/C Whatman filters (Whatman, Kent, UK). Filters were washed with an additional 12 ml of cold assay buffer and counted in ScintiVerse scintillation fluid (Fisher Scientific, Fairlawn, NJ) using a scintillation counter. Muscimol-stimulated uptake was calculated as the difference in uptake measured in the presence and absence of muscimol. Ethanol-enhanced ³⁶Cl[–] uptake was expressed as a percentage of uptake measured in the presence of muscimol alone.

Western Analysis of PKC ϵ S729 Phosphorylation

Pairs of mice were injected with ethanol (1.5 or 4 g/kg) or saline i.p. and were killed by decapitation at indicated times. Brains were rapidly removed, and frontal cortex, cerebellum, and striatum were dissected. Brain tissues were homogenized in lysis buffer containing 20 mM Tris, 10 mM EGTA, 2 mM EDTA, 250 mM sucrose, 1 mM PMSF, 1 \times Complete Protease Inhibitor Cocktail (Roche Applied Science, Indianapolis, IN), and a 1:100 dilution of serine/threonine phosphatase inhibitor cocktail 1 (Sigma-Aldrich, St Louis, MO) on ice. The homogenates were centrifuged at 10 000g for 10 min at 4°C and the supernatants were collected. Concentrated Laemmli sample buffer was added to yield a final solution containing 62.5 mM Tris-HCl (pH 6.8), 2% SDS, 5% β -mercaptoethanol, 10% glycerol, and 0.002% bromophenol blue. Samples were boiled at 90°C for 7 min and then passed through a 28-G needle. Equal amounts of protein (20 μ g/lane) were loaded onto 6% Tris-glycine SDS-polyacrylamide gels. In pilot experiments, we determined that 20 μ g of protein was optimal for detecting at least a two-fold increase in signal while remaining within the linear range of detection for each brain region that we tested. The separated proteins were then electrophoretically transferred to a nitrocellulose membrane (Hybond™-C Extra, Amersham Biosciences, Buckinghamshire, UK). Membranes

were blocked by incubation with 5% nonfat dry milk in 0.01 M PBS containing 0.1% Tween 20 for 1 h at 25°C. The membranes were then incubated with anti-phospho-PKC ϵ (Ser729) antibody (1:8000 dilution; Santa Cruz Biotechnology, Santa Cruz, CA) or anti-PKC ϵ (Choi *et al.*, 2002; 1:80 000 dilution), and anti-actin antibody (1:20 000 dilution; Sigma) in blocking solution overnight at 4°C. After being washed with 0.01 M PBS containing 0.1% Tween 20 (PBS-T) three times for 5 min, the membranes were incubated with goat anti-rabbit IgG-peroxidase-conjugated antibody (1:1000 dilution; Chemicon, Temecula, CA) in blocking solution for 1.5 h at 25°C. Antibody concentrations were based on the manufacturer's recommendations or were determined by pilot experiments conducted to identify concentrations that produced the highest specific immunostaining with the least nonspecific signal. We also confirmed the specificity of our anti-PKC ϵ antibody and the commercial anti-phospho-PKC ϵ antibody by Western blotting tissue from PKC ϵ (–/–) mice and no protein bands were observed (not shown).

After incubation in secondary antibody, the membranes were washed three times for 5 min in PBS-T. Immuno-reactive bands were visualized by chemiluminescence (Pierce, Rockford, IL) and recorded using X-ray film. The optical densities of the immunoreactive bands were quantified using a flat bed scanner and the program Image J (<http://rsb.info.nih.gov/ij>). Values for phospho-PKC ϵ and PKC ϵ immunoreactivity were normalized to actin immunoreactivity and the normalized values for each ethanol- and saline-treated pair were divided to yield an ethanol/saline ratio for phospho-PKC ϵ and total PKC ϵ . The experiment was repeated at least four times for each time point and each dose of ethanol.

RESULTS

Reduced AFT to Ethanol in PKC ϵ (–/–) Mice

PKC ϵ (–/–) mice show an increased duration of ethanol-induced LORR (Hodge *et al.*, 1999). To determine if this is related to increased initial sensitivity or altered AFT to ethanol, we compared LORR duration and the BEC upon recovery of the righting reflex with the ED₅₀ for ethanol-

induced loss of righting. As expected, PKC ϵ (–/–) mice showed a three-fold greater LORR duration when compared with PKC ϵ (+/+) littermates (Figure 1a) and the BEC at recovery from 4.0 g/kg was 16% lower in PKC ϵ (–/–) mice compared to PKC ϵ (+/+) mice (Figure 1b). In addition, the ED₅₀ for inducing loss of righting was decreased by about 8% in PKC ϵ (–/–) mice when compared with PKC ϵ (+/+) mice (Figure 1c). This indicates that PKC ϵ modulates initial sensitivity, and possibly AFT, to the hypnotic effect of 4 g/kg ethanol.

