Cellular/Molecular

JIP1-Mediated JNK Activation Negatively Regulates Synaptic Plasticity and Spatial Memory

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The c-Jun N-terminal kinase (JNK) signal transduction pathway is implicated in learning and memory. Here, we examined the role of JNK activation mediated by the JNK-interacting protein 1 (JIP1) scaffold protein. We compared male wild-type mice with a mouse model harboring a point mutation in the *Jip1* gene that selectively blocks JIP1-mediated JNK activation. These male mutant mice exhibited increased NMDAR currents, increased NMDAR-mediated gene expression, and a lower threshold for induction of hippocampal long-term potentiation. The JIP1 mutant mice also displayed improved hippocampus-dependent spatial memory and enhanced associative fear conditioning. These results were confirmed using a second JIP1 mutant mouse model that suppresses JNK activity. Together, these observations establish that JIP1-mediated JNK activation contributes to the regulation of hippocampus-dependent, NMDAR-mediated synaptic plasticity and learning.

Key words: fear; JIP1; JNK; LTP; memory; plasticity

Significance Statement

The results of this study demonstrate that c-Jun N-terminal kinase (JNK) activation induced by the JNK-interacting protein 1 (JIP1) scaffold protein negatively regulates the threshold for induction of long-term synaptic plasticity through the NMDA-type glutamate receptor. This change in plasticity threshold influences learning. Indeed, mice with defects in JIP1-mediated JNK activation display enhanced memory in hippocampus-dependent tasks, such as contextual fear conditioning and Morris water maze, indicating that JIP1-JNK constrains spatial memory. This study identifies JIP1-mediated JNK activation as a novel molecular pathway that negatively regulates NMDAR-dependent synaptic plasticity and memory.

Introduction

Human genetic studies have demonstrated that mutations in genes underlying the cJun NH₂-terminal kinase (JNK) signaling

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pathway are associated with neuropsychiatric, neurological, and neurodevelopmental disorders, including schizophrenia (MAP2K7; Winchester et al., 2012), epilepsy (MAPK10; Shoichet et al., 2006), autism spectrum disorder (MAPK8IP2 and TAOK2; Weiss et al., 2008; Giza et al., 2010; de Anda et al., 2012), and learning disability (MAPK10; Baptista et al., 2008; Kunde et al., 2013). These observations suggest that the JNK pathway has an important normal function in the CNS. Indeed, recent studies using *C*. elegans (Inoue et al., 2013) and murine (Sherrin et al., 2011) experimental models showed that JNK deficiency results in enhanced memory. This is exemplified by the observations that JNK1-deficient mice exhibit enhanced associative learning, including contextual fear conditioning (Sherrin et al., 2010) and altered synaptic plasticity (Li et al., 2007). Complementary studies using pharmacological inhibition of JNK demonstrate increased long-term depression (LTD) and loss of depotentiation

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(Yang et al., 2011). Moreover, JNK activation has been implicated in stress-mediated inhibition of long-term potentiation (LTP; Curran et al., 2003; Wang et al., 2004). Mechanisms that contribute to JNK-regulated synaptic plasticity include NMDAR-stimulated JNK activation (Mukherjee et al., 1999), AMPAR internalization (Zhu et al., 2005), and synaptic recruitment of PSD95 (Kim et al., 2007). Collectively, these data indicate that JNK plays a key role in the regulation of synaptic plasticity.

Although progress toward understanding the role of JNK in neuronal signaling has been achieved, little is known about the mechanisms that regulate JNK during behavioral responses. Previous studies have implicated roles for scaffold proteins in the control of MAP kinase (MAPK) signaling cascades, including the JNK signaling pathway (Morrison and Davis, 2003). Indeed, the JNK-interacting protein 1 (JIP1) scaffold protein can assemble a functional JNK signaling pathway (Whitmarsh et al., 1998, 2001). JIP1 is required for JNK activation caused by specific stimuli, including metabolic and excitotoxic stress (Whitmarsh et al., 2001; Morel et al., 2010). However, JIP1 is not required for JNK activation caused by other stimuli, including inflammatory cytokines (Whitmarsh et al., 2001).

JIP1 is highly expressed in the brain (Dickens et al., 1997; Whitmarsh et al., 1998) and localizes to synapses (Pellet et al., 2000). Interestingly, mice with JIP1 deficiency exhibit increased NMDAR signaling (Kennedy et al., 2007), implicating JIP1-mediated JNK activation in the regulation of NMDAR activity. The purpose of this study was to examine the impact of JIP1-regulated JNK activation on neuronal function and behavior. JIP1 may contribute to multiple biological processes, including microtubule motor protein function and JNK signaling (Morrison and Davis, 2003). The interpretation of studies using JIP1 knock-out mice (Whitmarsh et al., 2001) is therefore complicated by the presence of defects in JIP1-mediated JNK activation and defects in other JIP1-mediated biochemical activities. Therefore, we examined the effect of a point mutation in JIP1 (Thr 103 Ala) that selectively prevents JIP1-mediated JNK activation (Morel et al., 2010). This block in JNK activation is accounted for by a required role of JIP1 phosphorylation on Thr 103 for dynamic association with upstream MAP3K components of the signaling cascade (Nihalani et al., 2003; Morel et al., 2010). Our results demonstrated that JIP1mediated JNK activation regulates hippocampus-dependent, NMDAR-linked synaptic plasticity and memory. This conclusion was confirmed using a second mouse model with point mutations that disrupt the JNK-binding site on JIP1, which also prevents JIP1-mediated JNK activation and alters hippocampusdependent learning. Collectively, the data demonstrate that the JIP1-JNK signaling axis negatively regulates synaptic plasticity and spatial memory, possibly functioning to constrain and/or shape learning and memory under specific contexts.

Materials and Methods

Mice. C57BL/6J mice (stock number 000664) were obtained from The Jackson Laboratory and were established as a colony in our facility. The *Jip1 Thr* ¹⁰³*Ala* (JIP1 ^{TA}) mice have been described previously (Morel et al., 2010). Mice with a defect in the JNK-binding domain (JBD) of JIP1 (replacement of Leu ¹⁶⁰-Asn ¹⁶¹-Leu ¹⁶² with Gly ¹⁶⁰-Arg ¹⁶¹-Gly ¹⁶²) were established by homologous recombination in embryonic stem (ES) cells using standard methods. The mutated allele is designated as $Jip1^{\Delta JBD}$. Briefly, a targeting vector was constructed that was designed to introduce point mutations in exon 3 of the Jip1 gene to create the ΔJBD mutation. TC1 ES cells (strain129svev; RRID:CVCL_M350) were electroporated with this vector and selected with 200 μ g/ml G418 (Thermo Fisher Scientific, catalog #10131035) and 2 μ M ganciclovir (Millipore, catalog #345700). ES cell clones without (genotype +/*NeoR-Jip1* ^{WT}) and with

(genotype +/NeoR-Jip1 $^{\Delta$ JBD}) the Δ JBD mutation in exon 3 were identified. These ES cells were injected into C57BL/6J blastocysts to create chimeric mice that were bred to obtain germline transmission of the targeted Jip1 allele. The floxed Neo R cassette was excised using Cre recombinase. The full characterization of these mice has been described previously (Kant et al., 2017). All mice used in this study were back-crossed (10 generations) to the C57BL/6J strain (The Jackson Laboratory). All studies were performed using male mice. The mice were housed in a facility accredited by the American Association for Laboratory Animal Care. The Institutional Animal Care and Use Committee of the University of Massachusetts, University of Hawaii, and Morehouse School of Medicine approved all studies using animals.

Primary hippocampal neurons. Embryonic day 16.5 mouse embryos were used for isolation of primary hippocampal neurons (Whitmarsh et al., 2001). Briefly, hippocampi were placed in ice-cold Hank's buffered saline solution containing 20 mm HEPES (pH 7.3; HBSS, Thermo Fisher Scientific, catalog #15630080) and digested with 1% trypsin (Thermo Fisher Scientific, catalog #17075029) in the presence of 1 mg/ml DNase I (Sigma-Aldrich, catalog #11284932001). The trypsin solution was removed and the minced tissue was triturated in 1.0 ml of HBSS-20 mm HEPES containing DNase I (1 mg/ml; Sigma-Aldrich, catalog #11284932001) and soybean trypsin inhibitor (0.5 mg/ml; Thermo Fisher Scientific, catalog #17075029) to obtain a single-cell suspension. Dissociated neurons were centrifuged (180 × g, 10 min, 4°C) through a cushion of 4% bovine serum albumin (BSA; Thermo Fisher Scientific, catalog #B14) in HBSS. Hippocampal neurons were seeded in poly-D-lysine/laminin-coated chamber slides (BD Biosciences, catalog #354687) in Neurobasal medium (Thermo Fisher Scientific, catalog #21103049) containing B27 supplement (Thermo Fisher Scientific, catalog #17504044), 1% glutamine (Thermo Fisher Scientific, catalog #21051024), and 1% penicillin/streptomycin (Thermo Fisher Scientific, catalog #15140122).

Gene expression studies were performed using hippocampal neurons cultured for 14 d *in vitro* (DIV). The neurons were treated without or with 100 μ M NMDA/10 μ M Glycine (Sigma-Aldrich, catalog #M3262, catalog #G5417) in complete Neurobasal media (Thermo Fisher Scientific, catalog #21103049). The expression of mRNA was measured by quantitative RT-PCR assays.

Immunofluorescence analysis was performed using hippocampal neurons cultured for 16 d in vitro (16 DIV). Two different procedures were used to prepare neurons for immunofluorescence analysis. First, permeabilized fixed neurons were prepared by incubation with 4% (w/v) paraformaldehyde (Thermo Fisher Scientific, catalog #28906) at room temperature (20 min), followed by incubation with 0.1% Triton X-100 in PBS (5 min; Thermo Fisher Scientific, catalog #85112) and then blocking buffer (1% BSA (Thermo Fisher Scientific, catalog #B14)), 2% normal goat serum (Thermo Fisher Scientific, catalog #31872) in PBS for 1 h. Second, nonpermeabilized fixed neurons were prepared by incubation with 4% (w/v) paraformaldehyde/4% (w/v) sucrose (Thermo Fisher Scientific, catalog #28906; Fisher Scientific, catalog #BP-2201) at room temperature (8 min), followed by incubation (1 h) in blocking buffer. Neurons prepared by both methods were incubated overnight with primary antibodies to GluN1 (1:100, Millipore, catalog #05-432, RRID: AB_390129) and β-Tubulin (1:500), Covance Research Products, catalog #PRB-433C-200, RRID:AB_291636) in blocking buffer and then washed. The primary antibodies were detected by incubation with antimouse or anti-rabbit Ig conjugated to Alexa Fluor 488 or 633 (1:200, Invitrogen, catalog #A-11094, RRID:AB_221544; Thermo Fisher Scientific, catalog #A21100, RRID:AB_10374307). DNA was detected by staining with DAPI (Vectashield with DAPI, Vector Laboratories, catalog #H-1200). Fluorescence was visualized using a Leica TCS SP2 confocal microscope equipped with a 405 nm diode laser. The mean fluorescence intensity was quantitated using ImageJ software (RRID:SCR_001775).

Preparation of synaptosomes. Hippocampi from 4 mice (age 8–12 weeks) were isolated and homogenized in Syn-PER buffer (Thermo Fisher Scientific, catalog #87793). Briefly, neuronal tissue was homogenized in Syn-PER reagent, centrifuged following manufacturer's instructions, and the resulting synaptosome pellet was suspended in Syn-PER reagent to yield $3-4 \,\mu g/\mu l$ of synaptic protein. The synaptosomes contain the complete presynaptic terminal, including mitochondria and syn-

aptic vesicles, and also the postsynaptic membrane and the postsynaptic density (Villasana et al., 2006).

RNA analysis. The expression of mRNA was examined by qRT-PCR analysis using a 7500 Fast real-time PCR machine. TaqMan assays were used to quantitate cFos (Mm00487425_m1), cJun (Mm00495062_s1), Bdnf (Mm00432069_m1), GluN1 (Grin1, Mm00433800_m1), GluN2A (Grin2a, Mm00433802_m1), and GluN2B (Grin2b, Mm00433820_m1; Thermo Fisher Scientific). The relative mRNA expression was normalized by measurement of Gapdh (4352339E-0904021; Thermo Fisher Scientific) in each sample using TaqMan assays.

