Postsynaptic protein synthesis is required for presynaptic enhancement in persistent forms of long-term potentiation

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INTRODUCTION

Long-term potentiation (LTP) in the hippocampus is a fundamental process underlying learning and memory in the brain. At CA3-CA1 synapses, three discrete forms of LTP (LTP1, 2, and 3) have been differentiated on the basis of their persistence, maintenance mechanisms, Ca²⁺ signaling pathways, expression loci, and electrophysiological requirements. We previously showed that LTP2 and LTP3 involve a presynaptic expression component that is established in a translation-dependent manner. Here we investigate the locus of translation required for presynaptic expression. Neurotransmitter release rate was estimated via FM 1-43 destaining from CA3 terminals in hippocampal slices from male Wistar rats (6–8 weeks). Destaining was measured at sites making putative contact with CA1 dendritic processes in stratum radiatum that had been filled with a membrane impermeable translation inhibitor and a fluorescent indicator. Our results suggest that inhibition of postsynaptic translation eliminates the enhanced release ordinarily observed at 160 min post-LTP induction, and that this effect is limited to sites closely apposed to the filled postsynaptic cell. We conclude that postsynaptic translation is required for the presynaptic component of LTP2 and LTP3 expression. These data considerably strengthen the mechanistic separation of LTP1, 2, and 3 and provide evidence for an expanded repertoire of communication between synaptic elements.

Keywords: LTP, expression, presynaptic, hippocampus, electrophysiology
enhanced exocytosis associated with LTP2 and 3 is dependent on protein synthesis within the postsynaptic cell.

**MATERIALS AND METHODS**

**ETHICAL APPROVAL**

Experiments were performed in accordance with the Australian National University Animal Experimentation Ethics Committee guidelines.

**SLICE PREPARATION AND ELECTROPHYSIOLOGY**

Male Wistar rats (6–8 weeks) were anaesthetized with isoflurane, decapitated, and the brains submerged in ice-cold dissecting solution (mM): 124 NaCl, 3.2 KCl, 1.25 NaH2PO4, 26 NaHCO3, 0.5 CaCl2, 7 MgCl2, 2 ascorbate, 3 pyruvate and 10 D-glucose, equilibrated with 95% O2 – 5% CO2. Transverse hippocampal slices (300 μm) were prepared using a vibratome and area CA3 was removed to reduce potential hyperexcitability. Slices were transferred to a holding solution as above, except the Ca2+, Mg2+, and d-glucose concentrations were adjusted to 2.5 mM, 1.3 mM, and 25 mM respectively. Slices were maintained at 34°C for at least 30–40 min, then at room temperature for a further 30 min, or until required.

Slices were perfused in a continuous flow (~2 ml min⁻¹) of recording solution (as per holding solution, but without ascorbate and pyruvate) at 32–34°C. Whole-cell voltage-clamp recordings from CA1 pyramidal cells were made using glass electrodes (4–6 MΩ) filled with (mM): 127 KMeSO4, 8 KC1, 10 Hepes, 10 sodium phosphocreatine, 4 Mg float, 0.4 Na-GTP, 0.05 Alexa Fluor 488 and in some cases 0.0035 gelonin (Enzo Life Sciences). Somatic responses were evoked by stimulation of the Schaffer collateral/commissural axons (0.1 ms-pulse-width) with a Teflon-insulated tungsten bipolar electrode placed 100–200 μm from the cell of interest. The stimulation amplitude was adjusted in current-clamp to produce EPSPs approximately 1/3rd of action potential threshold. EPSC recordings (200–300 pA) were then obtained with somatic membrane potential clamped at −65 mV.