To investigate how PKC ϵ modulates the level of response to lower ethanol concentrations, we measured rotarod ataxia induced by 2.0 g/kg ethanol. At 5 min after the injection, the latencies to fall were similar ($p=0.29$) in PKC ϵ (+/+) mice (7 ± 2 s) and PKC ϵ (–/–) mice (4 ± 1 s) (Figure 2a). The BECs were also similar ($p=0.18$) in wild-type (247 ± 8 mg/dl; $n=14$) and PKC ϵ (–/–) mice (231 ± 8 mg/dl; $n=13$) at that time. At 15, 30, 45, 60, and 75 min postinjection, PKC ϵ (–/–) mice were less able to remain on the rotarod when compared with PKC ϵ (+/+) mice (Figure 2a). The BEC measured upon recovery, when mice were again able to remain on the rod for 3 min, was also significantly lower in PKC ϵ (–/–) mice compared with PKC ϵ (+/+) mice (Figure 2b). These findings suggested that the initial sensitivity to ethanol was similar, but that recovery, and therefore development of AFT, was more rapid in the wild-type animals. However, because the initial fall latencies measured 5 min after ethanol injection were very short (less than 10 s), we may have missed detecting a difference in initial sensitivity to 2.0 g/kg ethanol by this assay. Therefore, we also measured the ED₅₀ for inducing rotarod ataxia, and found it to be similar in PKC ϵ (+/+) and PKC ϵ (–/–) mice (Figure 2c). Taken together, these data suggest that PKC ϵ plays a greater role in AFT than in initial sensitivity to the ataxic effect of 2.0 g/kg ethanol.

The assays of LORR and rotarod ataxia provide only indirect information about AFT. AFT to ethanol-induced impairment of balance can be directly measured using a stationary dowel test (Erwin *et al.*, 2000; Wu *et al.*, 2001). To perform this test, we followed the method used by Erwin *et al.* (2000), which compares the response to two sequential doses of ethanol; however, we used lower ethanol doses

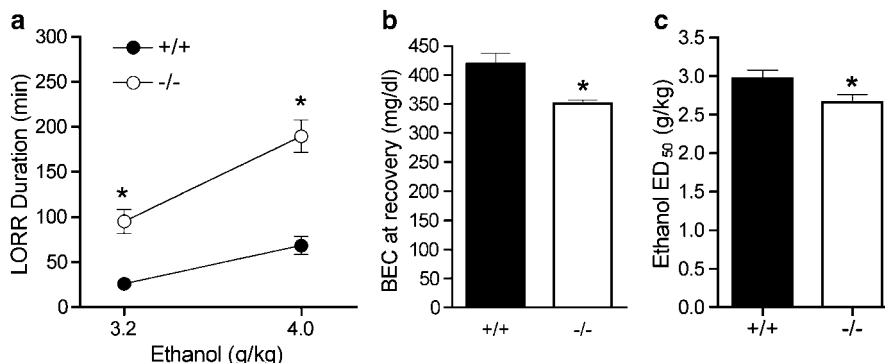


Figure 1 Ethanol-induced LORR. (a) LORR duration was increased in PKC ϵ (–/–) mice. Data are mean \pm SE values from seven to 13 mice per genotype. Two-way ANOVA showed main effects of genotype ($F(1,32) = 67.64$; $p < 0.001$) and dose ($F(1,32) = 35.13$; $p < 0.001$) with an interaction between these factors ($F(1,32) = 5.036$; $p = 0.0319$). * $p < 0.001$ compared with PKC ϵ (+/+) mice by *post hoc* Bonferroni tests. (b) The BEC upon regaining the righting reflex was higher in wild-type mice ($n = 7$) when compared with PKC ϵ (–/–) ($n = 8$) mice ($p < 0.003$, two-tailed *t*-test). (c) The ethanol ED₅₀ to induce loss of righting was lower in PKC ϵ (–/–) mice. Data are mean values \pm 95% CIs from six mice per genotype.

(1.5 g/kg) as two sequential higher doses of 2.0 g/kg produced loss of righting in some animals after the second dose. This method generates three measures: the magnitude of AFT, the rate at which AFT develops, and the level of initial sensitivity (see Materials and methods). We found that the magnitude of AFT and the rate at which AFT

developed were lower in PKC ϵ ($-/-$) mice when compared with PKC ϵ ($+/+$) littermates (Figure 3a and b). In contrast, the initial sensitivity to ethanol was similar between the genotypes (Figure 3c).

Chronic Tolerance to Ethanol

As PKC ϵ ($-/-$) mice showed decreased AFT, we investigated whether they also show reduced chronic tolerance to ethanol. Chronic tolerance is associated with accelerated development and increased magnitude of within-session tolerance in the stationary dowel test (Wu *et al*, 2001). Therefore, to measure chronic tolerance by this method, we treated both PKC ϵ ($+/+$) and PKC ϵ ($-/-$) mice with twice daily injections of 2.5 g/kg ethanol for 6 days and retested them on the stationary dowel. Following chronic ethanol exposure, the magnitude of within-session tolerance and the rate at which it developed increased significantly in both PKC ϵ ($+/+$) and PKC ϵ ($-/-$) mice to the extent that there was no longer a difference between the genotypes in these measures (Figure 3a and b). The change in within-session tolerance was greater in PKC ϵ ($-/-$) mice, which showed a 3.9-fold increase in AFT and a 6.8-fold increase in AFT rate, as compared with PKC ϵ ($+/+$) mice, which showed only a 1.4-fold increase in AFT and a 2.4-fold increase in AFT rate. Chronic ethanol exposure had no effect on initial sensitivity in either genotype (Figure 3c).