Immunoblot analysis. Tissue extracts were prepared from snap-frozen brain regions from adult mice (8–12 weeks old) using Triton X-100 lysis buffer (20 mm Tris-pH 7.4, 1% Triton X-100, 10% glycerol, 137 mm NaCl, 2 mm EDTA, 25 mm b-glycerophosphate, 1 mm sodium orthovanadate, 1 mm PMSF, 10 mg/ml Leupeptin, and 10 mg/ml Aprotinin). Extracts (20-50 µg of protein) were examined by protein immunoblot analysis by probing with antibodies to pSer 63-cJun (1:1000, Cell Signaling Technologies, catalog #2361, RRID:AB_490908), JNK (1:1000, Cell Signaling Technologies, catalog #9252, RRID:AB_2250373), GAPDH (Cell Signaling Technologies, catalog #2118, RRID:AB_561053), GluN2B (1:1000, Cell Signaling Technologies, catalog #4212S, RRID:AB_2112463), SAP102 (1:1000, Cell Signaling Technologies, catalog #3730S, RRID: AB_2092180), pERK1/2 (1:1000, Cell Signaling Technologies, catalog #5683P, RRID:AB_10841299), pSer 133 CREB (1:1000, Cell Signaling Technologies, catalog #9198, RRID:AB_2561044) and CREB (1:1000, Cell Signaling Technologies, catalog #9197, RRID:AB_321277), ERK2 (1:1000, Santa Cruz Biotechnology, catalog #sc-81457, RRID: AB_1122619), JNK1/2 (Santa Cruz Biotechnology, catalog #sc-137019, RRID:AB_2140722), pJNK (Santa Cruz Biotechnology, catalog #sc-6254, RRID:AB_628232) and GluN2A (1:500, Millipore, catalog #07-632, RRID: AB_310837), pY 1472 GluN2B (1:1000, Millipore, catalog #AB5403, RRID: AB_11210694); GluN1 (Millipore, catalog #05-432, RRID:AB390129), GluA1 (1:2000, Millipore, catalog #04-855, RRID:AB_1977216), GluA2 (1:1000, Millipore, catalog #07-261, RRID:AB_2116167), Synapsin I (1:1000, Millipore, catalog #AB 1543, RRID:AB_11210367), β -Tubulin (1:5000, Covance Research Products, catalog #PRB-435P-100, RRID:AB_291637), PSD-95 (1:2000, Sigma-Aldrich, catalog #P246, RRID:AB_260911) and KIF17 (1:1000, Sigma-Aldrich, catalog #K3638, RRID:AB_477148). Immunocomplexes were detected by fluorescence using anti-mouse (1:5000, LI-COR Biosciences, catalog #926-32210, RRID:AB_621842) and anti-rabbit secondary IRDye antibodies (1:5000, LI-COR Biosciences, catalog #827-08365, RRID:10796098) and quantitated using a LI-COR imaging system.

Multiplexed ELISA. Quantitative analysis of pSer⁶³-cJun (Bio-Rad, catalog #171-V50003M), cJun (Bio-Rad, catalog #171-V60002M), pJNK (Bio-Rad, catalog #171-V50011M), JNK (Bio-Rad, catalog #171-V60007M), pERK (Bio-Rad, catalog #171-V50006M), and ERK1/2 (Bio-Rad, catalog #171-V60003M) was performed using Bio-Plex Pro Cell Signaling Reagent kit (Bio-Rad, catalog #171-304006M) and a Luminex 200 instrument (Millipore-Sigma-Aldrich).

Kainate-induced excitotoxicity. JIP1 WT and JIP1 TA mice (8-12 weeks old) were injected intraperitoneally with 30 mg/kg kainic acid (Tocris Bioscience, catalog #0222; Yang et al., 1997). At 2 h posttreatment, the mice were perfused with 4% paraformaldehyde. Brains were harvested and fixed for an additional 24 h in 4% paraformaldehyde and then dehydrated and embedded in paraffin. Coronal sections (5 μ m) were cut, rehydrated, and subjected to heat-induced antigen retrieval (Vector Laboratories, catalog #H-3301). Sections were blocked for 1 h at room temperature (1% BSA, 2% normal goat serum, 0.4% Triton X-100 in PBS) and incubated overnight with primary antibodies to pSer⁶³-cJun (1:100, Cell Signaling Technologies, catalog #2361, RRID:AB_490908), or cFos (1: 200, Cell Signaling Technologies, catalog #4384S, RRID:AB_10698737) or cJun (1:100, Santa Cruz Biotechnology, catalog #sc-1694, RRID: AB_631263). The primary antibodies were detected by incubation with anti-rabbit Ig conjugated to Alexa Fluor 488 (1:200, Thermo Fisher Scientific, catalog #A11008, RRID:AB_143165). DNA was detected by staining with DAPI (Vectashield with DAPI, Vector Laboratories, catalog #H-1200). Fluorescence was visualized using a Leica TCS SP2 confocal microscope equipped with a 405 nm diode laser.

Surgery and infusions. Mice were anesthetized with 1.2% Avertin (25 g 2, 2, 2-Tribromoethanol, Sigma-Aldrich, catalog #T4, 840-2, 15.5 ml of tert-amyl alcohol (2-methyl-2-butanol), Fisher Scientific, catalog #A730-1) and implanted bilaterally with 26 gauge guide cannulae (Plastics One) into the dorsal hippocampus (AP -1.5 mm, lateral 1 mm, depth 1.3 mm). Mice were allowed to recover for at least 7 d before behavioral experiments. Intrahippocampal infusions were made using custom 28 gauge injectors (Plastics One, catalog #C232I/SPC) that extended 1 mm beyond the tips of the guide cannulae. On the day of the experiment, bilateral injections were performed using an infusion pump (CMA/100; CMA/Microdialysis) at a constant rate of 0.5 μ l/min (final volume: 0.25 μl/side). The competitive NMDAR antagonist D,L-2-amino-5-phosphonovaleric acid (APV; 10 μg/ml; Sigma-Aldrich, catalog #A5282) was dissolved in aCSF (aCSF) containing the following (in mm): 130 NaCl, 3.5 KCl, 10 glucose, 1.25 NaH₂PO₄, 2.0 CaCl₂, 1.5 MgSO₄, and 24 NaHCO₃. Controls received equal volumes of aCSF infused at the same rate. All infusions were made 15 min before contextual fear conditioning.

Analysis of dendritic spine density and morphology. Golgi staining was performed using the FD Rapid Golgi Stain Kit (FD Neurotechnologies, catalog #PK401) following the manufacturer's guidelines. Coronal sections $(150 \,\mu\text{m})$ were obtained using a microtome (Leica VT1000S). Spines examined were apical (stratum radiatum) and basal (stratum oriens) dendrites of CA1 pyramidal neurons. CA1 pyramidal neurons were traced using a Zeiss Axioskop 2 Plus microscope with a 100× oil-immersion objective. Only pyramidal neurons that exhibited complete impregnation and not obscured by other neurons or artifacts were examined. Five neurons per animal were 3D reconstructed using NeuroLucida Software (Micro-BrightField; RRID:SCR_001775). At least three apical (>50 μ m from soma) and three basal (>30 μ m from soma) dendritic segments (>25 μm length) were quantified in each neuron. Spine densities were calculated as mean numbers of spines per 10 µm per dendrite per neuron in individual mice. Dendritic arborization was performed using Sholl analysis of the apical and basilar dendrites of these neurons. Briefly, a series of increasingly large concentric circles centered at the cell body and separated by 10- μ m-radius intervals were superimposed upon traces of apical and basilar dendrites; the number of dendritic intersections with each concentric circle was recorded. On the basis of morphology, spines were classified into the following categories: thin, mushroom, and stubby (Korobova and Svitkina, 2010).

Analysis of tissue sections. Paraformaldehyde-fixed brains were cryoprotected in 30% sucrose in 0.1 M PBS for 48 h at 4°C. Immunohistochemical analysis was performed on free-floating sections cut at 30 μ m on a cryostat. The sections were washed once with PBS and blocked with 5% goat serum (Vector Laboratories, catalog #S-1000) in PBST (PBS + 0.3% Triton X-100) for 1 h. The sections were then incubated overnight with primary antibodies to JIP1 (1:500, BD Biosciences, catalog #611890, RRID:AB_399370), anti-NeuN antibody (1:500, Millipore, catalog #MAB377, RRID:AB_2298772), MAP2 antibody (1: 500, Millipore, catalog #AB5622, RRID:AB_91939), GAD67 antibody (1: 5000, Millipore, catalog #MAB5406, RRID:AB_2278725), or GFAP antibody (1:150, Promega, catalog #G5601, RRID:AB_430855) at 4°C. The sections were washed in PBST and incubated (1 h) with anti-mouse Ig or anti-rabbit Ig conjugated to Alexa Fluor 488 (Thermo Fisher Scientific, catalog #S11223, RRID:AB_2336881) or Alexa Fluor 546 (Thermo Fisher Scientific, catalog #S11225, RRID:AB_2532130). Nuclei were stained using DAPI (Vectashield with DAPI, Vector Laboratories, catalog #H-1200). Images were obtained with a Zeiss Axio Imager 2 microscope at 10× and 20× magnification. The mean fluorescence intensity was quantitated using ImageJ (RRID:SCR_003070) software. Sections were also examined using Nissl stain (Thermo Fisher Scientific, catalog #N21479).

Fear conditioning and extinction. Context-dependent and tone-dependent fear conditioning were performed using a computer-controlled fear conditioning system (TSE Systems; Todorovic et al., 2007). The fear-conditioning experiment was performed in a Plexiglas cage ($36 \times 21 \times 20$ cm) within a fear-conditioning box. The training (conditioning) consisted of a single trial. The mouse was exposed to the conditioning context (180 s), followed by a tone [conditioned stimulus (CS), 30 s, 10 kHz, 75 dB SPL, pulsed 5 Hz]. After termination of the tone, a foot shock [unconditioned

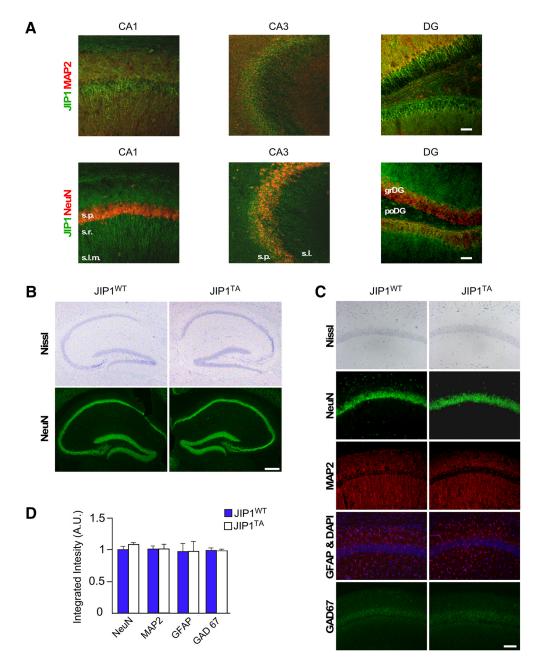


Figure 1. Analysis of JIP1 expression in the hippocampus. **A**, Fluorescent immunohistochemistry of the CA1, CA3, and DG regions of the hippocampus indicates that JIP1 (green) predominantly colocalizes with MAP2 (red) in wild-type brain sections (*top*), *suggesting* relative enrichment of the JIP1 protein in neuronal processes. Colocalization between JIP1 (green) and neuron-specific nuclear protein (NeuN; red) is sparse in the hippocampal subfields (bottom). CA1, Cornu ammonis 1; CA3, cornu ammonis 3; grDG, granular layer of DG; poDG, polymorphic layer of DG; s.l., stratum lucidem; s.l.m., stratum lacunosum moleculare; s.p., stratum pyramidale; s.r., stratum radiatum. Scale bar, 50 μ m. **B**, Nissl stain and NeuN stain of JIP1 ^{MT} and JIP1 ^{TA} coronal hippocampal sections. Scale bar, 200 μ m. **C**, **D**, Pyramidal cells of the CA1 region of JIP1 ^{WT} and JIP1 ^{TA} were stained with Nissl, NeuN, the dendritic marker MAP2, the astrocytic marker GFAP, and the inhibitory GABAergic marker GAD67 (**C**). The staining was quantitated (**D**) (mean \pm SEM; n=4; p>0.05; Student's t test). Scale bar, 50 μ m.

stimulus (US), 0.8 mA or 0.4 mA, 2 s, constant current] was delivered through a stainless steel grid floor. Under these conditions, the context served as background stimulus. Background contextual fear conditioning, but not foreground contextual fear conditioning, in which the tone is omitted during training, has been shown to involve the hippocampus (Phillips and LeDoux, 1992). A loudspeaker provided constant background noise. Contextual memory was tested in the fear-conditioning box for 180 s without CS or US presentation (with background noise) 24 h after contextual fear conditioning. The context-dependent extinction trials were performed at 24 h intervals and consisted of nonreinforced 3 min exposures (absence of a foot shock) to the same context. For fear response extinction, the aforementioned protocol was used to acquire contextual fear memory. Tone-dependent memory test was per-

formed in a novel context (context 2) 24 h after cued fear conditioning. Context 2 represented an identically sized cage with a plain floor in a light-surrounding environment (350–500 lux) outside the fear-conditioning box. No background noise was provided in context 2. During tone-dependent memory test, a 180 s pause without stimulation (pre-CS phase) preceded a 180 s period of auditory stimulation. Freezing, defined as a lack of movement except for respiration, was recorded every 10 s by a trained observer for a total of 18 sampling intervals. The mean number of observations indicating freezing was expressed as a percentage of the total number of observations. The exploration of the fear-conditioning box during the training and activity burst produced by electric foot shock were automatically detected by an infrared beam system and analyzed using TSE Systems software.

Morris water maze (MWM). The MWM paradigm (Morris et al., 1982) was performed in a circular tank (diameter 180 cm; height 75 cm) located in a room with various distal cues. The tank was filled with water (40 cm depth) maintained at 23°C, which was made opaque by the addition of a nontoxic white paint. Inside the pool was a removable, circular (12 cm in diameter) Plexiglas platform 0.5 cm below the surface of the water. On the first 2 d, each mouse received visible platform training that consisted of 4 consecutive trials of climbing onto the visible platform (with a black plastic brick placed above it) until each subject was able to climb without help. For the hidden platform task, the mice were given four consecutive trials per day starting from four different pseudorandomized start locations with a 15 min intertrial interval. Mice were allowed to search for the hidden platform for 60 s. If the mice did not find the platform within 60 s, they were guided to it. Mice were allowed to rest on the platform for 15 s after each trial. The hidden platform task was composed of two phases: (1) 10 d (acquisition phase days 3-13) with a hidden platform located in the center of the target quadrant and (2) reversal phase (day 14) with a hidden platform located in the center of the quadrant opposite to the original target quadrant. Reversal platform training was conducted without changing any distal visual cues. Probe trials in which the escape platform was removed from the pool were conducted on days 10 (target quadrant), 13 (target quadrant), and 15 (opposite quadrant). During the memory test (probe trials), the platform was removed from the tank and the mice were allowed to swim in the maze for 60 s. The swimming path of the mice was recorded by a video camera and analyzed with the computer-based tracking software Videomot 2 (TSE Systems, RRID: SCR_014334). The percentage of swim distance spent in the platform quadrant and the latency to find the platform were analyzed.