Baseline recordings (10 min) were initiated no later than 5 min after whole-cell configuration was obtained. Baseline synaptic responses were evoked by delivery of stimuli at a frequency of 0.066 Hz. Every 2.5 min a second pulse was administered 20 ms after the first to enable measurements of paired-pulse facilitation (PPF) and the paired-pulse ratio (PPR). Series resistance was monitored on-line and recordings were terminated if this varied by more than 20% from the baseline value, or alternatively rose above 25 MΩ. LTP was induced in current-clamp by theta-burst stimulation (TBS), consisting of trains of 10 × 100 Hz bursts (5 pulses/burst) with a 200 ms interburst interval (at the test pulse intensity). One train of ten bursts is denoted 1TBS. When multiple trains were delivered (i.e., 4TBS and 8TBS) they were separated by 30 s. See Figure 1A for pictorial representation of the induction stimulus. Synaptic responses were monitored for at least 160 min post-LTP induction, with a success rate of maintaining stable recordings over this period of approximately 15%.

Unless otherwise stated, two-tailed Student’s t-tests were used to determine statistical significance at the 95% confidence level.

**FM 1-43 LOADING AND UNLOADING**

After recording stable Schaffer collateral-evoked EPSCs for 160 min post-TBS, synaptic vesicles were loaded by delivering a train of 1200 stimuli (at 10 Hz) in the presence of FM 1-43 (5 μM, Biotium Inc.) and D-APV (100 μM, Tocris; Figure 1B). The dye application continued for 1 min beyond the duration of the loading stimulus to ensure completion of endocytosis and dye uptake. FM 1-43 was then washed out in the presence of ADVASEP-7 (0.5 mM, Biotium Inc.) for 20 min to remove any extracellularly bound dye (Kay et al., 1999). Terminals making putative contact with spines along an Alexa Fluor 488-labeled dendritic shaft (100–200 μm from the soma) were identified and destaining was achieved by stimulating at 1 Hz for 6 min in the presence of ADVASEP-7 and D-APV. Results from preliminary experiments demonstrated that this protocol maximally unloaded the dye at a rate that was dependent on Ca2+, temperature and stimulation frequency, consistent with it being a valid measure of exocytosis (data not shown).

**TWO-PHOTON IMAGING**

Fluorescence imaging (of labeled release sites and filled dendritic processes) was performed using a Zeiss LSM 510 two-photon laser-scanning microscope with a water immersion objective (Achroplan 40×/0.75 N.A.). Two-photon excitation was achieved with a Ti:Sapphire laser tuned to 810 nm. Imaging at greater depths than is normally ideal for monitoring FM 1-43 release was necessary in these experiments in order to track dendrites at distances of approximately 150–250 μm from the soma. All fields imaged were typically 70–100 μm deep, and were 40–80 μm away from the stimulating electrode. A series of 5–10 images (512 × 512, 0.15 mμm/pixel in the x-y axes) were taken at 0.2–0.4 μm intervals in the z-plane. These z-stacks were repeated every 30 s, beginning 1 min prior to and during the unloading stimulation. In offline analyses, maximal z-projections were generated from each image series to create one image per 30 s time point. Circular regions of interest were defined around the center of brightly stained punctate fluorescent spots and 2–16 terminals and 3 background regions were measured at each time point. A smaller number of puncta were sampled here than is usually possible with FM 1-43 loading as a result of the depth at which imaging was performed. The distance from each punctum to the nearest spine/dendritic shaft was also measured. Only puncta that satisfied four predetermined criteria were included in analysis. These were: (1) diameter of 0.5–1 μm; (2) approximately circular shape; (3) minimal x-y movement (such that any puncta that moved beyond the specified region of interest was excluded); and (4) activity-dependent destaining (i.e., no spontaneous destaining) that was well fit by a first order exponential decay function. A fluorescence time-course was generated by subtracting the average background fluorescence at each time point then normalizing the fluorescence of each punctum at each time point to the average of the prestimulus values. Photobleaching was corrected for by normalizing to the average background fluorescence at the corresponding time point throughout the unload stimulation. The halftime of decay of fluorescence intensity during unloading (t1/2) was calculated for each punctum from single exponential decay curves fitted to the 6 min of stimulus-induced destaining.
RESULTS
LTP2 AND LTP3 ARE DEPENDENT ON POSTSYNAPTIC TRANSLATION FOR THEIR PERSISTENCE