To determine if chronic tolerance to higher doses of ethanol also develops in PKC ϵ ($-/-$) mice, we measured the ethanol-induced LORR in mice treated with twice-daily injections of 2.5 g/kg ethanol for 6 days. Following chronic ethanol exposure, PKC ϵ ($+/+$) and PKC ϵ ($-/-$) mice both showed a decrease in LORR duration in response to acute administration of 3.6 g/kg ethanol, and there was no longer a difference in LORR duration between the genotypes (Figure 4a). The percent decrease in LORR duration that occurred with tolerance development was greater in PKC ϵ ($-/-$) than in PKC ϵ ($+/+$) mice (Figure 4b). These results indicate that absence of PKC ϵ does not impair the development of chronic tolerance to ethanol and may instead enhance it.

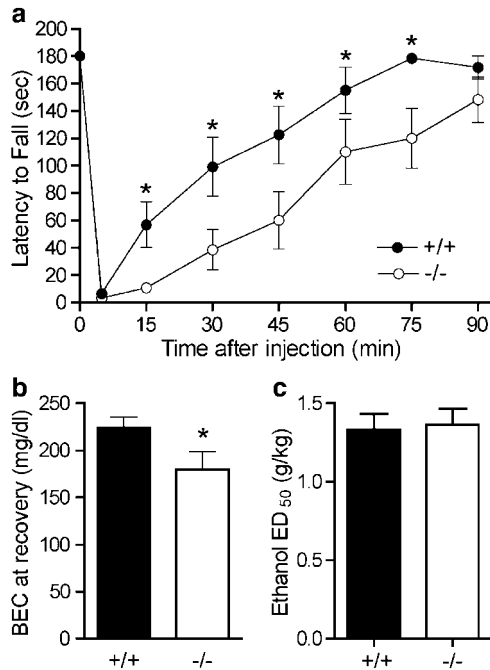


Figure 2 (a) PKC ϵ ($-/-$) mice ($n = 12$) were less able to remain on the rotarod than PKC ϵ ($+/+$) mice ($n = 13$) for 15–75 min after receiving 2.0 g/kg ethanol. Two-way, repeated measures ANOVA showed main effects of genotype ($F(1,154) = 7.90$; $p = 0.01$) and time ($F(7,154) = 52.80$; $p < 0.001$), with an interaction between these factors ($F(7,154) = 2.128$; $p = 0.044$). * $p < 0.05$ compared with PKC ϵ ($+/+$) mice at the same time by *post hoc* Newman–Keuls tests. (b) The BEC at recovery from rotarod ataxia was lower for PKC ϵ ($-/-$) mice ($n = 7$) compared with PKC ϵ ($+/+$) mice ($n = 9$). * $p < 0.05$ by two-tailed *t*-tests. (c) The ED₅₀ for ethanol-induced ataxia was similar for PKC ϵ ($-/-$) and PKC ϵ ($+/+$) mice. Data are mean values \pm 95% CIs from six mice per genotype.

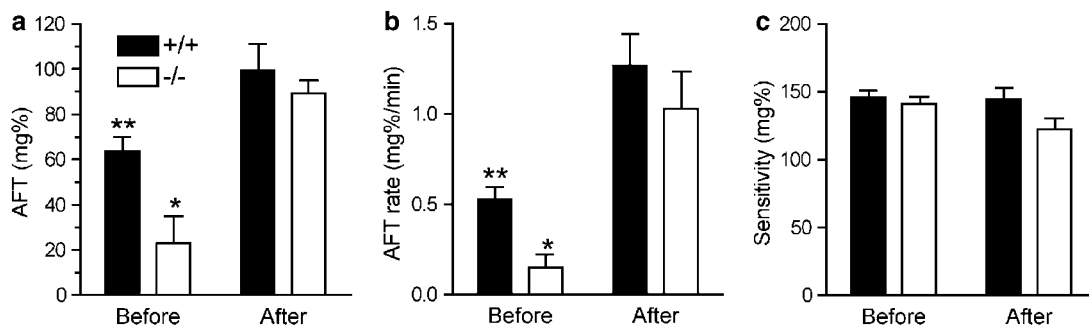


Figure 3 Ethanol-induced loss of balance on the stationary dowel before and after chronic ethanol exposure. AFT magnitude (a) and rate (b) were initially greater in wild-type mice compared with PKC ϵ ($-/-$) mice ('Before'), but chronic ethanol exposure increased AFT magnitude and AFT rate in both genotypes so that there was no longer a difference between the genotypes ('After'). Two-way ANOVA for AFT magnitude showed main effects of genotype ($F(1,57) = 7.56$; $p = 0.008$) and treatment ($F(1,57) = 30.08$; $p < 0.0001$). For AFT rate, there were also main effects of genotype ($F(1,51) = 4.92$; $p < 0.031$) and treatment ($F(1,51) = 34.04$; $p < 0.0001$). * $p < 0.05$ compared with PKC ϵ ($+/+$) mice before and with PKC ϵ ($-/-$) mice after chronic ethanol exposure; ** $p < 0.05$ compared with PKC ϵ ($+/+$) mice after chronic ethanol exposure (Bonferroni tests). (c) There was no change in calculated initial sensitivity to ethanol-induced ataxia in either genotype before or after chronic ethanol exposure. Data are mean \pm SEM values from 11 to 18 mice per genotype for each treatment condition.