Rotarod test. Motor coordination and skill learning were assessed using an accelerating Rotarod (Stoelting). Starting speed for the Rotarod began at 4 rpm and increased to 40 rpm over a 5 min period. Mice were tested 4 times daily for 2 consecutive days with an intertrial interval of 1 h between tests. The latency to fall off the rod was measured for each trial.

Elevated plus maze test. The elevated plus maze test for anxiety-related behaviors was performed as described previously (Todorovic et al., 2007). Briefly, mice were placed in the center platform of the elevated plus maze and allowed to explore for 5 min. Animal behavior was recorded by a video camera connected to a PC and analyzed by videotracking software (VideoMot 2, TSE Systems, RRID:SCR_014334). The percentage of time spent in the open and closed arms were recorded. Shift of preference from the open to the closed arms was interpreted as an increase of anxiety-like behavior. Locomotor activity was determined with this test by the distance traveled.

Open-field test. General exploratory activity and anxiety were assessed in an open-field test. Mice were placed in the center of an open-field apparatus (50 \times 50 cm) protected with 10-cm-high opaque walls and allowed to explore for 5 min. The field was divided into 16 equal squares (12.5 \times 12.5 cm) consisting of 12 outer squares and four inner squares. Animal behavior was recorded by a video camera connected to a PC and analyzed by video-tracking software (VideoMot 2, TSE Systems, RRID: SCR_014334). The amount of time spent in the inner and outer squares and the total distance traveled was measured.

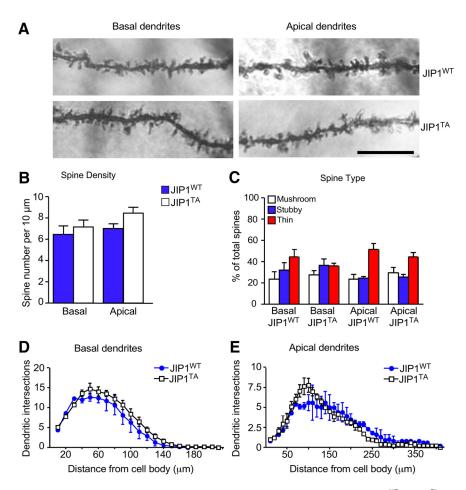


Figure 2. Neuronal spine density and dendritic arborization of CA1 pyramidal neurons are similar in JIP1 ^{MT} and JIP1 ^{TA} mice. **A**, Representative images of apical and basal dendrites spine morphology in JIP1 ^{TA} mice and JIP1 ^{WT} littermates. Scale bar, 10 μ m. **B**, Quantitation of basal and apical dendritic spine density (mean \pm SEM; n=5 slices from 5 mice per genotype; p>0.05, Student's t test). **C**, Quantitation of different spine types in basal and apical dendrites (mean \pm SEM, n=5 slices from 5 mice per genotype; p>0.05, Student's t test). **D**, **E**, Sholl analysis of dendritic arborization of CA1 pyramidal neurons. Values on the x-axis represent increasing distance from the soma of the pyramidal cells. Basal and apical dendrites of pyramidal cells from n=5 slices from 5 mice per genotype were examined (mean \pm SEM; p>0.05, Student's t test).

Acoustic startle and prepulse inhibition (PPI). Acoustic startle and PPI test were performed as described previously (Pitts et al., 2012). Mice were placed in the startle chamber (Responder-X; Columbus Instruments) and allowed a 5 min acclimation period with the background noise (70 dB) continuously present. After the acclimation period, two blocks of trials were administered to assess the acoustic startle response and PPI, respectively. The first block of trials consisted of eight sets of four types of trials that were distributed randomly. Startle stimuli (40 ms) of varying intensities were administered, with an interstimulus interval of 15 s. The stimulus intensities were 80, 90, 100, and 110 dB. Baseline activity was assessed by a set of no-stimulus trials. The startle amplitude was defined as the peak response during a 100 ms sampling window beginning with the onset of the startle stimulus. Mean startle amplitudes were derived by subtracting the average startle amplitudes of stimulus intensities used (80-110 dB) from the no-stimulus trial (70 dB). The second block of trials consisted of eight sets of five trial types, distributed randomly and separated by 20 s interstimulus intervals. The trial types were as follows: (1) no-stimulus/background noise (70 dB); (2) 40 ms, 110 dB startle alone; and (3-5) 110 dB startle preceded 100 ms by one of three 20 ms prepulses at the following intensities: 74, 80, or 86 dB. The startle amplitude for each subject at each of the different prepulse intensities was calculated using the following formula: PPI = $100 - 100 \times$ (response amplitude for prepulse stimulus paired with startle stimulus/response amplitude for startle stimulus alone).

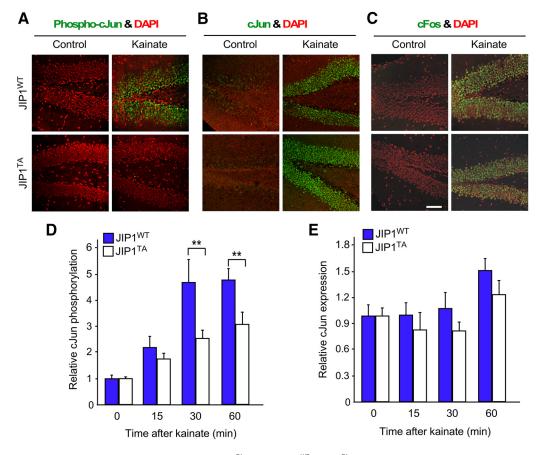


Figure 3. JIP1-dependent JNK activation in the hippocampus is suppressed in JIP1^{TA} mice. A-C, JIP1^{WT} and JIP1^{TA} mice were treated by systemic injection of kainate. At 2 h after treatment, sections of the brain were prepared and stained (green) with antibodies to pSer ⁶³ clun (A), clun (B), or cFos (C). DNA was stained with DAPI (red). Representative sections of the DG of the hippocampus are presented. Scale bar, 75 μ m. D, E, Extracts prepared from the hippocampus of JIP1^{WT} and JIP1^{TA} mice treated with kainate (0 – 60 min) were examined by multiplexed ELISA to measure the amount of pSer ⁶³-clun (D) and clun (E) normalized to the amount of JNK. Data are shown as mean \pm SEM. n = 5; **P < 0.01, two-way ANOVA followed by Bonferroni's *post hoc* test.

Electrophysiology. Extracellular recordings were performed as described previously (Lawrence et al., 2014). Hippocampi of wild-type or mutant mice (8-12 weeks old) were rapidly removed and briefly chilled in icecold aCSF containing the following (in mm): 130 NaCl, 3.5 KCl, 10 glucose, 1.25 NaH₂PO₄, 2.0 CaCl₂, 1.5 MgSO₄, and 24 NaHCO₃ (equilibrated with 95% $O_2/5\%$ CO_2 , pH 7.4). Transverse slices 350 μ m thick were prepared with a Vibratome (Leica, VT1200S) and maintained at least 1 h in a holding chamber containing aCSF. The slices were then transferred to a recording chamber and perfused (3 ml/min) with aCSF at 32°C. CA1 field EPSPs (fEPSPs) were recorded with a glass electrode filled with 3 M NaCl (resistance 1–1.5 M Ω) by stimulating the Schaffer collateral fibers through a bipolar stimulating electrode. The slope of the initial rising phase (20-60% of the peak amplitude) of the fEPSP was used as a measure of the postsynaptic response. Basal synaptic neurotransmission was studied by plotting stimulus strength or fiber volley against fEPSP slope to generate input/output (I/O) relationships. Pairedpulse facilitation was defined as the second fEPSP slope divided by the first at various interstimulus intervals (10, 50, 90, 130, 170, 210, and 250 ms). For the LTP and LTD measurements, a minimum of 20 min of baseline stimulation (0.05 Hz) was recorded every minute at an intensity that evoked a response 40% of the maximum response. The strong tetanic LTP induction protocol consisted of two 100 Hz tetani (1 s each), with an interval of 20 s between tetani. The weak LTP induction protocol consisted of 900 pulses given at 10 Hz. To induce NMDAR-dependent LTD, 1 Hz and 0.5 Hz single pulse stimuli were delivered for 15 and 30 min, respectively (900 stimuli). To induce mGluR-dependent LTD, slices were incubated with 100 μM (S)-3,5- dihydroxyphenylglycine (DHPG; Tocris Bioscience, catalog #0805) for 5 min. For the depotentiation study, LTP was evoked by high-frequency stimulation (HFS) with two trains of 100 Hz (20 s between tetani). Ten minutes after LTP induction, depotentiation was induced by low-frequency stimulation (LFS) consisting of 900 pulses delivered at 1 Hz for 15 min. fEPSP responses were recorded using a computer with WinLTP data acquisition software (WinLTP). For pharmacological studies, JNK-in-8 (Selleckchem, catalog #S 4901; 6 μ M in aCSF with 10% DMSO) or vehicle was applied for 20 min before testing and were maintained throughout the recording period.

For EPSC measurements, mice (6-10 weeks old) were anesthetized with isoflurane (Sigma-Aldrich, catalog #792632) and perfused with icecold sucrose solution containing the following (in mm): 200 sucrose, 20 glucose, 5 KCl, 1.2 NaH₂PO₄, 25 NaHCO₃, 0.5 CaCl₂, 7 MgCl₂, 1.3 ascorbic acid, and 2.4 sodium pyruvate. Transverse sections (350 μ m) of the hippocampus were made and then incubated in aCSF at 34°C (20 min) before storing at room temperature. The aCSF for these experiments was composed of the following (in mm): 125 NaCl, 10 glucose, 2.8 KCl, 1 NaH₂PO₄, 26 NaHCO₃, 2 CaCl₂, and 1.5 MgSO₄. Whole-cell voltage-clamp recordings were done at 32°C using an EPC-10 amplifier and Patchmaster acquisition software (HEKA Instruments, RRID: SCR_000034). Data acquisition and analysis was performed on a MacPro computer (Apple). Bicuculline (30 µm; Tocris Bioscience, catalog #0130) and CGP-55845 (1 µm; Tocris Bioscience, catalog #1248) were added to the aCSF to block GABA_A and GABA_B responses, respectively. Both sucrose and aCSF solutions were continually bubbled with 95% O₂/5% CO₂. Patch pipettes were filled with an intracellular solution containing the following (in mm): 130 Cs-gluconate, 5 NaCl, 0.5 CaCl₂, 2 MgCl₂, 10 HEPES, 5 Cs₄BAPTA, 2 MgATP, and 0.3 Na₂GTP. The intracellular solution was titrated to pH 7.2 and 280 mOsm. CA1 principal cells were held at either -70 or +50 mV to measure the AMPA and NMDAR components of EPSCs, respectively. Stimulation of the Schaffer collaterals synapsing onto the CA1 cells was performed using an A360 stimulus

isolation unit (World Precision Instruments) with a monopolar platinum iridium electrode placed in the stratum radiatum $\sim 350~\mu m$ from the cell body. The stimulation intensity was adjusted to be $\sim 75\%$ of the threshold for firing an action potential. Peak amplitudes were analyzed using Igor Pro version 6.3 (RRID:SCR000325).

Experimental design and statistical analysis. Data were analyzed statistically using StatView (SAS Institute) and GraphPad Prism 6 software. Student's t test was used for comparing two conditions and ANOVA was used with Bonferroni post hoc test for comparing more than two conditions. All data are expressed as means \pm SEM. The accepted level of significance was $p \leq 0.05$, indicated by a single asterisk in the figures; p-values ≤ 0.01 are indicated by double asterisks and p values ≤ 0.001 by triple asterisks.

Results

The aim of this study was to examine the impact of JIP1-regulated JNK activation on neuronal function and behavior. To accomplish this aim, we generated a novel mouse model harboring a T103A point mutation in the *Jip1* gene (also known as *Mapk8ip1*). These mutant *Mapk8ip1* T103A mice (JIP1 TA) exhibit a profound defect in JIP1-mediated JNK activation compared with control JIP1 TA animals. We therefore used this mutant mouse to determine the contribution of JIP1-mediated JNK activation to NMDA-dependent receptor signaling, memory, and synaptic plasticity.

JIP1 expression in hippocampal neurons

Our initial studies were designed to examine the expression of JIP1 protein in the hippocampus. Immunohistochemical stain-

ing of tissue sections demonstrated that JIP1 is expressed in the CA1, CA3, and the dentate gyrus (DG) regions of the hippocampus (Fig. 1A). JIP1 was primarily found on dendrites (colocalization with MAP2) and to a lesser extent on cell bodies (colocalization with NeuN) of the CA1/CA3 pyramidal and DG granule neurons (Fig. 1A).