In order to establish the locus of translation that is required for the persistence of LTP2 and LTP3, individual CA1 neurons were filled with a membrane impermeable translation inhibitor (gelonin, 3.5 μM). Gelonin inhibits protein synthesis by inactivating the 60S ribosome (Stirpe et al., 1980) and has previously been used to inhibit local postsynaptic protein synthesis required for learning-related facilitation (Sherff and Carew, 2004; Villareal et al., 2007) as well as local presynaptic translation required for synaptic plasticity in nerve-muscle cultures (Zhang and Poo, 2002).

LTP2 was significantly inhibited by the inclusion of gelonin in the whole-cell patch pipette (Figure 1B). Since LTP is usefully classified on the basis of persistence, the post-TBS data were fit with a double exponential decay function as described previously (Raymond et al., 2000; Raymond and Redman, 2002). The mean time-constant of decay of the slower exponential (τ) was used as a measure of LTP persistence and any instances in which data were not well fit were excluded from any further analysis. LTP2 τ was significantly reduced from 99 ± 4 min (n = 6) to...
34 ± 2 min (n = 9; p < 0.05) with gelonin in the patch pipette. The amplitude of the EPSC in the last 5 min of recording was also significantly reduced with the inclusion of gelonin, from 49 ± 1% (n = 6) to 17 ± 1% (n = 9; Figure 1D; p < 0.05). This is consistent with previously reported data demonstrating a role for translation in the persistence of LTP2 (Raymond et al., 2000; Johnstone and Raymond, 2011). Similarly, both LTP3 persistence and final magnitude were reduced by inhibition of translation with gelonin (Figure 1C). Interestingly, the initial magnitude of both LTP2 and LTP3 was decreased with gelonin in the patch pipette; a result that may provide insight into the critical role of early translation immediately post-LTP. A τ value for 8TBS LTP could not be determined in these experiments because all control LTP was essentially non-decremental over the recording period and could not be fit by exponential decay curves. We therefore restricted our analysis to comparison of LTP magnitude over the last 5 min of recording. Gelonin significantly reduced LTP3 magnitude at 160 min post-8TBS (Figure 1E; Control = 80 ± 6%, n = 4; gelonin = 5 ± 7%; n = 5; p < 0.05). Although inhibition of translation with gelonin also reduced the LTP2 and LTP3 EPSC amplitude immediately post-induction, we conclude that this is unlikely to be due to local toxicity effects of the drug. In the absence of TBS, EPSC amplitudes were unaffected by gelonin in the patch pipette over the entire recording period (n = 4; Figure 1B) and gelonin had no effect on the induction of translation-independent LTP1 (n = 4; Figure 1F). Importantly, application of gelonin extracellularly (100 μM) did not affect the persistence of LTP2 and LTP3 (n = 3, Figures 1B,C), confirming gelonin as a membrane-impermeable compound and supporting the postsynaptic specificity of effects described above. These results suggest that inhibition of protein synthesis in the postsynaptic compartment is sufficient to curtail the expression and maintenance of LTP2 and LTP3.

**PRESYNAPTIC LTP EXPRESSION REQUIRES POSTSYNAPTIC TRANSLATION**

To determine whether postsynaptic translation is also required for the enhanced presynaptic release associated with LTP2 and LTP3, FM 1-43 destaining was measured from CA3 terminals at various distances from CA1 apical dendrites filled with Alexa Fluor 488 and gelonin. This technique allowed for comparisons of presynaptic function between distant terminals and those in putative contact with the gelonin-treated cell. Figure 3 shows an example of a CA1 neuron filled with Alexa Fluor 488 (Figure 2A) and of FM 1-43 filled CA3 terminals surrounding filled postsynaptic dendritic branches (Figure 2B). Our previous data suggests that...
our sampled puncta were representative of CA3 terminals, and did not include large numbers of GABAergic inhibitory interneurons or astrocytes (Johnstone and Raymond, 2011). An example time series of FM 1-43 release under our conditions can be seen in Figure 2C.