Acute Tolerance to Ethanol Enhancement of GABA_A Receptor Function

PKC ϵ is abundant in the cerebellum (Choi *et al*, 2002) and ethanol's actions on cerebellar neurons appear to contribute to hypnotic and ataxic effects of ethanol (Palmer *et al*, 1987; Clark and Dar, 1988). A single exposure to ethanol *in vivo* reduces ethanol enhancement of GABA_A receptor function measured *in vitro* in cerebellar microsacs 1 h later; this effect has been proposed as a mechanism underlying AFT to ethanol (Allan and Harris, 1987). As GABA_A receptor modulation by ethanol is increased in PKC ϵ (–/–) mice (Hodge *et al*, 1999; Choi *et al*, 2002), we examined whether cerebellar GABA_A receptors from PKC ϵ (–/–) mice show acute tolerance to ethanol. We treated mice with 2.0 or

4.0 g/kg ethanol i.p. or an equivalent volume of saline and 45 min later isolated cerebellar microsacs to examine muscimol-stimulated $^{36}\text{Cl}^-$ uptake. As shown previously (Hodge *et al*, 1999), uptake stimulated by the direct GABA_A receptor agonist muscimol was similar in microsacs from PKC ϵ (–/–) and PKC ϵ (+/+) mice treated with saline or ethanol *in vivo* (Figure 5a). Also as shown previously (Hodge *et al*, 1999), muscimol-stimulated uptake was increased by the addition of 20 mM ethanol *in vitro* to a greater extent in microsacs from saline-injected PKC ϵ (–/–) mice when compared with microsacs from saline-injected PKC ϵ (+/+) mice (Figure 5b and c). *In vivo* treatment with 2.0 (Figure 5b) or 4.0 g/kg (Figure 5c) ethanol reduced the ability of *in vitro* 20 mM ethanol to enhance muscimol-stimulated uptake in PKC ϵ (+/+) microsacs but had no effect on uptake in microsacs from PKC ϵ (–/–) mice. These results indicate that PKC ϵ is required for acute tolerance of cerebellar GABA_A receptors to ethanol.

Ethanol Treatment Increases PKC ϵ S729 Phosphorylation in the Cerebellum

We next considered whether ethanol regulates PKC ϵ . As there are no specific substrates of PKC ϵ that have been identified, it is not yet possible to assay PKC ϵ activity in intact cells or tissues. PKC ϵ , like other PKC isozymes, undergoes a series of post-translational phosphorylation events that result in a mature, phosphorylated enzyme capable of being fully activated by lipid second messengers (Newton, 2003) and these events can be assayed using phosphorylation-site-specific anti-PKC ϵ antibodies (Parekh *et al*, 1999; Takahashi *et al*, 2000; Cenni *et al*, 2002). Unlike conventional PKC α , β , or γ isozymes, which are constitutively phosphorylated, there is a pool of unphosphorylated and inactive PKC ϵ that is stable within cells and can be recruited into cell signaling pathways by activation of phosphoinositide 3-kinase, which leads to the phosphorylation of the activation loop of PKC ϵ at T566 by phospholipid-

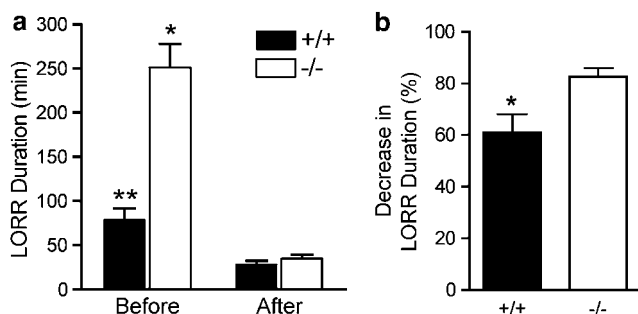


Figure 4 Duration of the ethanol-induced LORR before and after chronic ethanol treatment. (a) The duration of the LORR was greater in PKC ϵ (–/–) mice compared with PKC ϵ (+/+) mice before chronic ethanol exposure, but after chronic exposure, it was reduced in both genotypes and there was no longer a difference between them. Two-way ANOVA showed main effects of genotype ($F(1,27) = 28.02$; $p < 0.001$) and treatment ($F(1,27) = 62.04$; $p < 0.001$) with an interaction between these factors ($F(1,27) = 23.73$; $p < 0.001$). * $p < 0.05$ compared with PKC ϵ (+/+) mice before and with PKC ϵ (–/–) mice after chronic ethanol exposure and ** $p < 0.05$ compared with PKC ϵ (+/+) mice after chronic ethanol exposure (*post hoc* Tukey's tests). (b) The percent decrease in LORR duration that accompanied chronic treatment with alcohol was greater in PKC ϵ (–/–) than in PKC ϵ (+/+) mice (* $p < 0.05$ by two-tailed *t*-test).

