Limbic Epileptogenesis in a Mouse Model of Fragile X Syndrome

Fragile X syndrome (FXS), caused by silencing of the Fmr1 gene, is the most common form of inherited mental retardation. Epilepsy is reported to occur in 20–25% of individuals with FXS. However, no overall increased excitability has been reported in Fmr1 knockout (KO) mice, except for increased sensitivity to auditory stimulation. Here, we report that kindling increased the expressions of Fmr1 mRNA and protein in the forebrain of wild-type (WT) mice. Kindling development was dramatically accelerated in Fmr1 KO mice, and Fmr1 KO mice also displayed prolonged electrographic seizures during kindling and more severe mossy fiber sprouting after kindling. The accelerated rate of kindling was partially repressed by inhibiting N-methyl-D-aspartic acid receptor (NMDAR) with MK-801 or mGluR5 receptor with 2-methyl-6-(phenylethynyl)-pyridine (MPEP). The rate of kindling development in WT was not effected by MPEP, however, suggesting that FMRP normally suppresses epileptogenic signaling downstream of metabolic glutamate receptors. Our findings reveal that FMRP plays a critical role in suppressing limbic epileptogenesis and predict that the enhanced susceptibility of patients with FXS to epilepsy is a direct consequence of the loss of an important homeostatic factor that mitigates vulnerability to excessive neuronal excitation.

Keywords: epilepsy, FMRP, kindling, mGluR5, mossy fiber sprouting, NMDA

Introduction

Fragile X syndrome (FXS), the most common form of inherited mental retardation, results from a trinucleotide repeat (CGG) expansion in the 5'-untranslated region of the gene Fmr1 (Verkerk et al. 1991). The CGG repeat expansion and subsequent hypermethylation cause transcriptional silencing of Fmr1 and loss of its encoded protein, the fragile X mental retardation protein (FMRP) (O'Donnell and Warren 2002; Jin and Warren 2003). Murine FMRP is a brain-specific RNA-binding protein that suppresses translation of target RNAs. Through interactions with kinesin, FMRP is able to travel between distal neurites and the soma of neurons and thereby regulate protein synthesis locally within specific cellular compartments (Ashley et al. 1993; Warren and Nelson 1994; Eberhart et al. 1996; Brown et al. 1998; Darnell et al. 2001; Li et al. 2001; Ohashi et al. 2002; Ling et al. 2004; Davidovic et al. 2007).

Patients with FXS exhibit mental retardation, attention deficits, autistic behavior, macroorchidism, and facial abnormalities (Turner et al. 1980; Opitz and Sutherland 1984; Sobesky et al. 1996). Additionally, epilepsy is reported to occur in 20–25% of individuals with FXS and paroxysmal electroencephalography (EEG) abnormalities are present in about 50% of prepubescent boys with FXS (Musumeci et al. 1999; Saba-annam et al. 2001; Hagerman et al. 2002). These EEG pattern abnormalities resemble those of patients with benign focal epilepsy of childhood (Berry-Kravis 2002). FMRP deficiency is speculated to lead to increased neuronal excitability and susceptibility to seizures in humans. However, spontaneous seizures are never observed in Fmr1 knockout (KO) mice and sensitivity to convulsants (such as bicuculline, pentylentetrazol and kainic acid) is not significantly different from wild-type (WT) mice (Chen and Toth 2001). The most substantial deficit observed in Fmr1 KO mice is increased sensitivity to auditory seizures (Musumeci et al. 2000; Chen and Toth 2001), suggesting that Fmr1 KO mice do not have an overall increased excitability phenotype but rather auditory system specific hyperexcitability.

Recent studies have challenged this conclusion. In hippocampal slices from Fmr1 KO mice, increasing neuronal activity by blocking γ-aminobutyric acidergic (GABAergic) transmission by the GABA_A receptor antagonist bicuculline induces prolonged epileptiform discharges (Chuang et al. 2005). Furthermore, these prolonged discharges can be suppressed with the mGluR5 antagonist 2-methyl-6-(phenylethynyl)-pyridine (MPEP). These observations suggest that FMRP does influence neuronal excitability in the forebrain, possibly by repressing group I mGluR-mediated RNA translation (Chuang et al. 2005). Interestingly, MPEP could also reverse other symptoms of Fmr1 KO mice, including sound-induced seizures (Bear et al. 2004; Chuang et al. 2005; McBride et al. 2005; Yan et al. 2005). These findings support the mGluR theory of FXS (Bear et al. 2004; Ronesi and Huber 2008).