In control slices an increased rate of exocytosis was measured at all terminals following induction of LTP2 with 4TBS, regardless of distance to the dendrite (n = 40 terminals, six slices; Figures 3A,B). With gelonin in the postsynaptic recording pipette enhanced release was abolished only at sites very close to the postsynaptic cell (0–0.5 μm; n = 84 terminals, nine slices; Figures 3A,C; One-Way ANOVA, \( F(4, 77) = 35.02; \ p < 0.05 \)). Enhanced exocytosis remained intact at sites further removed from the treated postsynaptic cell. Example data depicting the time course of destaining from individual puncta with and without gelonin are shown in Figure 3D. The enhanced

**FIGURE 3 | Enhanced presynaptic function associated with LTP2 is dependent on postsynaptic translation.** (A) Scatter plot showing the half-life of FM 1-43 destaining versus distance to filled dendrite for individual terminals in slices stimulated with 4TBS with and without gelonin in the patch pipette. (B) Histogram summarizing destaining rate data from the “Control” group in (A) and from “No TBS” slices with distances binned. Destaining rate was enhanced in slices stimulated with 4TBS and there was no effect of distance to nearest postsynaptic dendrite in these untreated cells. (C) Histogram of data obtained from 4TBS gelonin-treated group in (A) and “No TBS” slices treated with gelonin, with distances binned. Inclusion of gelonin in the patch pipette abolished enhanced release induced by 4TBS only at sites close to (0–0.5 μm) the filled postsynaptic cell, but had no effect on release in the absence of TBS (One-Way ANOVA, \( \ast p < 0.05 \)). (D) Examples of destaining from two representative puncta in the 0.0–0.5 μm distance group. i, control terminal with no gelonin in the pipette shows normal, LTP-related enhanced rate of dye release. ii, terminal with gelonin in the postsynaptic cell shows reduced rate of dye release.
exocytosis associated with LTP3 was also found to be dependent on postsynaptic translation. In control slices enhanced release was observed following 8TBS regardless of distance to the nearest filled dendrite ($n = 39$ terminals; four slices; Figures 4A,B). However, inhibition of postsynaptic translation with gelonin ($n = 34$ terminals, five slices) blocked enhanced release at 160 min post-8TBS only at distances between 0 and 0.5 μm to the nearest filled dendrite (Figures 4A,C; One-Way ANOVA, $F(4, 29) = 37.06; p < 0.05$). As was the case with LTP2, enhanced release remained unaffected at sites further removed from the treated postsynaptic cell. Importantly, destaining rate in the absence of TBS remained unchanged both with ($n = 39$ terminals, four slices) and without gelonin ($n = 17$ terminals, three slices), regardless of distance to the nearest filled CA1 dendrite (Figures 3B,C).

**PAIRED-PULSE FACILITATION DATA SUPPORT A ROLE FOR PRESYNAPTIC EXPRESSION THAT IS DEPENDENT ON POSTSYNAPTIC TRANSLATION**

As a complementary experiment, levels of PPF were recorded with and without gelonin before and at various time points following induction of LTP2 and 3. A reduction in the PPR is thought to be indicative of increased $Pr$ (Schulz et al., 1994, 1995; Sokolov et al., 1998, 2002), therefore this analysis was valuable as it allowed for comparison of release probability with FM 1-43 destaining at 80 min and 160 min post-TBS.