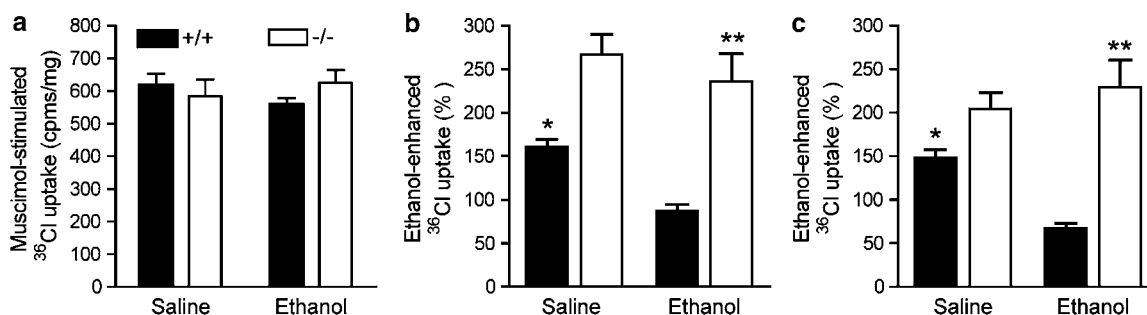


Figure 5 Acute tolerance of cerebellar GABA_A receptors to ethanol. (a) $^{36}\text{Cl}^-$ uptake stimulated by muscimol alone was not different between microsacs from PKC ϵ (–/–) mice (open bars) and PKC ϵ (+/+) mice (filled bars) and was not affected by prior *in vivo* exposure to 4 g/kg ethanol in either genotype. (b and c) Addition of 20 mM ethanol *in vitro* enhanced muscimol-stimulated $^{36}\text{Cl}^-$ uptake to a greater extent in microsacs prepared from saline-injected PKC ϵ (–/–) mice (open bars) as compared to microsacs from saline-injected PKC ϵ (+/+) mice (filled bars). Exposure to 2 g/kg (b) or 4 g/kg (c) ethanol *in vivo*, blocked enhancement of muscimol-stimulated $^{36}\text{Cl}^-$ flux by 20 mM ethanol *in vitro* in microsacs from PKC ϵ (+/+) mice but not in microsacs from PKC ϵ (–/–) mice. Two-way ANOVA for mice treated with 2.0 g/kg ethanol showed main effects of genotype ($F(1,33) = 26.87$; $p < 0.0001$) and *in vivo* treatment ($F(1,33) = 4.48$; $p = 0.042$). * $p < 0.05$ compared to microsacs from saline-treated PKC ϵ (–/–) mice and microsacs from ethanol-treated PKC ϵ (+/+) mice; ** $p < 0.05$ compared with microsacs from ethanol-treated PKC ϵ (+/+) mice (*post hoc* Newman–Keuls tests). Two-way ANOVA for mice treated with 4.0 g/kg ethanol showed a main effect of genotype ($F(1,13) = 51.78$; $p < 0.0001$) and an interaction between *in vivo* treatment and genotype ($F(1,13) = 12.10$; $p = 0.0041$). * $p < 0.05$ compared to microsacs from saline-treated PKC ϵ (–/–) mice and microsacs from ethanol-treated PKC ϵ (+/+) mice; ** $p < 0.05$ compared with microsacs from ethanol-treated PKC ϵ (+/+) mice (*post hoc* Bonferroni tests).

dependent kinase 1 (Cenni *et al*, 2002). Phosphorylation at this site is required for enzymatic activity (Parekh *et al*, 1999; Takahashi *et al*, 2000). This permits autophosphorylation at T710 and subsequent phosphorylation at S729 either by autophosphorylation (Cenni *et al*, 2002) or by a different kinase (Parekh *et al*, 1999). Phosphorylation at S729 permits full activity of PKC ϵ when the enzyme is stimulated by lipid second messengers (Parekh *et al*, 1999; Takahashi *et al*, 2000). Thus, PKC ϵ phosphorylation at S729 has been used as a marker for activation and post-translational processing of the immature pool of PKC ϵ that is not yet fully phosphorylated (Saitoh *et al*, 2001; Bayer *et al*, 2003; Zhou *et al*, 2003; Olive *et al*, 2005). Therefore, we used a phospho-specific antibody that detects phosphorylation of PKC ϵ at S729 to investigate the regulation of the immature PKC ϵ pool by ethanol.

Treatment with 1.5 or 4 g/kg ethanol *i.p.* increased phospho-S729-PKC ϵ immunoreactivity in the cerebellum of PKC ϵ (+/+) mice when compared to saline-treated mice (Figure 6a and b). Compared to the cerebellum, phospho-S729-PKC ϵ immunoreactivity in the frontal cortex and the striatum was relatively high in saline-treated mice and was not increased further by *in vivo* ethanol administration (Figure 6a). The ethanol-induced increase in cerebellar phospho-S729-PKC ϵ was detected 60 min, but not 30 min after *in vivo* ethanol administration (Figure 6b). Ethanol

treatment did not alter the amount of total cerebellar PKC ϵ immunoreactivity (Figure 6c).

Ethanol Preference Following Chronic Tolerance

We previously found that PKC ϵ (–/–) mice self-administer less ethanol than PKC ϵ (+/+) mice in both a two-bottle choice procedure (Hodge *et al*, 1999) and in an operant paradigm (Olive *et al*, 2000). We hypothesized that if chronic tolerance to the hypnotic and ataxic effects of ethanol is an important determinant of ethanol preference, then ethanol preference drinking might increase in PKC ϵ (–/–) mice following chronic ethanol exposure, when PKC ϵ (–/–) mice displayed chronic tolerance to ethanol. To test this, we assessed two-bottle choice drinking of 10% ethanol and water in PKC ϵ (–/–) and PKC ϵ (+/+) mice before and after exposure to twice daily injections of 2.5 g/kg ethanol for 6 days. Before chronic ethanol treatment, PKC ϵ (–/–) mice consumed less ethanol (Figure 7a) and showed reduced preference for ethanol (Figure 7b) when compared with PKC ϵ (+/+) mice. This pattern of drinking remained unchanged following parenteral administration of ethanol for 6 days.