To investigate seizure susceptibility in the limbic system of Fmr1 KO mice in vivo, we used a well-established animal model of limbic epilepsy, amygdala kindling, to explore the role of FMRP in limbic epileptogenesis. Kindling mimics epileptogenesis in which repeated administration of an initially subconvulsive electrical stimulation of the amygdala results in progressive intensification of behavioral seizures and prolongation of electrographic afterdischarges (AD) (Goddard et al. 1969; Racine 1972). We found that kindling stimulation increased the expression of Fmr1 mRNA and protein in the forebrain of WT mice. The development of behavioral seizures was dramatically accelerated in the Fmr1 KO mice, and AD durations were greatly exaggerated throughout kindling. Furthermore, exuberant mossy fiber sprouting (MFS) was observed in fully kindled Fmr1 KO mice. Behavioral seizures could be repressed by administering the N-methyl-D-aspartic acid receptor antagonist MK801 or mGluR5 antagonist MPEP in Fmr1 KO mice. Interestingly, MPEP did not alter kindling development in WT mice, suggesting that FMRP normally suppresses the epileptogenic effects of mGluR5 activity. Our
findings indicate that FMRP-mediated transcriptional repression plays a critical role in moderating susceptibility to limbic epileptogenesis.

Materials and Methods

Experimental Animals

Male Fmr1 KO mice of the FVB strain (The Dutch-Belgian Fragile X Consortium 1994) and their WT littermates were used. All animals were maintained on a 12:12 h light-dark cycle with a constant room temperature. Mice were group housed (3 to 4 per cage) and provided with ad lib food and water. Kindling experiments were conducted during the light phase of the cycle on 12-week-old adult mice. Mice were handled according to the guidelines prescribed by the Animal Care and Use Committee of the Institute of Neuroscience, Chinese Academy of Sciences. All efforts were made to minimize suffering and the number of animals used.

Drug Application

The NMDA antagonist MK801 (Sigma, St Louis, MO) and the mGluR5 antagonist MPEP (Tocris, Biotrend Chemikalien, Köln, Germany) were dissolved in sterile saline (0.9% NaCl). Drugs were administered by intraperitoneal injection (MK801, 0.5 mg/kg; MPEP, 30 mg/kg) 30 min before each stimulation. Application of the transcription inhibitor actinomycin D (Sigma) was performed on freely moving animals via a cannula inserted into ventricle (coordinates: 3 mm lateral and 2.3 mm posterior to bregma, 4 mm below dura) 30 min before a class 5 seizure-inducing stimulation. Actinomycin D was dissolved in saline (0.9% NaCl) to a final concentration of 50 μg/ml. Actinomycin D 5 μl was intracerebroventricular (i.c.v.) injected within 10 min.

Surgery and Kindling Procedure

A twisted bipolar electrode was implanted into the right amygdala (coordinates: 2.9 mm lateral and 1.2 mm posterior to bregma, 4.6 mm below dura) of animals under pentobarbital (60 mg/kg) anesthesia. Animals were then given 10 days to recover. The electrographic seizure threshold (EST) for each individual mouse was determined by applying 1-s train of 1-ms biphasic rectangular pulses at 60 Hz beginning at 50 μA. Additional stimulations increasing by 10 μA were administered at 2-min intervals until an electrographic seizure lasting at least 5 s was evoked. Stimulation at the EST intensity were subsequently applied once daily. EEGs and behavioral seizures were observed and recorded. The severity of the behavioral manifestations of seizures was classified according to the criteria of Racine (1972). Fully kindled is defined by the occurrence of 3 consecutive seizures of class 4 or greater. All surgery and kindling procedures were performed blind to genotype. Unstimulated control animals of each genotype underwent surgical implantation of an electrode in the amygdala and were handled identically but were not stimulated. Electrode placement was confirmed by methyl green pyronine-Y staining. Data derived from animals with correct electrode placement were analyzed.

RNA Extraction, cDNA Synthesis, Reverse Transcription Polymerase Chain Reaction, and Real-Time Polymerase Chain Reaction

Two weeks after being fully kindled, WT mice were killed either 3 h after a single class 5 seizure or without any further stimulation. Four microliters of first-strand cDNA preparation was used as template in polymerase chain reaction (PCR) reactions with the following sense and antisense oligonucleotide primers: 1) P1 (5'–CTCAAAGGGACGCCCATC) and P2 (5'–TACCATCCTGGTCCAGAC), these primers direct the amplification of nucleotide 793 to 1456 of mouse Fmr1 cdNA and 2) P3 (5'–CTGCCCAGAATCATCCTT) and P4 (5'–CCACACCCTGTTGC. TTAGT), these primers direct the amplification of nucleotide 579 to 947 of mouse Gapdh cDNA. The cDNA was amplified with Taq polymerase under the following PCR conditions for Fmr1: 28 cycles of denaturation at 94 °C for 30 s, annealing at 55 °C for 30 s, and extension at 72 °C for 1 min and for Gapdh 20 cycles of 94 °C for 30 s, 60 °C for 30 s, and 72 °C for 1 min. PCR products were analyzed in 1% agarose gels.