We next examined the hippocampal architecture of JIP1 ^{TA} and JIP1 ^{WT} mice. Qualitative analysis of Nissl-stained (Fig. 1*B*, *C*) and Golgi-stained (Fig. 2*A*) coronal sections did not reveal differences in hippocampus morphology between JIP1 ^{TA} and JIP1 ^{WT} mice. Moreover, analysis of basal and apical dendrites from Golgi-stained CA1 pyramidal neurons revealed no significant differences in dendritic branching, spine density, or spine type (Fig. 2*A*–*E*). In addition, the intensity and distribution of the neuronal marker NeuN ($t_{(6)} = 0.89$, p = 0.41; two-tailed unpaired Student's t test), the dendritic marker MAP2 ($t_{(6)} = 0.71$, p = 0.51; two-tailed unpaired Student's t test), the GABAergic interneuron marker GAD67 ($t_{(6)} = 0.27$, p = 0.79; two-tailed unpaired Student's t test), and the glial marker GFAP ($t_{(6)} = 0.44$, t = 0.67; two-tailed unpaired Student's t test) were similar in JIP1 ^{TA} and JIP1 ^{WT} mice (Fig. 1*B*–*D*). Collectively, these data

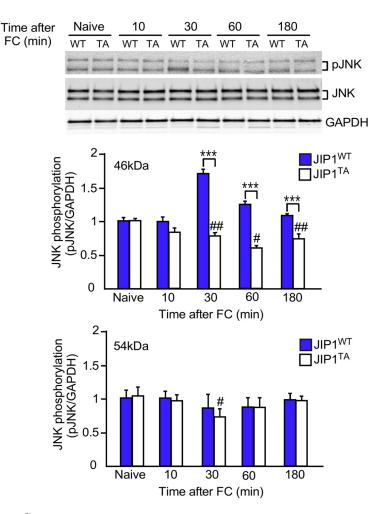


Figure 4. JIP1^{TA} mice exhibit reduced JNK activation in the dorsal hippocampus after contextual fear conditioning. Dorsal hippocampal tissue was isolated from naive mice and from mice at different times after contextual fear conditioning (FC) and examined by immunoblot analysis by probing with antibodies to phospho-JNK, JNK, and GAPDH. The amount of 46 and 54 kDa phospho-JNK was quantitated and normalized to the amount of JNK in each sample. Data are shown as mean \pm SEM. n=5; ****p<0.001, for JIP1 TA compared with JIP1 WT mice; #p<0.01, #p<0.001 compared with the naive control, two-way ANOVA, followed by Bonferroni's post hoc test.

indicate that the Thr 103Ala mutation in JIP1 did not cause detectable changes in basic hippocampus morphology.

JIP1-mediated JNK activation in the hippocampus

We have demonstrated previously that JIP1 TA mice exhibit defects in metabolic stress-induced JNK activation in adipose tissue (Morel et al., 2010). To test whether the Thr 103 Ala mutation in JIP1 also caused defects in JNK activation in neural tissue, we examined a model of JIP1-dependent JNK activation (kainate excitotoxicity) in the hippocampus. Examination of the DG after short-term exposure to kainate caused JIP1-dependent JNK activation and phosphorylation of the JNK substrate cJun and JIP1independent increased expression of cJun and cFos (Whitmarsh et al., 2001). We found that kainate caused a similar increase in cJun and cFos expression in JIP1 TA and JIP1 WT mice (Fig. 3 B, C). In contrast, kainate caused increased cJun phosphorylation in JIP1 WT but not JIP1 TA mice (Fig. 3A). These data demonstrate that the Thr ¹⁰³Ala mutation suppresses JIP1-mediated JNK activation in the hippocampus. Time course analysis confirmed that cJun phosphorylation in the hippocampus of kainate-stimulated JIP1 WT mice was strongly suppressed in JIP1 TA mice after exposure to kainate (two-way ANOVA: genotype $F_{(1,32)} = 15.11$, p <0.001; time $F_{(3,32)} = 34.26$, p < 0.001; genotype × time $F_{(3,32)} =$

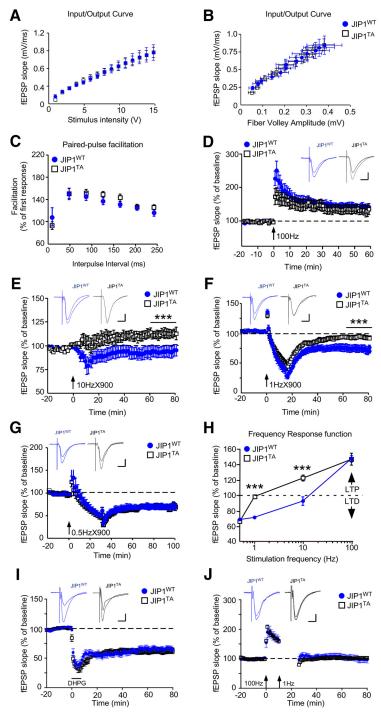


Figure 5. The threshold for LTP induction is reduced in JIP1 TA mice. A, B, B, as al synaptic transmission at Schaffer collateral—CA1 synapses, as assessed by measuring the fEPSP I/O relationship (A) and the fEPSP slope to fiber volley relationship (B), was similar in JIP1 TA slices (n=16 slices, 13 mice) compared with slices obtained from JIP1 TH littermates (n=16 slices, 12 mice). No statistically significant differences were found (p>0.05, two-way repeated-measures ANOVA). C, fEPSPs from JIP1 TA (n=16 slices, 13 mice) and JIP1 TA (n=16 slices, 12 mice) slices exhibited similar paired pulse facilitation. No statistically significant differences were found (p>0.05, two-way repeated-measures ANOVA). D, HFS LTP was induced by two trains of 100 Hz stimulation (separated by a 20 s interval) to the Schaffer collaterals in slices from JIP1 TA and JIP1 TA mice (n=10 slices, 8 mice/genotype). Stimulation was delivered at time 0 (arrow). No statistically significant differences were found (p>0.05, Student's t test). E, An intermediate stimulation LTP protocol involved 900 pulses of 10 Hz stimuli delivered at time 0. LTP induced at intermediate frequencies was significantly facilitated in slices taken from JIP1 TA mice when compared with JIP1 TH controls (n=10 slices, 9 mice/genotype). Statistically significant differences are indicated (***p<0.001, Student's t test). E, LTD induced by LFS (1 Hz, 900 pulses, 0 –15 min time) was significantly reduced in JIP1 TA slices compared with JIP1 TH slices (n=10 slices, 10 mice/genotype). Statistically significant differences are indicated (***p<0.0001, Student's t test). E, LTD induced by 0.5 Hz stimulation (0.5 Hz, 900 pulses, 0 –30 min time) was similar in JIP1 TA and JIP1 TA slices (n=10 slices, 11 mice/genotype). No statistically significant differences were found (p>0.05, Student's t test). E, Frequency—response function in JIP1 TA and JIP1 TA and JIP1 TA and JIP1 TA and JIP1 TA slices (t=10 slices). The statistica

5.05, p = 0.005; Fig. 3D,E). These data extend our previous finding that JIP1^{TA} mice exhibit defects in JIP1-dependent JNK activation in adipose tissue (Morel et al., 2010) to demonstrate that JIP1^{TA} mice also exhibit profound defects in JIP1-mediated JNK activation in the hippocampus.

To test how JIP1-mediated JNK activation contributes to a physiologically relevant response, we examined the effect of contextual fear conditioning, which was shown previously to cause transient JNK activation in the dorsal hippocampus (Sherrin et al., 2010). Immunoblot analysis using antibodies to JNK and phospho-JNK was performed to assess changes in JNK activation in dorsal hippocampal extracts from JIP1 WT and JIP1 TA mice that were subjected to single-trial contextual fear conditioning. Extracts were prepared at various times after the single trial to correspond to the consolidation phase of contextual fear (Igaz et al., 2002). JNK immunoblot analysis detects the 46 kDa (JNK1 α 1 and JNK1 β 1) and 54 kDa isoforms (JNK2 α 2, JNK2 β 2, and JNK3 α 2) in brain (Davis, 2000). We found that contextual fear conditioning caused an increase in 46 kDa phospho-JNK in JIP1 $^{\mathrm{WT}}$ mice at 30 min (176 \pm 12% of naive control mice; Bonferroni post hoc test, p < 0.001) and 60 min (127 \pm 8% of naive; Bonferroni *post hoc* test, p = 0.09; Fig. 4). In contrast, we found that JIP1 TA mice exhibited decreased 46 kDa phospho-JNK at 30 min (75 \pm 11% of naive; Bonferroni post hoc test, p = 0.011), 60 min (53 \pm 7% of naive; p < 0.001) and 180 min (79 \pm 6% of naive; p = 0.009) after contextual

mice. The percentage change in synaptic strength from baseline in JIP1 TA and JIP1 WT mice at 50 – 60 min after stimulation at the indicated frequency is presented. Data are shown as mean \pm SEM. Magnitudes of LTP/LTD were calculated as the ratio of the average fEPSPs between 50 and 60 min and average baseline fEPSPs between $-20 \, \text{min}$ and $0 \, \text{min}$. Statistically significant differences are indicated (***p < 0.001, Student's t test). I. mGluR-dependent LTP in hippocampal slices from JIP1 TA and JIP1 WT mice, mGluR-LTD was induced by incubation of JIP1 ^{TA} and JIP1 ^{WT} slices with DHPG (100 μ M) for 5 min (n = 10 slices, 10 mice/genotype). Basal fEPSPs were recorded before LTD induction with DHPG. No statistically significant differences were found (p > 0.05, Student's t test). J, Depotentiation is not affected in JIP1 TA mice. HFS (100 Hz twice for 1 s with 20 s interval) followed by 1 Hz (15 min) stimulation 10 min later to the Schaffer collaterals produced similar depotentiation in slices from JIP1 TA and JIP1 WT mice (n = 10 slices, 10 mice/genotype). No statistically significant differences were found (p > 0.05, Student's t test). The insets in **D**-**J** show representative fEPSP responses obtained before and after LTP, LTD and depotentiation inducing stimuli. Calibration: 0.2 mV/10 ms.

←

fear conditioning (two-way ANOVA: genotype $F_{(1,50)}=15.11,\ p<0.001;$ time $F_{(4,50)}=10.42,\ p<0.001;$ genotype \times time $F_{(4,50)}=22.11,\ p<0.001;$ Fig. 4). These data demonstrate that the JIP1 Thr ¹⁰³Ala mutation suppresses JNK activation caused by contextual fear conditioning.

JIP1-mediated JNK activation suppresses NMDA-dependent synaptic plasticity

JNK activation is associated with mechanistic changes in synaptic plasticity in the hippocampus (Wang et al., 2004; Chen et al., 2005; Zhu et al., 2005; Li et al., 2007; Yang et al., 2011). Therefore, synaptic plasticity may be altered by JIP1-dependent JNK activation in the context of fear conditioning (Fig. 4). We therefore examined synaptic transmission in JIP1 WT and JIP1 TA mice.

Basal synaptic transmission at Schaffer collateral—CA1 synapses in JIP1 WT and JIP1 TA mice was characterized through the I/O relationship of fEPSPs. Analysis revealed that the fEPSP slopes, plotted against the stimulation strength, were comparable in slices from JIP1 WT and JIP1 TA mice (two-way repeated-measures ANOVA: genotype $F_{(1,420)}=0.23, p=0.63$; amplitude $F_{(14,442)}=79.27, p<0.001$; genotype × amplitude $F_{(14,420)}=0.35, p=0.63$; Fig. 5A). Similarly, when the fEPSP slopes were plotted as a function of increasing fiber volley amplitudes, no differ-

ences between JIP1 WT and JIP1 TA mice were detected (Fig. 5*B*). Moreover, PPF, indicative of presynaptic plasticity (Zucker and Regehr, 2002), at Schaffer collateral–CA1 synapses from JIP1 TA mice was similar to JIP1 WT mice at several interpulse intervals (two-way repeated-measures ANOVA: genotype $F_{(1,223)} = 0.08$, p = 0.78; interval $F_{(6,223)} = 13.56$, p < 0.001; genotype × interval $F_{(8,208)} = 0.15$, p = 0.98; Fig. 5*C*). These data demonstrate similar basal synaptic transmission in the hippocampi of JIP1 TA and JIP1 WT mice.

To determine whether the JIP1 Thr 103 Ala mutation affects NMDAR-dependent forms of synaptic plasticity (Citri and Malenka, 2008), LTP was measured using a tetanic HFS protocol (2 trains of 1 s 100 Hz, separated by 20 s). Similar potentiation was produced in both JIP1 WT and JIP1 TA mice (Fig. 5*D*; fEPSPs were potentiated to 145 \pm 7% for JIP1 WT and 146 \pm 4% for JIP1 TĀ; $t_{(18)} = 0.12$, p = 0.89; two-tailed unpaired Student's ttest), indicating that the ability to induce LTP with strong stimulation was unimpaired in JIP1 TA mice. We next investigated whether the threshold stimulation for induction of LTP (Hu et al., 2007) differed between JIP1 WT and JIP1 TA mice. A 900 pulse train of 10 Hz stimulation resulted in a modest, but detectable, degree of LTP in slices obtained from JIP1 $^{\text{TA}}$ mice (Fig. 5E; 122 \pm 4% of baseline at 50-60 min after LTP induction), though with minimal posttetanic potentiation, whereas fEPSPs in JIP1 WT slices were not potentiated (Fig. 5E; 92.1 \pm 5.5% of baseline; $t_{(18)} =$ 7.61, p < 0.001; two-tailed unpaired Student's t test), with a trend toward depression. These data demonstrate that LTP could be induced at lower frequencies in JIP1 TA mice compared with JIP1 WT mice.