Ethanol Clearance Following Chronic Ethanol Exposure

Genotype-related changes in ethanol-induced behaviors could be owing to changes in ethanol clearance rather than in tolerance to ethanol. PKC ϵ (–/–) and PKC ϵ (+/+) mice do not differ in their rate of ethanol clearance when examined in the ethanol-naïve state (Hodge *et al*, 1999). To determine if this is true following repeated ethanol exposure, we analyzed BECs following injection of 3.6 g/kg ethanol before and after chronic exposure to ethanol, 2.5 g/kg, twice daily, for 6 days (Figure 8). We found no difference between genotypes or any effect of chronic ethanol exposure on these measurements.

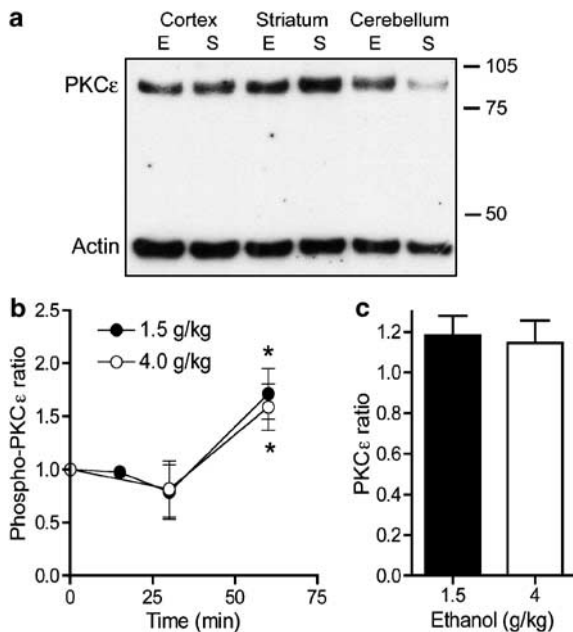


Figure 6 Phospho-PKC ϵ immunoreactivity following *in vivo* ethanol treatment. (a) Pairs of mice were injected with 4 g/kg ethanol (E) or saline (S) and killed 1 h later. Shown is a representative Western blot demonstrating an ethanol-induced increase in phospho-PKC ϵ immunoreactivity (92 kDa bands) relative to actin (43 kDa bands) in the cerebellum, but not in the striatum or the frontal cortex. (b) Time course of ethanol-induced increases in phospho-PKC ϵ immunoreactivity in mice injected with 1.5 or 4 g/kg ethanol. One-way ANOVA showed an effect of time in mice treated with 1.5 g/kg ($F(3,22) = 5.91$; $p = 0.005$) or 4 g/kg ($F(2,22) = 5.148$; $p = 0.0147$) ethanol. * $p < 0.05$ compared with zero time (Dunnett's tests). (c) Injection with 1.5 or 4 g/kg ethanol *in vivo* did not alter the abundance of total cerebellar PKC ϵ immunoreactivity relative to actin immunoreactivity measured 1 h later.

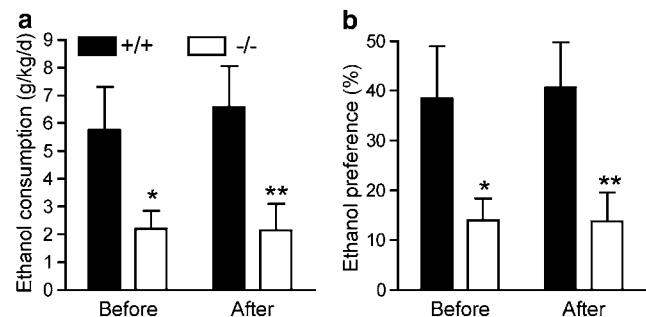


Figure 7 Two-bottle choice ethanol consumption and preference before and after chronic ethanol exposure. PKC ϵ (–/–) mice showed decreased consumption of 10% (v/v) ethanol (a) and preference for the ethanol containing solution (b) before and after chronic ethanol treatment when compared with PKC ϵ (+/+) mice. For ethanol consumption, two-way ANOVA showed a main effect of genotype ($F(1,25) = 9.41$; $p = 0.0051$) but not of treatment ($F(1,25) = 0.088$; $p > 0.05$). For ethanol preference, there was also a main effect of genotype ($F(1,25) = 9.24$; $p = 0.0055$) but not of treatment ($F(1,25) = 0.014$; $p > 0.05$). * $p < 0.05$ compared with PKC ϵ (+/+) mice before and ** $p < 0.05$ compared with PKC ϵ (+/+) mice after chronic treatment with ethanol (Newman–Keuls tests).