Real-time PCR analysis of Fmr1 expression was performed using SYBR Green I in a 20-μl reaction with an ABI Prism 7000 thermocycler (Applied Biosystems, Foster City, CA). Primers used for the amplification of Fmr1 were P5 (5'–GTGGTTA GCTAAAGTGAGGATGAT) and P6 (5'–CAGGTTGTGTTGAGTTAACAGATC). Primers used to amplify Gapdh were the same as those listed above (P3 and P4). Products were amplified with HotStart ExTaqDNA polymerase under the following PCR conditions: denaturation at 95 °C for 5 min followed by 45 cycles of 10 s at 95 °C and 31 s at 60 °C. Dissociation curve analysis was performed after amplification to confirm the absence of nonspecific amplification products and primer-dimers. Real-time PCR results were analyzed with ABI Prism SDS 7000 software (Applied Biosystems).

In Situ Hybridization

Two weeks after being fully kindled, WT mice were perfused transcardially for 10 min with 4% paraformaldehyde in 0.1 M phosphate buffer (PB, pH 7.4) either 3 h after a single class 5 seizure or without any further stimulation. Brains were removed and postfixed in the same fixative for 4 h at 4 °C. After postfixation, brains were immersed in 20% sucrose in PB overnight at 4 °C. Coronal cryostat sections (20 μm) were mounted onto Fisher Super Frost Microscope slides and stored at -80 °C.

Antisense probes against Fmr1 nucleotides 1126-1670 were designed. Reverse transcription for polymerase chain reaction (RT-PCR) products were subcloned into pEGM-T Easy vector (Promega, Madison, WI) and digoxigenin (DIG)-labeled sense and antisense ribonucleotide probes were generated by in vitro transcription using T7 and Sp6 RNA polymerases. Slides were prehybridized in hybridization buffer for 4 h at 65 °C and then hybridized with probes (1 μg/ml) in hybridization buffer at 65 °C overnight. After extensive washing, slides were incubated with preabsorbed alkaline phosphatase-conjugated anti-DIG antibody (Roche, Switzerland) at 4 °C overnight. Signals were developed by nitroretrazolium blue-5-bromo-4-chloro-3-indoly phosphate color reaction in the dark for 6-8 h. Samples were analyzed using an Nikon E600FN upright microscope.

Western Blot

Frozen forebrains extracted at the time points described in the Results section were thawed on ice and homogenized in lysis buffer containing 50 mM Tris-HCl (pH 7.4), 20 μg/ml leupeptin, 20 μg/ml aprotonin, 0.1 mM phenylmethylsulfonyl fluoride, 1 mM EDTA, 1 mM sodium orthovanadate, 1.0% deoxycholate. The homogenate was incubated on ice for 30 min to allow complete lysis and then centrifuged for 20 min at 14 000 x g (4 °C) to remove nuclear fractions, and protein concentrations were then determined using the bicinchoninic acid kit (BCA kit; Pierce, Rockford, IL). Proteins were separated by 9% sodium dodecyl sulfate-polyacrylamide gel and transferred to nitrocellulose membranes with a transblot semi-dry apparatus (Bio-Rad, Hercules, CA). Membranes were blocked in Tris-Buffered saline Tween-20 (TBST; 0.05% Tween-20 in TBS) containing 5% nonfat dry milk for 1 h and then incubated with anti-FMRP monoclonal antibody 1C3 (1:500; Chemicon, Temecula, CA) or anti-Glyceraldehyde 3-phosphate dehydrogenase monoclonal antibody (1:5000, KangChen, Shanghai, China) at 4 °C overnight. Membranes were washed in TBST buffer and incubated for 1 h with horseradish peroxidase-conjugated goat anti-mouse secondary antibody (1:5000, Chemicon) at room temperature. Immunoblots were developed with enhanced chemiluminescence (Pierce). Quantitative
analysis was conducted using Image Quant software (Molecular Dynamics, Sunnyvale, CA).

**Immunohistochemistry**

Free-floating sections were treated in 1% H2O2 for 30 min at room temperature and incubated for 5 min in 0.5% Triton X-100, washed in phosphate-buffered saline (PBS), and blocked with 5% NGS in PBS for 1 h at room temperature. Slices were then incubated with anti-FMRP (1:200; Chemicon) overnight at 4 °C, washed, and incubated with biotinylated goat anti-mouse IgG (1:200; Boster, Wuhan, China) for 1 h at room temperature. After being washed with PBS, slices were reacted with peroxidase-conjugated avidin-biotin complexes (Vector, Burlingame, CA; 1:100) for 1 h at room temperature. Signals were developed with 0.03% 3,3-diaminobenzidine tetrahydrochloride dihydrate (Sigma) in 50 mM Tris buffer containing 0.003% H2O2, pH 7.6. Negative control sections were prepared by omitting primary antibodies. Samples were analyzed under a Nikon E600FN upright microscope.