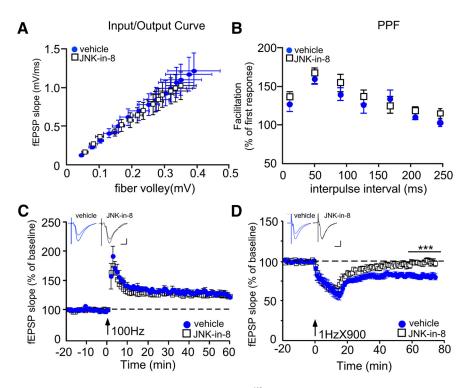


Figure 6. Inhibition of JNK signaling mimics the effect of JIP1 (Thr 103 Ala) mutation on NMDAR-dependent LTD. *A, B, JNK-in-8* treatment did not affect baseline synaptic transmission or PPF in wild-type slices. I/O curves, as assessed by the fEPSP slope to fiber volley relationship, were similar in vehicle-treated slices (n=12 slices) compared with wild-type slices treated with JNK-in-8 (n=12 slices; *A*). fEPSPs from vehicle-treated slices (n=12 slices) and JNK-in-8 treated (n=12 slices) slices exhibited similar PPF (\boldsymbol{B} ; p>0.05, two-way repeated-measures ANOVA). *C,* HFS-LTP was induced with a pair of 100 Hz tetani in the presence of either vehicle (n=12) or 6 μ M JNK-in-8 (n=12). LTP was unaffected by JNK inhibition. (p>0.05, Student's t test). *D,* LFS-LTD was induced (1 Hz, 900 pulses) in the presence of either vehicle (n=10) or 6 μ M JNK-in-8 (n=10). LTD was impaired by JNK inhibition (****p<0.001, Student's t test). The insets in *C* and *D* show representative fEPSP responses obtained before and after LTP/LTD-inducing stimuli. Calibration: 0.2 mV/10 ms.

We also examined the induction of NMDAR-dependent LTD (Collingridge et al., 2010) in JIP1 WT and JIP1 TA mice. We found that LFS consisting of 900 single pulses at 1 Hz for 15 min generated robust LTD in JIP1 WT mice, but LTD was absent in JIP1 TA mice (Fig. 5F; JIP1 WT mice, 72 \pm 2%; JIP1 TA mice, 98 \pm 3%; p < 0.01; $t_{(18)} = 7.05$, p < 0.001; two-tailed unpaired Student's t test). In view of the similar posttetanic depression (initial dip) between JIP1 WT and JIP1 TA brain slices, we determined whether JIP1 TA mice exhibit other forms of LTD. We therefore induced LTD by delivering the same number of single pulses (900) at 0.5 Hz for 30 min (Fig. 5G). This form of LTD also relies on NMDARs (Dudek and Bear, 1992), but depends to a greater degree on release of calcium from intracellular stores (Nakano et al., 2004). This LTD protocol produced equivalent depression in both JIP1 TA and JIP1 WT mice (Fig. 5*G*; JIP1 WT, 68 \pm 3%; JIP1 TA, 66 \pm 2%; $t_{(26)}$ = 0.55, p = 0.58; two-tailed unpaired Student's t test). These data indicate that disruption of JIP1-mediated JNK activation reduces the optimal frequency stimulation for LTD while enabling LTP at lower stimulation frequencies at Schaffer collateral-CA1 synapses (Fig. 5H).

Two different forms of LTD coexist at Schaffer collateral–CA1 synapses in the hippocampus, one dependent on NMDARs and another dependent on the activity of metabotropic glutamate receptors (mGluRs), specifically mGluR5 (Huber et al., 2001). Therefore, we next examined a possible effect of the JIP1 TA mutation on mGluR-dependent LTD. We found that a 5 min bath application of 100 μ M DHPG (a group I mGluR agonist) to acute slices resulted in a persistent depression of Schaffer

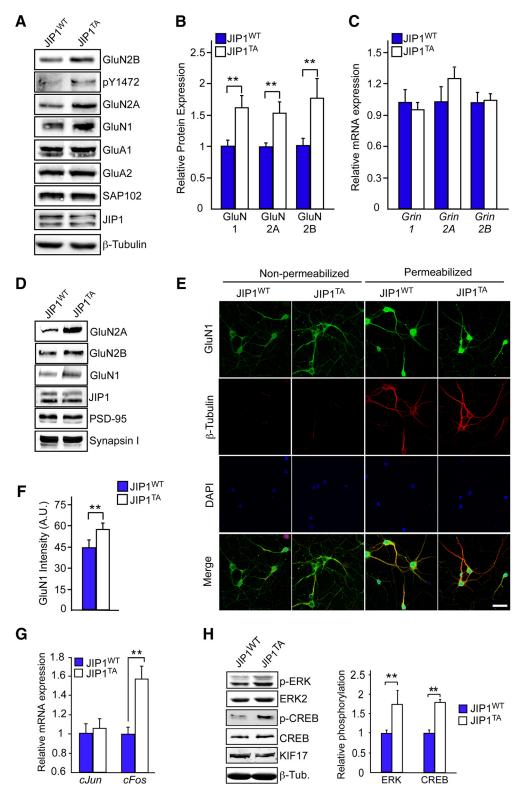


Figure 7. The JIP1 TA mutation promotes increased NMDAR expression and activity. A-C, Lysates prepared from the hippocampio of JIP1 TA and JIP1 TA mice were examined by immunoblot analysis by probing with antibodies to NMDA and AMPAR subunits, SAP102, JIP1, and β-tubulin (A). The number of NMDAR subunits in the hippocampus was determined and normalized to the amount of β-tubulin in each sample (B, mean \pm SEM, n=5; **P<0.01, Student's P<0.01, St

collateral-evoked responses that was comparable between JIP1 TA and JIP1 WT slices (Fig. 5I; 62 \pm 1% of baseline at 60 min after drug application for JIP1 TA vs 65 \pm 8% for JIP1 WT; $t_{(18)} = 0.68$, p = 0.49; two-tailed unpaired Student's t test). These results suggest that, whereas the JIP1 TA mutation blocks the NMDAR-dependent forms of LTD, it does not affect the mGluR-dependent form.

Another important aspect of synaptic modification is synaptic depotentiation, namely that the synaptic potentiation could be subsequently reversed by LFS. In view of the finding that two trains of HFS at 100 Hz induced significant LTP in response to a 20 s intertetanus interval in both groups of mice (Fig. 5D), we used this HFS protocol followed by LFS 10 min later (900 pulses at 1 Hz) to induce depotentiation (Fig. 5J). One hour after the LFS, the evoked responses in both JIP1 $^{\mathrm{WT}}$ and JIP1 TA mice dropped to the baseline level (100 \pm 4% compared with baseline before the HFS for JIP1 WT, and $101 \pm 3\%$ for JIP1 ^{TA}; $t_{(18)} = 0.23$, p = 0.82; twotailed unpaired Student's t test), indicating effective LFS-induced depotentiation in both genotypes.

We next tested whether application of a selective inhibitor of JNK would mimic the effects of the JIP1 Thr 103 Ala mutation on NMDA-receptor dependent forms of LTP and LTD. Slices were obtained from wild-type mice and exposed either to the selective JNK inhibitor JNK-in-8 (6 μM) or solvent (aCSF) for 20 min before stimulation protocols. The drug application was maintained throughout the recording period. Basal synaptic transmission was not affected by JNK-in-8, as indicated by the unchanged I/O relation (twoway repeated-measures ANOVA: genotype $F_{(1,308)} = 0.88$, p = 0.35); amplitude $F_{(14,308)} = 61.73, p < 0.001$; genotype × amplitude $F_{(14,308)} = 0.35$, p = 0.63) and unchanged PPF (two-way repeated-measures ANOVA: genotype $F_{(1,132)} = 0.46$, p = 0.56); interval $F_{(6,132)} = 42.12$, p <0.001; genotype × interval $F_{(6,132)} = 1.17$, p = 0.11; Fig. 6A, B). In addition, inhibiting JNK activation in slices did not affect HFS LTP (Fig. 6C; 60 min after 2×100 Hz tetanization: $127 \pm 6\%$ of baseline for vehicle-treated slices vs 125 ± 5% for JNK-in-8-treated slices; $t_{(22)} = 0.25$, p =0.81; two-tailed unpaired Student's *t* test).

However, JNK inhibition prevented the induction of LTD using the 1 Hz stimulation protocol (Fig. 6*D*; vehicle-treated slices, 77 \pm 3%; JNK-in-8-treated slices, 101 \pm 4%; p < 0.05; $t_{(18)} = 4.81$, p < 0.001; two-tailed unpaired Student's t test). The latter result closely parallels the result obtained for the JIP1 ^{TA} slices (cf. Fig. 5*F*). Therefore, these data confirm that JIP1-mediated JNK

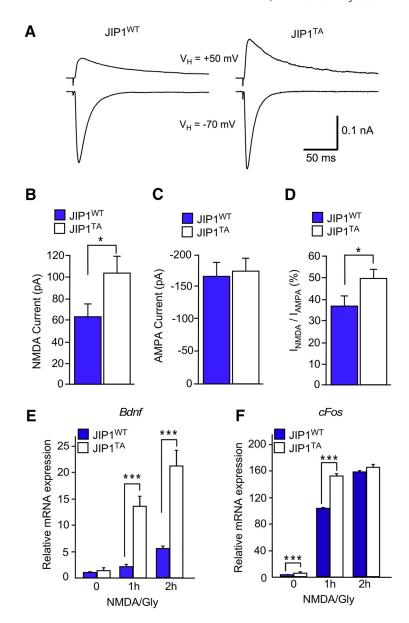


Figure 8. Increased synaptic NMDAR activity in hippocampal slices from JIP1 TA mice. A, Whole-cell voltage-clamp traces depicting typical EPSCs elicited by stimulating the Schaffer collaterals while recording CA1 pyramidal cells from JIP1 WT and JIP1 TA hippocampal slices. Strong blockade of NMDARs at a holding potential of -70 mV by magnesium isolates the AMPA component (bottom traces). The \pm 50 mV upper traces primarily represent the NMDA component because the traces were recorded 30 \pm 50 ms after stimulation when \sim 90% of the AMPAR response had decayed. The traces show the increase in the NMDAR-mediated component for the JIP1 TA group relative to JIP1 WT, whereas AMPAR mediated responses were not different. Traces depicted are averages of 20 sweeps for both NMDA and AMPAR-mediated components of JIP1 WT recorded from the same pyramidal neuron; 10 and 15 sweeps were used to produce the averages depicted for the NMDA and AMPAR-mediated components recorded from a JIP1 TA pyramidal neuron. B, C, Comparison of average NMDA and AMPAR currents. NMDAR-mediated currents (B) were significantly greater in the JIP1 TA group compared with the JIP1 WT group (mean \pm SEM; $n=16\sim$ 17 cells; *p<0.05, Student's t test), whereas average AMPAR-mediated currents (C) did not differ. D, NMDA to AMPAR current ratios were evaluated on a cell-by-cell basis. The JIP1^{TA} ratios (I_{NMDA}/I_{AMPA}) were larger than ratios measured from neurons in the JIP1^{WT} group (mean \pm SEM; n=16 \sim 17 cells; *p < 0.05, Student's t test). **E**, **F**, NMDA-stimulated gene expression in primary cultures of JIP1 WT and JIP1 TA hippocampal neurons was studied by treating neurons with 100 μ m NMDA plus 10 μ m glycine. The expression of $Bdnf(\textbf{\textit{E}})$ and cFos(F) mRNA was quantitated by qRT-PCR and normalized to Gapdh (mean \pm SEM; $n=5\sim$ 6; ***p<0.001, two-way ANOVA followed by Bonferroni's post hoc test).

activation plays an essential, but differential, role in certain forms NMDA-receptor dependent synaptic plasticity.

JIP1 TA mice exhibit increased NMDAR signaling

It is established that JIP scaffold proteins influence NMDAR activity (Kennedy et al., 2007). The induction of synaptic plasticity

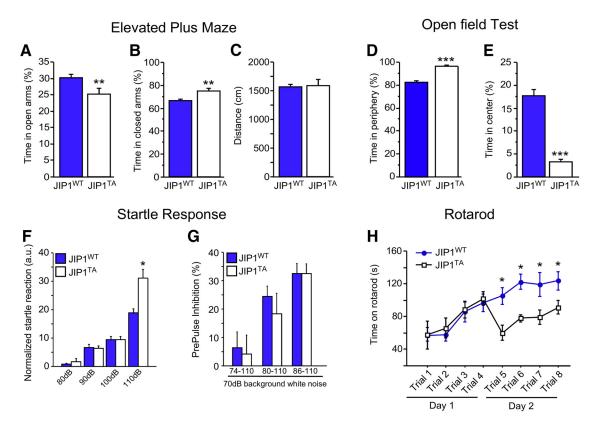


Figure 9. JIP1 TA mice display normal locomotor function, motor coordination, elevated anxiety-like behavior, and increased acoustic startle response. **A–C**, Results of elevated plus maze test. JIP1 TA mice show decreased time spent in open arms (**A**) and increased time spent in closed arms (**B**) relative to wild-type mice, indicative of elevated anxiety-like behaviors. In addition, JIP1 TA mice show normal activity as measured by total distance traveled (**C**). Data are shown as mean ± SEM. n = 10; ***p < 0.01, Student's t test. **D**, **E**, Open-field test. JIP1 TA mice show increased anxiety-like behavior in an open-field test. Mice were allowed to explore an open field for 5 min. JIP1 TA mice spent more time in the periphery (**D**) and less time in the center region of the open field (**E**), both indicators of increased levels of anxiety-like behavior in this test. Data are shown as mean ± SEM. n = 10; ***t = 10, ***t

at lower stimulation frequencies detected in JIP1 TA mice compared with JIP1 WT mice (Fig. 6) may therefore be mediated by changes in NMDAR signaling. Indeed, we found increased expression of NMDAR subunits GluN1 ($t_{(8)} = 2.66$, p = 0.029; two-tailed unpaired Student's t test), GluN2A ($t_{(8)} = 2.85$, p =0.037; two-tailed unpaired Student's t test), and GluN2B proteins $(t_{(8)} = 2.65, p = 0.027;$ two-tailed unpaired Student's t test; but not mRNA) in the whole hippocampus of JIP1 TA mice compared with JIP1 WT mice (Fig. 7A-C). Control studies detected no differences in the expression of the AMPAR subunits GluA1 and GluA2 (Fig. 7A). The increased expression of NMDAR subunits was also detected in hippocampal synaptoneurosomes (Fig. 7D) and by increased cell surface detection of GluN1 by immunofluorescence analysis of hippocampal neurons ($t_{(16)} = 2.85$, p =0.012; two-tailed unpaired Student's t test; Fig. 7E,F) cultured from JIP1 TA mice compared with JIP1 TH mice. These changes were associated with biochemical evidence of increased NMDAR activity through assessment of downstream signaling changes, including increased GluN2B Tyr ¹⁴⁷² phosphorylation (Salter and Kalia, 2004; Fig. 7A), increased cFos mRNA expression ($t_{(9)} =$ 3.31, p = 0.009; two-tailed unpaired Student's t test; Bading et al., 1993; Fig. 7G), ERK phosphorylation ($t_{(8)} = 3.11$, p = 0.017; two-tailed unpaired Student's t test; Xia et al., 1996; Fig. 7H), and