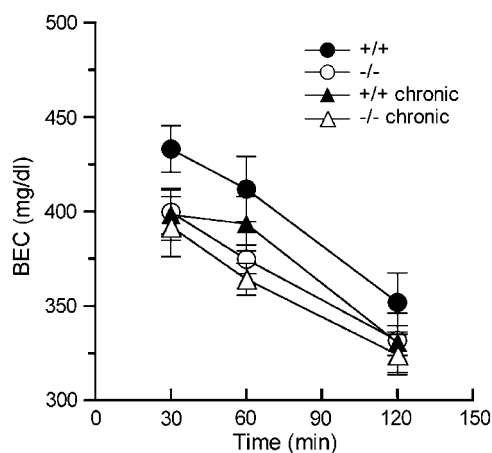


Figure 8 Ethanol clearance before and after chronic treatment. BECs in samples from PKC ϵ (-/-) (○, △) and PKC ϵ (+/+) (●, ▲) mice obtained following an acute injection of 3.6 g/kg ethanol before (○, ●) or after (△, ▲) chronic ethanol exposure. Analysis of these data by three-way ANOVA with genotype and prior chronic ethanol treatment as between-subject factors and time as a repeated measure revealed a main effect of time ($F(2,28) = 33.6$; $p < 0.001$) without significant effects of genotype ($F(1,14) = 4.18$; $p > 0.05$) or prior ethanol treatment ($F(1,14) = 3.91$; $p > 0.05$), or an interaction between these variables.

DISCUSSION

Our findings establish that PKC ϵ contributes to AFT to the ataxic and hypnotic effects of ethanol. We found that the duration of the ethanol-induced LORR and the extent of ethanol-induced rotarod ataxia were increased in mice lacking PKC ϵ . Likewise, recovery from each behavioral task occurred at lower BECs in PKC ϵ (-/-) than in PKC ϵ (+/+) mice. This was associated with a decrease in the ED₅₀ for the initiation of ethanol-induced loss of righting indicating that PKC ϵ contributes to hypnotic sensitivity to ethanol. In contrast, the ED₅₀ concentration of ethanol that produced rotarod ataxia was not different in wild-type and PKC ϵ (-/-) mice, suggesting that the major effect of PKC ϵ is to regulate AFT rather than initial sensitivity to ethanol-induced ataxia. We observed a similar effect of PKC ϵ on AFT to ethanol-induced loss of balance on the stationary dowel where we found that both the magnitude and rate at which AFT developed were reduced in PKC ϵ (-/-) mice by about 50% compared with wild-type littermates. In contrast, the calculated initial sensitivity to ethanol-induced loss of balance on the stationary dowel was similar between genotypes. Together, these results suggest that PKC ϵ participates in the development of AFT to the motor-impairing effects of ethanol in mice.

In contrast to its role in AFT, PKC ϵ does not appear to mediate chronic tolerance to the hypnotic or ataxic effects of ethanol. PKC ϵ (+/+) and PKC ϵ (-/-) mice treated with 2.5 g/kg ethanol twice daily for 6 days developed chronic tolerance to ethanol-induced LORR and to ethanol-induced ataxia on the stationary dowel such that at the end of the treatment period, there was no longer a difference between the genotypes in these behaviors. We therefore conclude that PKC ϵ contributes to AFT in ethanol-naïve mice but does not promote the development of chronic tolerance. Rather, our results suggest that PKC ϵ impairs the develop-

ment of chronic tolerance as PKC ϵ (-/-) mice showed a greater decrease in LORR than wild-type mice after chronic treatment and therefore developed chronic tolerance to a greater extent than wild-type mice (Figure 4b). Likewise, after chronic ethanol exposure, PKC ϵ (-/-) mice showed a greater increase in AFT and in AFT rate than PKC ϵ (+/+) mice on the stationary dowel. These results resemble those described for lines of mice selected for high (HAFT) and low (LAFT) AFT to ethanol (Wu *et al.*, 2001). These mice show genotypic differences in AFT when alcohol naïve but, following chronic ethanol exposure, show equivalent responses to ethanol-induced loss of balance on the stationary dowel and ethanol-induced LORR. Together with our findings, these results indicate that AFT and chronic tolerance can be genetically dissociated.

Tolerance to ethanol has been proposed to promote increased ethanol consumption through two mechanisms: tolerance to the rewarding effects of the drug, requiring that a subject drink more to achieve a pleasurable effect, and tolerance to its aversive properties, which reduces a disincentive to drink (Kalant, 1996). However, we found that despite treatment that produced chronic tolerance to motor impairing and hypnotic effects of ethanol as well as increased AFT, both PKC ϵ (+/+) and PKC ϵ (-/-) showed no change in ethanol preference drinking. These results indicate that AFT and chronic tolerance to ataxic and hypnotic effects of ethanol can be dissociated from ethanol preference drinking and suggest that these behaviors, at least in part, involve distinct neural systems.

As the PKC ϵ null allele is present from birth, the phenotypes we observed could be owing to changes in the nervous system resulting from the absence of PKC ϵ during development. In addition, since the null allele was generated on a 129S4 background and we studied hybrid C57BL/6 \times 129S4 mice, it is possible that the behavioral phenotypes we observed were due to 129S4 alleles neighboring the PKC ϵ locus, and not to the PKC ϵ null mutation. However, we previously showed that conditional rescue of PKC ϵ using a tetracycline-regulated transgene restores hypnotic responses to ethanol in PKC ϵ (-/-) mice to wild-type levels (Choi *et al.*, 2002). Thus, differences in behavior on the LORR test are unlikely to be owing to developmental effects or neighboring genes. We have not examined AFT to ethanol-induced ataxia using transgenic rescue or inbred 129S4 mice and therefore cannot entirely eliminate the possibility that our results are owing to neighboring genes or developmental changes for these phenotypes.