**The Fluoro-Jade B Staining**

For cell death analyses, both kindled WT and Fmr1 KO mice were sacrificed 24 h after being fully kindled. WT and Fmr1 KO brain sections were prepared as described in the In Situ Hybridization. Mounted sections were rehydrated in an ethanol series (3 min 100% ethanol, 1 min 70% ethanol, 1 min 50% ethanol, and 1 min distilled water) and then incubated in a solution of 0.06% potassium permanganate for 10 min on a rotating platform, rinsed in distilled water for 2 min, and transferred to fluoro-jade B (FJB) staining solution (0.0004% FJB in 0.1% acetic acid, prepared 10 min before use) for 20 min. Labeled sections were imaged with a confocal laser-scanning microscope (Olympus LSM-GB200) under green (FJB) fluorescence. Staining and data analysis were done by an individual blinded to the genotype and treatment.

**Timm Staining**

Fully kindled WT and Fmr1 KO mice were killed either 5 or 28 weeks later after the last stimulation. Mice were perfused transcardially with saline for 2 min, followed by 0.16% sodium sulfide in PBS for 4 min, and followed by saline for 1 min and 4% paraformaldehyde in PBS for 10 min. Brains were then removed and postfixed overnight at 4 °C and immersed in 20% sucrose at 4 °C the following day. Horizontal sections (20 μm) were collected in PBS (pH 7.4) and mounted onto gelatin-coated slides and developed in the dark at room temperature for 30-50 min in a 12:6:2 mixture of gum arabic (50% w/v), hydroquinone solution (5.67% w/v), citric acid-sodium citrate buffer (26% citric acid w/v; 24% sodium citrate, w/v), and silver nitrate (17% w/v). After washing thoroughly with running water, sections were dehydrated, cleared, and coverslipped. To assess MSB, the Timm index was measured using Image-Pro Plus 5.0 software (Media Cybernetics, Bethesda, MD) by dividing the total area of Timm granules by total length of dentate gyrus (DG; Watanabe et al. 1996). The Timm index of each mouse was calculated as the mean of 3 serial sections through the hippocampus of each animal.

**Nissl Staining and Cell Number Counting**

Cell death was assessed in serial horizontal sections (20 μm) through the hippocampus. Every third slice between 4000 and 5300 μm from the dorsal surface of neocortex were placed into 1:1 alcohol:chloroform for 30 min and then rehydrated in an ethanol series (3 min each in 100%, 95%, 80%, 70%, 60%, 50%, and 0% ethanol in distilled water). Rehydrated slices were stained in cresyl violet solution for 30 min at room temperature then dehydrated (3 min each in 50%, 60%, 70%, 80%, 95%, and 100% ethanol), cleared in xylene, and mounted with resinous medium, and analyzed under a Nikon E600FN upright microscope. Cells in the hilus field were counted using Image-Pro Plus 5.0 software (Media Cybernetics).

**Statistical Analysis**

Statistical analyses were carried out with GraphPad Prism Software (San Diego, CA). For statistical analyses of kindling development, Timm indices, and cell number, analyzers were blinded to the genotypes and treatment. Kindling development data and real-time PCR result were compared with unpaired t test analysis, Timm indices were compared with 2-way ANOVA, post hoc Bonferroni’s test. Cell number and western blots were compared with 1-way ANOVA, post hoc Dunnett’s test.

**Results**

The Expressions of Fmr1 mRNA and FMRP are Upregulated by Seizure Activity

Previous studies have demonstrated that the expression of FMRP can be regulated by synaptic activity and experience (Weiler et al. 1997; Todd and Mack 2000; Todd et al. 2003; Gabel et al. 2004; Hou et al. 2006). To examine whether seizure activity regulates Fmr1 transcription, we investigated the level of Fmr1 mRNA expression after kindling-induced seizures (see Materials and Methods for a description of kindling procedures) by RT-PCR, real-time PCR, and in situ hybridization. Two weeks after being fully kindled, WT mice were killed either 3 h after a single stimulation, which was given to trigger a class 5 seizure, or without any further stimulation. RT-PCR and real-time PCR were performed using RNA from the forebrain of both unstimulated kindled WT mice and stimulated kindled WT mice. As shown in Figure 1A, the level of Fmr1 mRNA in the forebrain normalized to Gapdh increased significantly 3 h after kindling-induced seizure. Real-time PCR analysis showed that a single class 5 seizure caused a 2-fold increase of Fmr1 mRNA (n = 8) relative to unstimulated kindled mice (n = 8, P < 0.05).

In situ hybridization revealed that Fmr1 expression was strongly induced in the cortex from baseline levels 3 h after seizure activity. Fmr1 expression was also upregulated in the hippocampal CA1 region and DG, albeit to a lesser degree and from a higher baseline (Fig. 1B). All changes in Fmr1 mRNA expression after kindling stimulations were bilateral with no hemispheric bias. No difference in Fmr1 mRNA expression was observed between unstimulated fully kindled WT mice and unkindled, sham-stimulated controls (data not shown).