CREB phosphorylation ($t_{(8)} = 5.25$, p = 0.008; two-tailed unpaired Student's t test; Ginty et al., 1993; Fig. 7H). Control experiments demonstrated that the expression of KIF17, a microtubule motor that transports GluN2B vesicles and regulates synaptic plasticity (Yin et al., 2011, 2012), was similar in JIP1 $^{\rm WT}$ and JIP1 $^{\rm TA}$ mice (Fig. 7H). Together, these data indicate that JIP1 $^{\rm TA}$ mice may exhibit increased NMDAR signaling in the hippocampus compared with JIP1 $^{\rm WT}$ mice.

 0.28, p = 0.77; two-tailed unpaired Student's t test; Fig. 8A, C). These findings were reflected in the NMDAR/AMPAR peak current ratio calculated for each cell tested. JIP1 WT mice had a NMDA/AM-PAR ratio of 37 \pm 4%, whereas JIP1 ^{TA} mice yielded a significantly larger ratio of $50 \pm 4\%$ ($t_{(31)} = 2.21$, p = 0.038; twotailed unpaired Student's t test; Fig. 8D). This analysis demonstrated that NMDAR signaling is increased in JIP1 TA hippocampal neurons compared with those of JIP1 WT mice. To confirm this conclusion, we examined NMDAR-stimulated gene expression in primary JIP1 TA and JIP1 WT hippocampal neurons. We found that NMDA-induced expression of brainderived neurotrophic factor (Bdnf) mRNA (Ghosh et al., 1994; Tabuchi et al., 2000) and cFos mRNA (Bading et al., 1993; Xia et al., 1996) by JIP1 TA neurons were increased when compared with JIP1 WT neurons ([two-way ANOVA: genotype $F_{(1,29)} = 70.26$, p < 0.001; time $F_{(2,29)} =$ 39.7, p < 0.001; genotype × time $F_{(2,29)} =$ 14.91, p < 0.001; for *Bdnf* mRNA]; [two-way ANOVA: genotype $F_{(1,30)} = 17.67, p < 0.001;$ time $F_{(2,30)} = 182.1$, p < 0.001; genotype \times time $F_{(2,30)} = 7.98$, p < 0.001; for *cFos* mRNA]; Fig. 8E,F). Together, these data indicate that the increased NMDAmediated signaling detected in the JIP1 TA hippocampus is due to enhanced NMDAR activity in individual neurons.

JIP1-mediated JNK activation in locomotor, sensory and emotional responses

To address the possibility that increased NMDAR signaling in JIP1^{TA} mice compared with JIP1^{WT} mice may cause behavioral changes, we first performed a battery of basic behavioral tests of CNS function on JIP1^{WT} and JIP1^{TA} mice. We found that anxiety-related behavior was increased in JIP1^{TA} mice. For example, JIP1^{TA} mice in an elevated plus maze spent less time in the open arms, compared with JIP1^{WT} mice $(t_{(18)} = 2.79, p = 0.012;$ two-tailed unpaired Student's t test), with no changes in locomotor activity $(t_{(18)} = 0.76, p = 0.45;$ two-tailed unpaired Student's t test; Fig.

9A–C). Moreover, JIP1 ^{TA} mice spent significantly less time in the center ($t_{(18)} = 5.40$, p < 0.001; two-tailed unpaired Student's t test) and more in the periphery ($t_{(18)} = 6.46$, p < 0.001; two-tailed unpaired Student's t test) during an open-field test compared with JIP1 ^{WT} animals (Fig. 9 D, E). Consistent with previous reports (Grillon et al., 1998), elevated anxiety in JIP1 ^{TA} mice was accompanied by an enhancement of the startle response to strong acoustic stimuli (two-way repeated-measures ANOVA: genotype $F_{(1,42)} = 5.42$, p = 0.035; stimulus $F_{(3,42)} = 105.42$, p < 0.001; genotype × stimulus = 9.25, = 0.001; Fig. 9= 0.001; No changes in sensorimotor gating (pre-pulse inhibition) were observed (two-way)

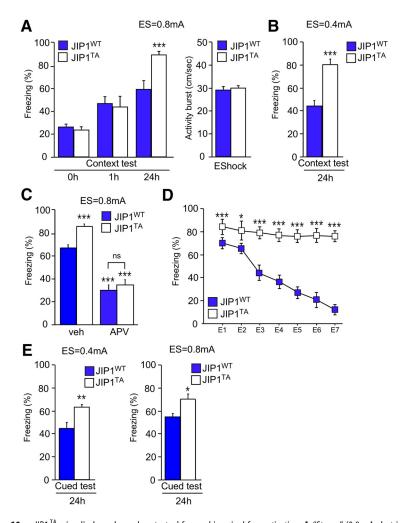


Figure 10. JIP1 TA mice display enhanced contextual fear and impaired fear extinction. A, "Strong" (0.8 mA electric shock) training demonstrated that JIP1^{TA} and JIP1^{WT} littermate mice exhibited similar contextual freezing when tested immediately after training or 1 h later, but the JIP1 TA mice froze more than JIP1 WT mice at 24 h after training (left). Foot-shock reactivity during fear-conditioning training did not significantly differ between JIP1 TA and JIP1 WT mice (right). Data are shown as mean \pm SEM. $n = 10 \sim 11$; *** p < 0.001, Student's t test. **B**, "Weak" (0.4 mA electric shock) training demonstrated that JIP1 TA mice (n = 14) exhibited contextual freezing that was similar to the "strong" training schedule, but JIP1 $^{\rm WT}$ mice (n=14) displayed significantly less contextual fear conditioning at 24 h after training. Data are shown as mean \pm SEM. n=14; ***p<0.001, Student's t test. $\boldsymbol{\zeta}$, JIP1 TA and JIP1 WT littermate mice were infused with vehicle or the selective NMDAR antagonist APV (10 μ g/ml) before "strong" (0.8 mA) contextual fear conditioning. The following day, there was a similar impairment in both genotypes in freezing levels to the conditioning context. Data are shown as mean \pm SEM. n=10; ***p<0.001 vs JIP1 WT+ veh; n.s.; JIP1 WT+ APV vs JIP1 TA+ APV, two-way ANOVA followed by Bonferroni's post hoc test. **D**, JIP1 And JIP1 TA and JIP1 WT littermate mice were trained by "strong" (0.8 mA) electric shock) contextual fear conditioning. Extinction began 24 h later and consisted of daily 3 min reexposures of mice to the conditioning context in the absence of shock. When compared with JIP1 WT littermates, JIP1 TA mice showed increased freezing behavior throughout extinction days 1–7 (E1–E7), indicating impaired extinction process in JIP1^{TA} mice. Data are shown as mean \pm SEM. n=10; *p<0.05, ****p<0.001; two-way repeated-measures ANOVA followed by Bonferroni's post hoc test. E, "Weak" (0.4 mA electric shock; left) and "strong" (0.8 mA; right) cued fear training, consisting of a single pair of cue (tone) and shock, demonstrated that JIP1 TA and JIP1 WT mice exhibited enhanced conditioned freezing to a cue (tone) when tested 24 h after training. Data are shown as mean \pm SEM. n=11; *p<0.05, **p<0.01, Student's t test.

repeated-measures ANOVA: genotype $F_{(1,28)}=0.13, p=0.722$; Fig. 9*G*). We also found that "fast" improvement in motor coordination on the accelerating rotarod was comparable between JIP1 WT and JIP1 TA mice. However, during the second day of rotarod training JIP1 TA mice did not display improved "slow" skill learning, unlike JIP1 WT mice, as indicated by significant genotype × trial interaction value (two-way repeated-measures ANOVA: genotype $F_{(1,98)}=4.84, p=0.045$), trial $F_{(7,98)}=7.73, p<0.001$; genotype × trial $F_{(7,98)}=3.34, p=0.003$; Fig. 9*H*). Together, these data demonstrate that JIP1 TA mice have normal sensory and motor activity and attention function. However,

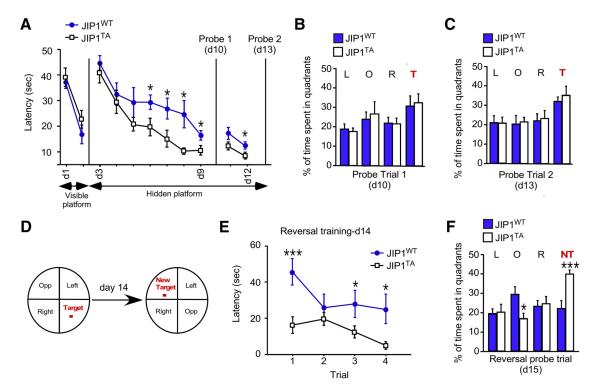


Figure 11. JIP1 TA mice exhibit enhanced acquisition and reversal learning in the MWM test. A-C, JIP1 TA and JIP1 WT littermate mice learned the visible platform task (day 1 and 2), as indicated by reductions in escape time during training. The mice were then trained to find a hidden platform during the next 7 d. JIP1 TA mice showed faster escape latencies at days 6-9 of training compared with JIP1 WT littermates (A). A first probe test (day 10) was conducted 24 h after the completion of training. No significant differences in percentage time spent in the target quadrant (T) between JIP1 TA and JIP1 WT mice were observed (B). The mice were then subjected to 2 d of additional training (days 11-12) and a second probe trial was performed 24 h later. No significant differences between JIP1 TA and JIP1 WT mice were observed during second probe trial (C). Data are shown as mean \pm SEM. D-F, Twenty-four hours after the second probe test, the platform was moved to the opposite quadrant in the pool and mice were trained for four trials (day 14, reversal learning). In this new setting, JIP1 TA mice displayed shorter escape time to find newly placed platform (NT) compared with JIP1 WT littermate mice (E). The probe test for reversal training was conducted 24 h after the completion of new platform training (day 15). Analysis of the time spent in the quadrants revealed that JIP1 TA mice spent significantly more time in the NT than JIP1 WT mice (F). Data are shown as mean E SEM. E 14; E 20.05, E 20.05, E 30.10, two-way repeated-measures ANOVA (F) and two-way ANOVA (F) followed by Bonferroni's post hoc tests.

JIP1 TA mice displayed increased levels of anxiety-related behaviors and altered skill learning.

Improved extinction-resistant contextual fear memory in IIP1 $^{\rm TA}$ mice

Contextual fear conditioning triggers JIP1-dependent JNK activation in the hippocampus (Fig. 4). Moreover, JNK activation in the hippocampus is implicated in some forms of hippocampusdependent memory (Sherrin et al., 2011). We therefore compared JIP1 WT and JIP1 TA mice to determine whether JIP1-mediated JNK activation is required for the behavioral response to contextual fear conditioning. The mice received either "weak" (0.4 mA shock) or "strong" (0.8 mA shock) training and were then reexposed to the chamber 24 h later to assess long-term fear memory (Shalin et al., 2006). The two training procedures were used to adjust for nociceptive sensitivity and also to test whether a "weak" stimulation protocol similar to one that facilitated synaptic plasticity (cf. Fig. 5E) results in enhanced hippocampal memory in JIP1 TA mice. JIP1 TA and JIP1 WT mice showed no differences in freezing or activity during the exploration period before the footshock (data not shown), or after the 0.8mA foot-shock delivery $(t_{(19)} = 0.46, p = 0.65;$ two-tailed unpaired Student's t test; Fig. 10A, right). Significantly enhanced contextual freezing of JIP1 TA mice compared with JIP1 WT controls was found 24 h after "strong" training (Fig. 10A, left; mean percentage freezing: JIP1 TA = $85 \pm 3\%$; JIP1 WT = $59 \pm 4\%$; $t_{(19)} = 5.11$, p < 0.001; two-tailed unpaired Student's t test). This enhancement of contextual fear was more pronounced in JIP1 TA mice when the

"weak" paradigm was used (Fig. 10B; mean percentage freezing: JIP1 TA = $81 \pm 4\%$; JIP1 WT = $44 \pm 5\%$; $t_{(26)} = 5.26$, p < 0.001; two-tailed unpaired Student's t test). We next investigated short-term memory by testing JIP1 TA and JIP1 WT mice immediately or 1 h after the "strong" conditioning paradigm. We found that JIP1 TA and JIP1 WT mice displayed similar responses in the both contextual tests ($t_{(20)} = 0.18$, p = 0.85; 0 h after conditioning, two-tailed unpaired Student's t test; $t_{(26)} = 0.11$, p = 0.75; 1 h after conditioning, two-tailed unpaired Student's t test; Fig. $t_{(26)} = 0.11$, $t_{(26)} = 0$

We then tested whether contextual fear memory is affected differentially in JIP1 ^{TA} and JIP1 ^{WT} mice after antagonism of the NMDARs. JIP1 ^{TA} and JIP1 ^{WT} mice were implanted with cannulae into the dorsal hippocampus and subjected to contextual fear conditioning. A selective NMDAR antagonist, APV (10 μ g/ml; 0.5 ml) or solvent (aCSF) was infused into the dorsal hippocampus 15 min before training. Both JIP1 ^{TA} and JIP1 ^{WT} mice given APV showed substantially reduced freezing to context compared with mice that received aCSF before conditioning (two-way ANOVA: genotype $F_{(1,36)} = 8.76$, p = 0.0054; treatment $F_{(1,36)} =$

143.06, p < 0.001; genotype × treatment = 5.29, p = 0.027; Fig. 10*C*). However, no differences between APV-treated JIP1 TA mice and APV-treated JIP1 TA mice were observed (Bonferroni post hoc test: p = 0.62; Fig. 10*C*), indicating that the memory-facilitating effects observed in JIP1 TA were mediated via activation of NMDARs. Together, these experiments show that blockade of JIP1-mediated JNK activation can improve contextual learning induced by a relatively mild learning protocol and that enhanced NMDAR function is required for this facilitatory effect of JIP1 TA mutation on learning.