An additional finding in this study was that *in vivo* exposure to 2.0 or 4.0 g/kg ethanol led to GABA_A receptor resistance to ethanol measured 60 min later in cerebellar microsacs from PKC ϵ (+/+) mice. This was similar to findings reported by Allan and Harris (1987), who detected acute tolerance of cerebellar GABA_A receptors to ethanol *in vitro* 60 min after ethanol injection *in vivo*. Our finding that this response was absent in tissue from PKC ϵ (-/-) mice suggests that it is PKC ϵ -dependent. PKC ϵ -dependent, acute tolerance of GABA_A receptors could contribute to the accelerated recovery from ethanol intoxication that we observed in wild-type mice in assays of LORR and ataxia 60 min after ethanol injection. We did not examine GABA_A receptor function at time points less than 1 h after ethanol exposure, but Allan and Harris (1987) found no difference

in ethanol response at 30 min after ethanol injection. This suggests that other PKC ϵ -dependent mechanisms that do not involve cerebellar GABA $_A$ receptors also contribute to recovery from ethanol-induced rotarod ataxia in wild-type mice, particularly at early time points.

In the stationary dowel test, we injected mice with 1.5 g/kg ethanol, which is lower than the doses (2.0 and 4.0 g/kg) that we administered to mice used for GABA $_A$ receptor studies. However, we used two injections of ethanol in the stationary dowel test and the mean BEC 5 min after the second injection was 311 ± 11 mg/dl ($n = 11$). This was significantly greater ($p < 0.0001$) than the mean BEC measured 5 min after injection of a single dose of 2.0 g/kg ethanol (247 ± 8 mg/dl; $n = 14$). Therefore, the BECs achieved during the stationary dowel test after the second ethanol injection were within the range of concentrations attained in animals that showed GABA $_A$ receptor resistance to ethanol *in vitro*. Moreover, it took an average of 2.32 ± 0.13 h for wild-type mice ($n = 9$) to recover on the stationary dowel test after the second ethanol injection, which is much greater than the time (1 h) needed for development of GABA $_A$ receptor resistance to ethanol. Therefore, it is likely that acute tolerance of GABA $_A$ receptors to ethanol contributed to AFT in wild-type mice during the stationary dowel test.

Another finding in this study was that injection of either 1.5 or 4.0 g/kg ethanol increased PKC ϵ phosphorylation at S729 in the cerebellum of wild-type mice. As increased phosphorylation of S729 reflects an increase in the amount of PKC ϵ that can be fully activated by lipid second messengers (Saitoh *et al*, 2001; Bayer *et al*, 2003; Zhou *et al*, 2003; Olive *et al*, 2005), these results suggest that ethanol stimulates a signaling cascade that increases the pool size of fully phosphorylated PKC ϵ in the cerebellum. Although ethanol-stimulated PKC ϵ phosphorylation at S729 in wild-type mice, we were only able to measure this increase 60 min following ethanol injection and not at 30 min or earlier time points. However, in our study of rotarod ataxia, we observed some recovery in fall latency in wild-type mice as early as 15 min after injection of 2 g/kg ethanol. As we could not detect an increase in PKC ϵ S729 phosphorylation at this time, these results suggest that there are additional PKC ϵ -dependent mechanisms that contribute to AFT at earlier times (< 60 min), that do not involve post-translational processing of PKC ϵ . One possibility is that ethanol activates another signaling pathway that requires the presence of a basal level of PKC ϵ activity, or that ethanol increases the production of lipid second messengers, which activate the pre-existing pool of fully phosphorylated, mature PKC ϵ without increasing PKC ϵ processing.

In summary, when taken together with previous studies using PKC ϵ ($-/-$) mice and peptide inhibitors of PKC ϵ , which suggest that PKC ϵ attenuates the GABA $_A$ receptor response to ethanol (Hodge *et al*, 1999), our data provide evidence for a negative feedback loop that contributes to acute tolerance of GABA $_A$ receptors to ethanol. Given the role of cerebellar GABA $_A$ receptors in hypnotic and ataxic responses to ethanol (Palmer *et al*, 1987; Clark and Dar, 1988), such a PKC ϵ -GABA $_A$ receptor signaling pathway could promote AFT to ethanol.

Why PKC ϵ ($-/-$) mice show reduced ethanol self-administration remains elusive. It will be important in future work to measure rewarding and aversive properties

of ethanol in these animals, especially since ethanol does not elicit dopamine release from the nucleus accumbens of PKC ϵ ($-/-$) mice (Olive *et al*, 2000), suggesting that these mice might experience less ethanol-induced reward than wild-type animals. We also do not yet know how ethanol stimulates PKC ϵ phosphorylation in the cerebellum or which proteins are phosphorylated by PKC ϵ to regulate the sensitivity of GABA $_A$ receptors to ethanol. It will be of interest to identify upstream and downstream mediators of these events, as mapping of this signaling pathway could uncover additional proteins besides PKC ϵ that regulate the development of AFT to ethanol, ethanol self-administration, and contribute to risk of alcoholism in humans.

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