We next examined if the upregulation of Fmr1 mRNA by seizure activity resulted in a corresponding increase in protein expression. Total protein lysates from individual forebrains were assayed by western blot. The mean level of FMRP normalized to Gapdh in the forebrain of unstimulated fully kindled mice was essentially identical to that of unkindled controls (data not shown). As shown in Figure 2A, FMRP expression increased about 5-fold 3 h after class 5 seizure activity in kindled mice (n = 8) relative to unstimulated fully kindled mice (n = 12, P < 0.01), before declined back to baseline levels by 12 h (Fig. 2B).

To investigate the spatial distribution of seizure-induced FMRP expression, we performed immunohistochemical staining on hippocampal sections. In unstimulated fully kindled mice, FMRP was primarily expressed in the soma of CA1 pyramidal neurons and DG granule cells, similar to unkindled controls (Fig. 2C and data not shown). However, 3 h after a single class 5 seizure, the intensity of FMRP immunostaining was greatly increased in the soma of CA1 pyramidal neurons and in the soma of DG granule cells (Fig. 2C, right panel). No obvious immunostaining of either the dendrites or the nucleus of neurons in the hippocampus was observed in control or stimulated animals.

To investigate whether the expression of FMRP correlates with the development of kindling, we examined the expression of FMRP at different stages during kindling development. Protein lysates from the forebrain were extracted 3 h after the
first stimulation on the first day of kindling (1 stimulation; AD duration, 8.1 ± 0.5 s), 3 h after the first class 2 seizure (5 ± 1 stimulations; AD duration of class 2 seizure, 19.2 ± 2.7 s), or 3 h after the third consecutive class 4/5 seizure (12 ± 1 stimulations; AD duration of final class 4/5 seizure, 26.9 ± 2.2 s). Relative to sham-stimulated controls, no increase in FMRP was observed 3 h after the initial subconvulsive stimulation (data not shown), but FMRP expression in the forebrain significantly increased after the first class 2 seizure and was further enhanced after full kindling (Fig. 2D). Statistical analysis showed a 3.5-fold increase of FMRP expression after class 2 seizure (n = 5) and 5.2-fold increase of FMRP expression in fully kindled mice (n = 8) compared with controls (n = 17; P < 0.05 and P < 0.001, respectively) (Fig. 2E). Taken together, these data reveal that both mRNA and protein expression of Fmr1 gene are upregulated by kindling-induced seizure activity in vivo.

Finally, in order to decipher the transcriptional versus translational mechanisms of the seizure-induced upregulation of FMRP, we i.c.v. injected the transcription inhibitor actinomycin D 30 min before a class 5 seizure-inducing stimulation and then assayed Fmr1 mRNA or protein levels in the forebrain 3 h after by RT-PCR or western blotting. Actinomycin D did block the seizure-induced increase in the level of Fmr1 mRNA (n = 3; P < 0.05; Fig. 2F), but the increase in protein level was unaffected (Fig. 2G). This data suggests that seizure activity-induced upregulation of FMRP protein in WT mice is largely due to changes in the rate of translation.

**Accelerated Development of Kindling in Fmr1 KO Mice**

To study the functional significance of seizure-regulated FMRP expression, seizure behaviors and AD durations of WT and Fmr1 KO mice after daily amygdala stimulations were compared. There was no significant difference in EST between the 2 groups (Fig. 3A). The mean EST for WT and Fmr1 KO mice were 141.4 ± 9.77 μA (n = 25) and 139.7 ± 10.71 μA (n = 31), respectively. Once-daily stimulation of the amygdala resulted in progressive increase in the intensity of behavioral seizures in Fmr1 KO mice as in WT mice. However, the rate at which behavioral seizures progressed was dramatically accelerated in Fmr1 KO mice (Fig. 3B) with the number of stimulations required to induce 3 consecutive class 4/5 seizures (5.45 ± 0.40, n = 25) reduced to about half of the number of stimulations required for WT mice (11.5 ± 0.45, n = 31, P < 0.001) (Fig. 3C).

The differences in behavioral seizure intensity were paralleled by differences in electrophysiological measures of seizure duration. Fmr1 KO mice displayed prolonged AD after repeated kindling stimulations (Fig. 4A). By the third day of stimulation, the AD duration was doubled compared with that of WT mice (Fig. 4A). In WT mice, the duration of AD reached about 30 s by the 14th stimulation and additional stimulation did not induce any further prolongation of AD duration. In sharp contrast, the AD duration reached 30 s after only 3 stimulations in Fmr1 KO mice and climbed up to over 45 s by the 14th stimulation (Fig. 4A). Representative electrographs are shown in Figure 4B.