Previous studies have demonstrated that activation of hippocampal NMDAR regulates fear extinction (Szapiro et al., 2003; Suzuki et al., 2004; Leaderbrand et al., 2014) and that NMDAR blockade can prevent fear extinction in a variety of fearrelated tasks (Santini et al., 2001; Sotres-Bayon et al., 2007). Therefore, we extended our cognitive evaluation of JIP1 TA mice by assessing contextual fear extinction to further assess the link of JIP1-JNK activation through NMDARs. Conditioned JIP1 TA and JIP1 WT mice were subjected to brief (3 min) daily extinction trials by nonreinforced placement in the training context for 7 d after contextual conditioning. During the extinction trials, JIP1 TA mice exhibited almost unchanged freezing responses, indicating nearly fully impaired extinction compared with JIP1 WT mice (repeated-measures ANOVA: genotype $F_{(1,108)} = 388.76, p < 0.001$; time $F_{(6,108)} =$ 36.26, p < 0.001; $F_{(6,108)}$ genotype \times time = 25.35, p < 0.001; Fig. 10D). Over-

all, these results indicate that the enhanced contextual fear observed in JIP1 TA mice is strongly resistant to extinction.

Finally, cue-dependent fear conditioning was assessed in JIP1 TA mice and JIP1 WT controls. As with contextual fear conditioning, mice received either a "strong" or a "weak" training protocol, both consisting of a single pairing of tone and shock at 0.8 and 0.4 mA, respectively. In both protocols, JIP1 TA mice, compared with JIP1 WT controls, displayed significantly more freezing when reexposed to the tone 24 h after training (Fig. 10*E*, left; mean percentage freezing: "weak" cued test- JIP1 TA = 66 \pm 3%; JIP1 WT = 44 \pm 4%; $t_{(20)}=3.71$, p=0.01; two-tailed unpaired Student's t test; Fig. 10*E*, right; mean percentage freezing: "strong" cued test-JIP1 TA = 70 \pm 3%; JIP1 WT = 55 \pm 4%; $t_{(20)}=2.65$, p=0.015; two-tailed unpaired Student's t test), indicating that JIP1 TA mice display memory enhancement in the fear learning tasks in a manner involving both amygdala- and hippocampus-dependent memory systems.

Improved spatial learning in JIP1 TA mice

We next tested JIP1^{TA} and JIP1^{WT} mice using the MWM protocol, a hippocampus-dependent spatial memory task (Morris et al., 1982). During the hippocampus-independent visible portion

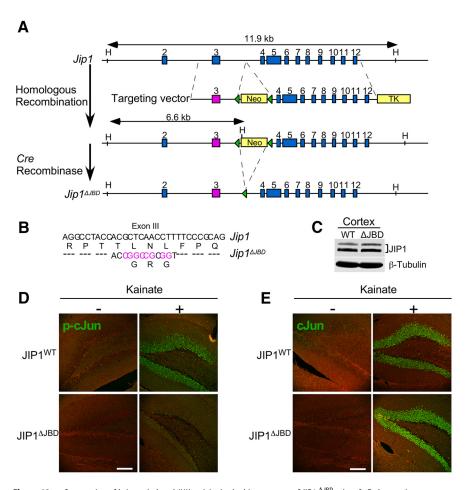


Figure 12. Suppression of kainate-induced JNK activity in the hippocampus of JIP1 $^{\Delta JBD}$ mice. **A**, **B**, A targeting vector was designed to replace JIP1 residues Leu 160 -Asn 161 -Leu 162 with Gly 160 -Arg 161 -Gly 162 in exon 3 of the $^{\it Jip1}$ gene by homologous recombination in ES cells. The *floxed Neo* $^{\it R}$ cassette inserted in intron 3 and used for selection was deleted with *Cre* recombinase. H, HindIII restriction enzyme. **C**, Lysates prepared from the cerebral cortex of $^{\it Jip1}$ $^{\it L}$ (WT) and $^{\it Jip1}$ $^{\it L}$ $^{\it JBD/\Delta}$ $^{\it JBD}$ ($^{\it L}$ JBD) mice were examined by immunoblot analysis using antibodies to JIP1 and $^{\it B}$ -Tubulin. **D**, **E**, JIP1 $^{\it WT}$ and JIP $^{\it L}$ Bod mice were treated without and with kainate. Representative sections of the DG stained (green) with antibodies to phospho-cJun (**D**) or cJun (**E**) are presented. DNA was stained with DAPI (red). Scale bar, 75 $^{\it L}$ m.

of the training, both JIP1 TA and JIP1 WT mice readily learned to find the marked platform (Fig. 11A). Mice were then trained to swim to a hidden platform located in a fixed location of the pool (Fig. 11 A, D; target quadrant, T). JIP1 TA mice had significantly decreased escape latencies compared with JIP1 WT mice on days 6-9 and day 12 of training (two-way repeated-measures ANOVA: genotype $F_{(1,208)} = 11.03$, p = 0.0049; time $F_{(8,208)} = 21.01$, p < 0.001; genotype \times time $F_{(8,208)} = 0.74$, p = 0.65; Fig. 11A). However, JIP1 ^{TA} and JIP1 ^{WT} mice did not differ in the time spent in the platform-containing quadrant during probe trials on day 10 (Fig. 11*B*; 36 \pm 3% of time in quadrant T for JIP1 ^{TA}; 35 \pm 2% JIP1 WT; $t_{(26)} = 0.27$, p = 0.783; two-tailed unpaired Student's ttest) and day 13 (Fig. 11*C*; 38 \pm 5% of time in quadrant T for JIP1^{TA} mice; 36 \pm 3% JIP1^{WT} mice; $t_{(26)} = 0.34$, p = 0.734; two-tailed unpaired Student's t test). Although JIP1 TA mice did not display preference for the target quadrant during probe trials, the decrease in escape latency during training suggests that the presence of the JIP1 Thr 103 Ala mutation appears to affect gradual learning of the hidden platform MWM task moderately.

To test whether JIP1 TA mice display enhanced fast spatial learning, we performed a reversal of the hidden platform in the MWM. The hidden platform was moved to the opposite quadrant of the pool (new target quadrant, NT) and the mice were

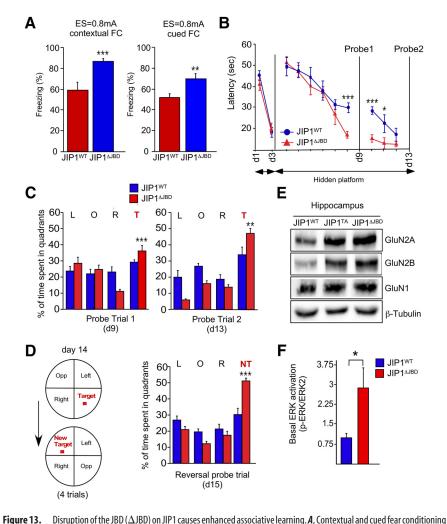


Figure 13. Disruption of the JBU (ΔJBD) on JPT (Causes enhanced associative learning, **A**, contextual and cued rear conditioning of JIP1 $^{\Delta JBD}$ and JIP1 WT mice consisted of one exposure to cue (context + tone) and 0.8 m Ashock (mean \pm SEM; $n=10\sim11$; **p<0.01, Student's t test ****p<0.001, Student's t test ****p<0.001, Student's t test). **B**, **C**, MWM tests of mean latencies to escape to a visible (days 1–2) or a hidden platform (days 3–12) are presented for JIP1 $^{\Delta JBD}$ or JIP1 WT mice (**B**). Probe trials were performed on days 9 and 13 of water maze training (**C**). JIP1 $^{\Delta JBD}$ mice spent significantly longer time in the target quadrant compared with JIP1 WT littermates (mean \pm SEM; n=10; **p<0.05; ***p<0.05; ***p<0.01; ****p<0.001, two-way repeated-measures ANOVA (**B**) and two-way ANOVA (**C**) followed by Bonferroni's post hoc tests). **D**, The water maze platform was moved to the opposite quadrant in the pool and mice were trained for four trials (day 14, reversal learning). The probe test for reversal training was conducted 24 h after the completion of new platform training (day 15). Analysis of the time spent in the quadrants during the probe trial revealed that JIP1 $^{\Delta JBD}$ mice spent significantly more time in the NT than JIP1 WT mice (mean \pm SEM; n=10; ****p<0.001, two-way ANOVA followed by Bonferroni's post hoc test). **E**, Hippocampus lysates of JIP1 WT , JIP1 TA , and JIP1 $^{\Delta JBD}$ mice were examined by immunoblot analysis by probing with antibodies to NMDAR subunits and β -Tubulin. **F**, The amount of phospho-ERK in hippocampus lysates of naive JIP1 WT and JIP1 $^{\Delta JBD}$ mice was quantified by multiplexed ELISA and normalized to the amount of ERK2 in each sample. Data are shown as mean \pm SEM. n=5; *p<0.05, Student's t test.

trained to swim to this new location for four trials separated by 15–20 min (Fig. 11D). JIP1 TA mice learned significantly faster to swim to the new platform location compared with control mice (two-way repeated-measures ANOVA: genotype $F_{(1,78)}=16.37$, p=0.012; time $F_{(3,78)}=1.98$, p=0.13; genotype × time $F_{(3,78)}=1.86$, p=0.15; Fig. 11E). Moreover, the JIP1 TA mice displayed increased preference for the NT during a probe trial given on probe day 15 (Fig. 11F; time spent in NT: 41 \pm 2% for JIP1 TA; $23\pm4\%$ for JIP1 TY; $t_{(26)}=4.02$, p<0.001; two-tailed unpaired Student's t test). This enhancement in spatial reversal learning of the new platform location was accompanied by a decrease in the percentage of time JIP1 TA mice spent searching for the platform in the previously correct quadrant (O; $t_{(26)}=2.34$, p=0.028; two-tailed unpaired Student's t test; Fig. 11F). These data indi-

cate that the Thr 103 Ala mutation in JIP1 allows for improved fast spatial learning.

JNK binding is required for JIP1-mediated NMDAR regulation of learning and memory

To confirm that JIP1-mediated JNK activation plays a critical role in NMDAR-dependent, hippocampus-dependent learning and memory, we tested an independent mouse model of defective JIP1-mediated JNK activation: JIP1 lacking the JNK binding domain (JBD). To create a mouse model lacking the JBD of JIP1 (JIP1 ^{ΔJBD} mice), the core of the JBD (Leu¹⁶⁰-Asn 161-Leu 162) mediating a hydrophobic interaction of JIP1 with JNK (Whitmarsh et al., 2001; Heo et al., 2004) was replaced with Gly ¹⁶⁰-Arg ¹⁶¹-Gly ¹⁶² in JIP $^{\Delta JBD}$ mice (Fig. 12 A, B). Immunoblot analysis of brain extracts from JIP1 WT and JIP1 $^{\Delta JBD}$ mice indicated similar levels of JIP1 protein expression (Fig. 12C). To establish that the ΔJBD mutation blocks JIP1mediated JNK activation in vivo, we compared kainate-induced excitotoxicity in the hippocampus of JIP1 WT and JIP1 ΔJBD mice. This analysis demonstrated that kainate caused similar increases in cJun expression in the DG of JIP1WT and JIP1 $^{\Delta \rm JBD}$ mice, but kainate-stimulated cJun phosphorylation was markedly suppressed in JIP1 $^{\Delta JBD}$ mice compared with JIP1 WT mice (Fig. 12 D,E). These data confirm that JIP1 $^{\Delta JBD}$ mice represent a model of defective JIP1-mediated JNK signaling.