**Effects of NMDA Receptor and mGluR5 Antagonists on Epileptogenesis**

Studies of several animal models have shown that activation of ionotropic (Croucher et al. 1995) and metabotropic glutamate receptors (Wong et al. 1999) can promote epileptogenesis. We speculated that FMRP might function to suppress seizure-promoting actions of either one or both classes of glutamate receptors. To test this, we treated Fmr1 KO mice with the
noncompetitive NMDA receptor open-channel blocker MK801 or the metabolic glutamate receptor mGluR5 antagonist MPEP. Consistent with previous studies (McNamara et al. 1988; Sato et al. 1988; Löscher and Hönack 1991), the rate of kindling in WT mice was slowed down during the first 2 weeks of stimulation when MK801 was injected 30 min before each daily stimulation relative to controls injected with saline (Fig. 5A). Interestingly, for the first 9 days of stimulation, the acceleration of kindling development in Fmr1 KO mice was also suppressed by MK801 (Fig. 5B). During this period, there was no significant difference in the average seizure class between MK801 treated WT mice (n = 13) and MK801 treated Fmr1 KO mice (n = 13). After the 10th stimulation, kindling development in MK801-treated Fmr1 KO mice dramatically accelerated compared with the MK801-treated WT mice. These observations implicate the involvement of NMDARs in the progression of kindling in both WT and Fmr1 KO mice, and FMRP ablation does not obviously alter the suppressive effect of inhibiting NMDAR on kindling development.

**Figure 2.** FMRP expression is upregulated after seizure activity. (A) Representative western blot showing a transient increase of FMRP in the forebrain of kindled mice after an evoked seizure. Two weeks after fully kindling, a single class 5 seizure was induced in WT mice and forebrains were isolated 3 h or 12 h later. Each lane was loaded with an equal amount of protein extract from a single forebrain sample (mouse #1-6). Lane 1 and 2, unstimulated fully kindled WT mice; lane 3 and 4, fully kindled mice 3 h after a class 5 seizure; lane 5 and 6, fully kindled mice 12 h after a class 5 seizure. After a class 5 seizure, FMRP expression increases before returning to baseline levels by 12 h. (B) Quantitative analysis of western blot band intensities. FMRP immunoreactivity was normalized to Gapdh immunoreactivity and mean ± standard error of the mean (SEM) values are presented as a percentage of the mean level in unstimulated fully kindled mice; *P < 0.05; **P < 0.01; ***P < 0.001; 1-way ANOVA followed by post hoc Dunnett’s test. (C) Seizure activity leads to increased FMRP expression in the soma of CA1 pyramidal cells and granule cells in the DG. Coronal sections through the hippocampus of unstimulated fully kindled WT mice (left panel) and fully kindled WT mice 3 h after a class 5 seizure-inducing stimulation (right panel) were immunolabeled with anti-FMRP. Lower panels show higher magnification of upper panel, note that no staining is observed in neurites and nucleus of hippocampal neurons. Scale bars: 400 μm for upper 1 row; 100 μm for lower 2 rows. (D) Upregulation of FMRP correlates with kindling development. Forebrains of WT mice isolated 3 h after sham stimulation (lane 1), 3 h after the first class 2 seizure during kindling (lane 2), or 3 h after the third class 5 seizure during kindling (lane 3). (E) Quantitative analysis of FMRP expression in (D). FMRP levels were normalized to Gapdh and mean ± SEM values are presented as a percentage of the mean level in sham-stimulated mice; *P < 0.05; **P < 0.01; ***P < 0.001; 1-way ANOVA with post hoc Dunnett’s test. (F) Real-time PCR of Fmr1 RNA from control mice, saline-treated stimulated mice, actinomycin D-treated stimulated mice. Fmr1 transcript levels were upregulated at 3 h after stimulation in saline-treated mice and application of actinomycin D 30 min before stimulation repressed the increase of Fmr1. Gapdh transcript levels were used to normalize the levels of Fmr1 and values are presented as group mean ± SEM as a percentage of unstimulated samples; *P < 0.05; 1-way ANOVA, post hoc Dunnett’s test. (G) Representative western blot showing that application of actinomycin D did not obviously repress seizure-induced increases of FMRP.
In WT animals, pretreatment with MPEP (30 mg/kg) 30 min before each daily stimulation did not significantly affect the development of kindling (Fig. 6A), consistent with previous study (Nagaraja et al. 2004). There was no significant difference in the average seizure class between the saline-treated WT mice (n = 25) and MPEP-treated WT mice (n = 15) at any time point throughout kindling (Fig. 6A). In contrast, application of MPEP significantly decelerated the rate of kindling progression in Fmr1 KO mice up to the ninth day of stimulation. The average seizure class from the fourth through to the eighth stimulation point in MPEP-treated Fmr1 KO mice (n = 13) was significantly lower than saline-treated Fmr1 KO mice (n = 31) (Fig. 6B). Together, this data suggests that FMRP normally suppresses epileptogenic mGluR5 signal transduction and this activity is disinhibited in Fmr1 KO mice, thus accelerating kindling development.
Fmr1 mice and Fmr1 KO mice. Saline or MK801 was administered 30 min before each stimulation. Asterisks indicate statistically significant differences in the average behavior seizure class at the indicated time point between the 2 presented groups. *P < 0.05; **P < 0.01; ***P < 0.001; 2-tailed unpaired t-test.