We then examined contextual and cued fear conditioning to assess the effect of the JIP1 $^{\Delta \text{JBD}}$ mutation on learning and memory. The JIP1 $^{\Delta \text{JBD}}$ mice displayed a significantly increased conditioned freezing response in both learning tasks compared with the JIP1 $^{\text{WT}}$ mice when tested at 24 h after training (Fig. 13A, left: mean percentage contextual freezing, JIP1 $^{\text{WT}}$ = $58 \pm 3\%$; JIP1 $^{\Delta \text{JBD}}$ = $89 \pm 3\%$; $t_{(18)}$ = 6.30, p < 0.001; two-tailed unpaired Student's t test; Fig. 13A, right: mean percentage cued

test; Fig. 13A, right: mean percentage cued freezing; JIP1 WT = 51 \pm 1%; JIP1 $^{\Delta JBD}$ = 69 \pm 6%; $t_{(20)}$ = 2.97, p = 0.007; two-tailed unpaired Student's t test), a result very similar to that observed for the JIP1 TA mice (Fig. 10). JIP1 $^{\Delta JBD}$ mice exhibited enhanced learning in the MWM (two-way repeated-measures ANOVA: genotype $F_{(1,162)}$ = 26.04, p < 0.001; time $F_{(8,162)}$ = 126.7, p < 0.001; genotype × time $_{(8,162)}$ = 1.75, p = 0.84), and spent significantly more time in the target quadrant compared with control JIP1 The mice when subjected to probe trials on day 9 ($t_{(18)}$ = 4.02, p < 0.001; Fig. 13C, left; 41 \pm 2% of time in quadrant T for JIP1 To JIP1 Jabb Jabb; 32 \pm 1% of time in quadrant T for JIP1 To JIP1 AJBD; 36 \pm 3% of time in quadrant T for JIP1 To JIP1 AJBD; 36 \pm 3% of time in quadrant T for JIP1 To JIP

not show enhanced memory retention in the hidden platform MWM task (cf. Fig. 11B,C). The larger phenotype in the JIP1 $^{\Delta \text{JBD}}$ mice in the MWM compared with the JIP1 $^{\text{TA}}$ mice may result from a greater effect of the JNK-binding site mutation on JIP1-mediated JNK activation than the mutation of the T103 phosphorylation site on JIP1.

We then examined reversal learning by moving the hidden platform to the opposite quadrant (Fig. 13D, left). Similar to JIP1 TA mice, the JIP1 $^{\Delta JBD}$ mice showed shorter latencies than control mice in finding the new platform location during a 1 d trial period (data not shown). Moreover, JIP1 ^{AJBD} mice spent significantly more time in the NT compared with JIP1 WT control mice (Fig. 13D, right; mean percentage of time spent in the NT was 49 \pm 3%, for JIP1 $^{\Delta JBD}$; 28 \pm 4% for JIP1 WT; $t_{(18)} = 4.21$, p < 0.001; two-tailed unpaired Student's t test). Furthermore, like JIP1 TA mice, we found significantly increased expression of GluN1, GluN2A, and GluN2B in the hippocampus of JIP1 $^{\Delta JBD}$ mice compared with hippocampi from

JIP1 WT mice (Fig. 13*E*) and increased ERK activation ($t_{(8)} = 2.57$, p = 0.033; two-tailed unpaired Student's *t* test; Fig. 13*F*).

Together, these analyses of JIP1 TA and JIP1 DIPD mice confirm that loss of JIP1-mediated JNK activation enhances NMDA-dependent hippocampus-dependent memory.

Discussion

The JIP1 scaffold protein can assemble a functional JNK signaling module formed by members of the mixed-lineage protein kinase family of MAP3K, the MAP2K family member MKK7, and JNK (Whitmarsh et al., 1998). However, JIP1 also functions as an adapter protein that mediates transport by microtubule motor proteins (Fu and Holzbaur, 2014), including kinesin-1 (Verhey et al., 2001; Whitmarsh et al., 2001) and dynein (Standen et al., 2009; Fu and Holzbaur, 2013). These two functions of JIP1 compromise the interpretation of loss-of-function studies focused on the analysis of JIP1 knock-out mice (Whitmarsh et al., 2001; Kennedy et al., 2007). To overcome this limitation, we studied two mouse models with germline mutations in the *Jip1* gene that prevent JIP1-mediated JNK activation. First, a point mutation of the JIP1 phosphorylation site Thr 103 (by replacement with Ala) in JIP1 TA mice suppresses JIP1-mediated JNK activation (Fig. 3) by disrupting the regulated interaction of mixed-lineage protein kinases with JIP1 (Morel et al., 2010). Second, a three-residue mutation of the JIP1 site that binds JNK in JIP1 $^{\Delta \rm JBD}$ mice prevents JIP1-mediated JNK activation (Fig. 12). These complementary mouse models therefore provided an opportunity to disrupt JIP1-mediated JNK activation selectively in vivo.

It is established that the JIP1 scaffold protein mediates JNK activation in the neuronal response to excitotoxin (Whitmarsh et al., 2001) and in adipose tissue during metabolic stress responses (Jaeschke et al., 2004; Morel et al., 2010). The results of the present study extend these findings to neural functioning by demonstrating that JIP1-linked JNK activation in the hippocampus regulates contextual fear conditioning in a NMDAR-dependent fashion. This finding builds on the previous demonstration that

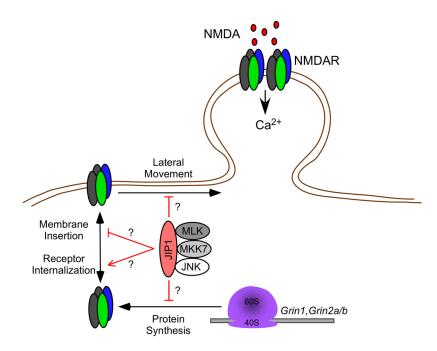


Figure 14. A model of how JIP1-mediated JNK signaling regulates synaptic NMDAR expression. JIP1-dependent JNK activation by the NMDAR may suppress translation of NMDAR subunit mRNA (*Grin1*, *Grin2a*, *Grin2b*). Alternatively, the same pathway regulates cell surface insertion or retrieval of NMDARs and/or lateral diffusion of extrasynaptic NMDARs into synaptic sites.

JNK1-deficient mice exhibit enhanced contextual fear conditioning (Sherrin et al., 2010) and altered synaptic plasticity (Li et al., 2007).

We further show that JIP1-mediated JNK activation regulates NMDAR signal transduction associated with an altered threshold for LTP, decreased long-term fear memory, and decreased spatial memory (Figs. 5, 6, 8, 10, 11, 13). These observations are consistent with the conclusion that JNK normally functions to regulate negatively mechanisms responsible for learning and memory (Sherrin et al., 2011). We found that the enhanced learning in JIP1 mutant mice was associated with an increase in the NMDAR component of the synaptic response, and enhanced activity of downstream pathways that facilitate induction of NMDAR-dependent LTP. This is consistent with previous reports that have separately implicated both JNK signaling (Sherrin et al., 2010) and CA1 hippocampal NMDARs in contextual fear conditioning, spatial learning, and synaptic plasticity (Kutsuwada et al., 1996; Tsien et al., 1996; Tang et al., 1999; Liu et al., 2004; Lau and Zukin, 2007; Yashiro and Philpot, 2008; Brigman et al., 2010). These data indicate that JIP1-mediated JNK activation may constrain synaptic plasticity, learning, and memory through attenuation of NMDAR function. Furthermore, decreases in JIP1 level and/or localization affecting JNK activity, perhaps resulting from distinct signaling pathways (e.g., glutamate-mediated downregulation of JIP1 level in growth cones, Dajas-Bailador et al., 2014; Ca²⁺-dependent degradation of JIP1, Allaman-Pillet et al., 2003), would thus be predicted to reduce this constraint, leading to enhanced learning and memory.

The increased NMDAR signaling caused by loss of JIP1-mediated JNK activation in JIP1 TA mice is associated with increased expression of the NMDAR subunits GluN2A and GluN2B (Fig. 7). This increase in the levels of NMDAR subunits is significant because it is established that changes in GluN2A and GluN2B expression cause altered plasticity and memory (Tang et al., 1999; Brigman et al., 2010; Chao et al., 2013). This may be mediated by extending the integration time window for NMDAR signaling coincident with presynaptic and postsynaptic activity and de-

creasing the threshold for inducing long-term synaptic changes. Indeed, a constraint by JIP1-JNK on plasticity thresholds may, in turn, regulate information processing and learning (Kiyama et al., 1998; Hawasli et al., 2007; Hu et al., 2007). This is consistent with the observation that loss of JIP1-mediated JNK activation in JIP1 TA mice enables the establishment of LTP at lower stimulation frequencies (Fig. 5H), with the converse being a requirement for higher stimulation frequencies needed in the presence of JIP1-JNK activation. The mechanism of JIP1-dependent regulation of NMDAR subunit expression (Fig. 7A,B) remains to be determined and may include changes in NMDAR membrane insertion, internalization, or lateral movement into synapses (Fig. 14). In addition, JIP1 may regulate NMDAR subunit expression through a posttranscriptional mechanism (Fig. 14). Indeed, it is known that GluN1, GluN2, and GluN2B protein expression can be regulated by CPEB3 (Chao et al., 2013); that GluN2A protein expression can be regulated by CPEB1 (Udagawa et al., 2012; Swanger et al., 2013); and that GluN2B expression can be regulated by a microRNA (Harraz et al., 2012). Strikingly, the learning and memory phenotypes of Cpeb3 -/- mice associated with increased NMDAR expression (Chao et al., 2013) are similar to the phenotypes of the mice with defects in JIP1 function examined in the present study (JIP1 TA and JIP1 DIBD mice).

It is possible that increased expression of NMDAR subunits only partially accounts for the learning and memory phenotypes of JIP $^{\rm TA}$ and JIP1 $^{\rm \Delta JBD}$ mice. Indeed, it has been shown that cFyn mediates phosphorylation of the NMDAR subunit GluN2B on Y¹⁴⁷², resulting in increased NMDAR activity (Salter and Kalia, 2004) by attenuating NMDAR internalization (Roche et al., 2001; Prybylowski et al., 2005), increasing the proper localization of the GluN2B NMDARs at synapses (Nakazawa et al., 2006), and enhancing GluN2B NMDAR-mediated currents at CA1 synapses (Yang et al., 2012). In the present study, increased GluN2B Y 1472 phosphorylation resulted from disruption of JIP1-mediated JNK activation in JIP1 TA mice (Fig. 7A), also perhaps contributing to the observed increase in NMDAR signaling. This change in GluN2B Y 1472 phosphorylation may be caused by JIP1-mediated recruitment of cFyn (Kennedy et al., 2007) or by JNK-mediated recruitment of PSD-95/Fyn complex (Kim et al., 2007). Another potential contributing factor may be the binding of JIP1 to LRP8, a protein that regulates NMDAR signaling (Stockinger et al., 2000; Beffert et al., 2005). Finally, it is possible that JIP1-mediated interactions with the exchange factors Ras-GRF1 and Tiam1 may contribute to increased NMDAR-dependent activation of the ERK pathway and activity-dependent actin remodeling critical for synaptic plasticity and memory (Buchsbaum et al., 2002; Krapivinsky et al., 2003; Tolias et al., 2007).

Contextual fear learning recruits both the hippocampus and amygdala, whereas cued fear learning relies on the amygdala (Phillips and LeDoux, 1992). JIP TA and JIP1 DIPTE mice displayed enhancement in both contextual and cued fear conditioning (Figs. 10, 13). Therefore, these learning enhancements suggest that JIP1-mediated JNK activation is also important in the amygdala. Although JIP1 is expressed in amygdala (unpublished preliminary observations), JNK signaling and the importance of JIP1 in the amygdala have yet to be investigated thoroughly. We hypothesize that the observed contextual-learning enhancement is at least in part due to a lack of hippocampal JIP1-mediated JNK signaling. Not only is signal transduction altered in this region, but JIP TA and JIP1 DIPD mice also demonstrate an improved spatial water maze memory, which classically relies on the hippocampus.

Genetic anomalies within the JNK pathway have also been associated with a subset of other psychiatric disorders (Coffey,

2014). However, the degree to which and the mechanism by which JNK is involved is unknown. JIP1 TA mutant mice display a range of behaviors, including exaggerated fear responses to cues associated with the danger, difficulty suppressing fear behavior even when these cues no longer predict danger, elevated acoustic startle response, and anxiety-like behaviors that may represent rodent homologs of the symptoms that are diagnostic for trauma- and stressor-related disorders, such as posttraumatic stress disorder (PTSD) (Shalev et al., 2017). These responses may be regulated by JIP1-mediated JNK signaling in the hippocampus, the amygdala, or in various cortical regions that interconnect to form the neural circuits that promote adaptation to stress and fear conditioning. Interestingly, we have shown that intrahippocampal infusion with a JNK inhibitor prevents stress-induced changes in fear conditioning (Sherrin et al., 2010). Therefore, it is possible that drugs that target the function of JIP1 to regulate JNK activity or NMDAR function positively (Myers and Davis, 2007; Feder et al., 2014; Ori et al., 2015; Mataix-Cols et al., 2017) may therefore be useful for the treatment of PTSD or anxiety disorders marked by abnormal fear learning and maladaptive processing of information related to threat. Our study provides a proof-of-concept that validates this approach using a model organism. An exciting future possibility is the application of this strategy to the treatment of human fear and anxiety.

Overall, the results of this study suggest that JNK activation caused by the JIP1 scaffold protein constrains learning and memory in an NMDAR-dependent fashion. This role of JIP1 starkly differs from the related protein JIP2 that acts to promote NMDAR signaling by a JNK-independent mechanism (Kennedy et al., 2007). Our studies of JIP1 therefore establish a role for the JIP1-JNK pathway in NMDAR-dependent regulation of memory acquisition, consolidation, and retention.

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