Enhancement of Mossy Fiber Sprouting in Fully Kindled fmr1 KO Mice

MFS, characterized by an abnormal projection of the axon collaterals of granule cells into the inner molecular layer (IML) of the DG of the hippocampus (Sutula et al. 1989), is a feature of neuronal circuit reorganization associated with limbic epilepsy. We examined axon projections of hippocampal granule cells by Timm staining of WT and Fmr1 KO mice, 5 weeks or 28 weeks after 22 consecutive days of stimulation. Representative Timm-stained hippocampal sections from both kindled and age-matched, sham-stimulated WT and Fmr1 KO mice are shown in Figure 7A. The innervation’s pattern in sham-stimulated Fmr1 KO mice (n = 13) was not obviously different from that in sham-stimulated WT mice (n = 8) (Fig. 7A, d vs. a and j vs. g; Fig. 7B). Interestingly, striking enhancement of MFS in the IML was observed 5 weeks after 22 daily stimulations in the fully kindled Fmr1 KO mice, but no change was observed in fully kindled WT mice (Fig. 7A, e vs. b, k vs. h; Fig. 7B). Twenty-eight weeks after 22 daily stimulations, a slight increase of Timm staining in the IML was detected in WT kindled mice (Fig. 7A, i vs. g; Fig. 7B) but was less severe than that in Fmr1 KO mice 5 weeks after stimulation. At the 28 week time point, MFS in fully kindled Fmr1 KO mice was further enhanced (Fig. 7A, l vs. k). Statistical analyses revealed a 10-fold increase of Timm index in the IML of the DG in the 5 weeks later killed fully kindled Fmr1 KO mice and a 40-fold increase in 28 weeks later killed fully kindled Fmr1 KO mice compared with their age-matched fully kindled WT mice (Fig. 7B). The enhancement of Timm staining after kindling in Fmr1 KO mice was selective to the IML region. To determine whether the enhanced MFS in Fmr1 KO mice was accompanied with an increase in neuronal cell loss, we performed Fluoro-Jade B staining and Nissl staining (Fig. 8) of brain sections from kindled mice. No obvious neuronal degeneration was detected in either fully kindled WT or Fmr1 KO mice.

Discussion

Four principal findings emerged from this study. First, kindling induced by repeated electrical stimulations of the amygdala resulted in upregulation of the Fmr1 gene in the forebrain. Second, the development of kindling, as measured by behavioral and electrophysiological indices, was accelerated in Fmr1 KO mice. Third, inhibiting NMDAR slowed the early
progression of kindling development in both WT and Fmr1 KO mice. By contrast, inhibiting mGluR5 partially repressed the development of kindling in Fmr1 KO mice but had little effect on kindling development in WT mice. Forth, kindling-induced MFS into the IML region of the DG was dramatically enhanced in Fmr1 KO mice.

FMRP is upregulated in response to neuronal activity (Weiler et al. 1997; Todd and Mack 2000; Todd et al. 2003;)

**Figure 7.** Axon projections of granule cells in the hippocampus of sham-stimulated and fully kindled WT and Fmr1 KO mice. (A) Representative Timm staining of horizontal brain sections of sham-stimulated (sham) WT mice (a, g); sham Fmr1 KO mice (d, j); fully kindled (kindled) WT mice 5 weeks (b, h) or 28 weeks (c, i) after 22 daily stimulation; kindled Fmr1 KO mice 5 weeks (e, k) or 28 weeks (f, l) after 22 daily stimulations. Under control conditions (sham), Fmr1 KO mice show slightly more Timm-stained granules in the granule cell body layer compared with WT mice. Five weeks after 22 daily stimulations, MFS is obvious in the IML of the DG in Fmr1 KO mice. Twenty-eight weeks after 22 daily stimulations, MFS is dramatically more severe in Fmr1 KO mice. In contrast, WT mice show little MFS 28 weeks after 22 daily stimulations. Scale bars: upper 2 panels, 250 μm; lower 2 panels, 50 μm. (B) Statistical analysis of Timm index in the IML of the DG. Bars represent mean ± standard error of the mean. ***P < 0.001; 2-way ANOVA with post hoc Bonferroni’s test.

**Figure 8.** Fluoro-Jade B and Nissl staining reveal no obvious neuronal cell loss in Fmr1 KO mice after kindling. (A) Fluoro-Jade B staining. A WT hippocampal section 24 h after pilocarpine-induced seizure (340 mg/kg, intraperitoneally) with significant cell death in the hilus of the DG is shown as a positive control (n = 3). No cell death is detected in sections from sham WT (n = 3), kindled WT (n = 6), or kindled Fmr1 KO (n = 6) mice 24 h after the third consecutive class 5 seizure. Scale bar: 400 μm. (B) Nissl staining. Nissl-stained brain sections from sham WT mice, kindled WT mice, sham Fmr1 KO mice, and kindled Fmr1 KO mice reveal similar cell densities in the CA1, CA3, cortex, and hilus of the DG. Scale bars: top, 500 μm for upper one row; bottom, 250 μm for other rows. (C) Statistical analysis of cell number in the hilus. No significant difference was found among the 4 groups. Bars represent mean ± standard error of the mean; 1-way ANOVA, post hoc Dunnett’s test.
Gabel et al. 2004; Irwin et al. 2005; Hou et al. 2006) and plays an important role in synaptic plasticity by regulating the translation, subcellular localization, and stability of specific mRNA targets (Brown et al. 2001; Darnell et al. 2001; Zhang et al. 2001; Miyashiro et al. 2003; Zalfa et al. 2003; Antar et al. 2004). Although FMRP can be translocated to distal compartments of neurons and regulate local translation of target mRNA, we observed increased FMRP immunoreactivity only in the cell body of CA1 and DG cells after seizure activity and did not detect FMRP within the neurites or nucleus of these cells.

Several findings have pointed a role for both ionotropic (Croucher et al. 1995) and metabotropic (Wong et al. 1999) glutamate receptors in promoting seizure activity in various animal models of epilepsy. To test the involvement of glutamate receptors in the expedited development of kindling, we treated Fmr1 KO mice with the NMDA receptor antagonist MK801 or the mGluR5 antagonist MPEP. The loss of Fmr1 did not obviously influence the dynamics of kindling development under conditions of NMDA receptor inhibition. However, blocking mGluR5 dramatically slowed the rate of kindling development exclusively in Fmr1 KO mice. This suggests that enhanced seizure susceptibility in the Fmr1 KO mouse is at least partially mediated by abnormal activation of mGluRs-dependent signal transduction which is largely suppressed in WT mice. This is consistent with the former finding that Group I mGluR and FMRP function in a convergent way in regulating mRNA translation and subsequently regulate synaptic plasticity and neuronal excitability (Greenough et al. 2001; Bear et al. 2004; Chuang et al. 2005; Hou et al. 2006; Nosyreva and Huber 2006; Pan and Broadie 2007; Pan et al. 2008).

The development of kindling in Fmr1 KO mice have 3 interesting characteristics: normal EST, accelerated development of kindling, and prolonged electrographic seizures. These features share strong similarity to those observed in α2A adrenergic receptor (α2A-AR) null mice in which the number of stimulations required to achieve full kindling is reduced by about half and the duration of AD is nearly doubled throughout the development of kindling compared with WT mice (Janumppalli et al. 1998). The α2A-AR is coupled to Gt/o subset of heterotrimeric G-proteins, which may function against the activation of group I mGluRs which interact with Gq proteins. It is possible that loss of FMRP and inactivation of α2A-AR may lead to epileptogenesis by allowing excessive activation of a common signaling cascade. On the other hand, it is possible that the 2 mutants display similar phenotype of accelerated kindling by employing 2 parallel cellular cascades.

The formation of new synaptic connections is an important process in epileptogenesis. MFS is a major hallmark of abnormal synapse formation in the epileptic brain (Margerison and Corsellis 1966; McNamara 1999). As FMRP has been shown to regulate the formation of new synaptic connections (Comery et al. 1997; Irwin et al. 2000; Galvez et al. 2003; Pan et al. 2004; Antar et al. 2006), we investigated MFS in naive and kindled Fmr1 KO mice. Whereas, the naive Fmr1 KO brain exhibited no obvious abnormalities in mossy fiber innervation, MFS was dramatically enhanced in the IML region of Fmr1 KO mice compared with WT mice 5 or 28 weeks after 22 daily stimulations (Fig. 7). This difference was not likely to be simply the result of enhanced seizure activity in the Fmr1 KO brain because WT mice could be stimulated an additional 8 times (thus 36% more stimulations than Fmr1 KO mice) and still show little evidence of increased MFS (data not show). Because no obvious neuronal cell death was detected in either the kindled WT or Fmr1 KO mice 24 h after the completion of kindling development, this enhanced axon reorganization indicates that FMRP may exert an important role in preventing activity-induced neuronal network reorganization in the adult brain.

It has long been known that the cellular and molecular changes that underlie kindling are opposed by refractory periods, but the nature of these changes are not clear (Mucha and Pinel 1977). The upregulation of FMRP expression by kindling-induced seizure as well as the accelerated development of kindling and pervasive seizure-evoked synaptogenesis in Fmr1 KO mice that we observed together support the idea that FMRP is an important intrinsic buffer against epileptogenesis. It is in this light that we predict that the enhanced susceptibility of patients with FXS to epilepsy is the direct consequence of the absence of a critical homeostatic factor that mitigates vulnerability to excessive neuronal excitation.

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Notes

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Address correspondence to Dr Zhi-Qi Xiong. Institute of Neuroscience, Chinese Academy of Sciences, 320 Yue Yang Road, Shanghai 200031, China. Email: xiongzhiqi@ion.ac.cn.

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