LTP2 was associated with a reduction in PPR at 160 min after induction (PPR Baseline = 2.13 ± 0.07, $n = 5$ cells; PPR 160 min = 1.26 ± 0.05, $n = 5$ cells; $p < 0.05$), but not at 80 min (PPR 80 min = 1.82 ± 0.28, $n = 5$ cells; Figures 5A,C). LTP3 was associated with a reduction in PPR at both 80 min (PPR Baseline = 1.85 ± 0.08, $n = 4$ cells; PPR 80 min = 1.28 ± 0.01, $n = 4$ cells) and 160 min post-TBS (PPR 160 min = 1.28 ± 0.005, $n = 4$ cells; Figures 5A,D; $p < 0.05$). Importantly, the PPR remained stable over the entire recording period in the absence of TBS (Figure 5B). These data are consistent with our previously reported FM 1-43 destaining data, and support the finding that LTP2 involves a presynaptic component that is not expressed until 80–160 min post-TBS, whereas LTP3 involves a presynaptic component that may be recruited earlier and is apparent as early as 80 min post-TBS.

In further accordance with the FM 1-43 destaining data, inclusion of gelonin in the patch pipette internal solution abolished the elevated $Pr$ that was associated with LTP2.

**FIGURE 4 | Enhanced presynaptic function associated with LTP3 is dependent on postsynaptic translation.** (A) Scatter plot showing the half-life of FM 1-43 destaining versus distance to filled dendrite for individual terminals in slices stimulated with 8TBS with and without gelonin in the patch pipette. (B) Histogram summarizing destaining rate data obtained from the “Control” group in (A) except with distances binned. There was no effect of distance on mean half-life of destaining in untreated cells. (C) Histogram summarizing data obtained from “gelonin” group in (A) except with distances binned. Inclusion of gelonin in the patch pipette abolished enhanced release induced by 8TBS only at sites close to the treated postsynaptic cell (One-Way ANOVA, *$p < 0.05$*).
Presynaptic LTP requires postsynaptic translation

FIGURE 5 | PPR measurements demonstrate increased release probability with LTP2 and LTP3. (A) Paired-pulse ratio (PPR) in slices stimulated with 4TBS and 8TBS (at time point indicated by arrow) showing a reduction in the ratio over time for both groups. (B) PPR was stable in the absence of TBS. (C) Bar graph showing mean PPR during the first 10 min of recording (Baseline) and at 80 min and 160 min post-4TBS. PPR was significantly reduced at 160 min post-4TBVS, consistent with the enhanced FM 1-43 destaining data for LTP2. (D) Bar graph as for (C) except for LTP3 data induced with 8TBS. PPR was significantly reduced at both 80 min and 160 min post-TBS, consistent with the FM 1-43 destaining data for LTP3. * Indicates 0.05

(ppr baseline = 1.60 ± 0.07, n = 9 cells; ppr 80 min = 1.56 ± 0.05, n = 9 cells; Figures 6a,b) and LTP3 (PPR baseline = 1.57 ± 0.02, n = 5 cells; PPR 80 min = 1.57 ± 0.10, n = 5 cells; PPR 160 min = 1.45 ± 0.14, n = 5 cells; Figures 6a,c). These results support FM 1-43 destaining data suggesting that persistent forms of LTP recruit a presynaptic mode of expression that is dependent on translation in the postsynaptic compartment.

Discussion

LTP1, 2, and 3 have previously been differentiated on the basis of persistence, Ca\(^{2+}\) signaling mechanisms, expression locus, electrophysiological requirements, and maintenance mechanisms (Abraham and Otani, 1991; Raymond et al., 2000; Raymond and Redman, 2002, 2006; Raymond, 2007, 2008; Johnstone and Raymond, 2011). The findings described here allow for further expansion of the LTP1, 2, and 3 model and provide additional evidence that certain classes of synapses are capable of supporting multiple discrete forms of LTP.

We now show that the enhanced exocytosis and P\(_r\) associated with LTP2 and LTP3 is eliminated in the presence of a translation inhibitor in the postsynaptic cell. Enhanced release was only abolished at terminals very closely apposed to the postsynaptic spines of treated neurons, whereas it remained intact at sites more distal to the treated dendrite. These data suggest the effects of gelonin were indeed confined to the treated cell and selectively affected the LTP response of terminals making synaptic contact with that cell.

Synaptic cascade from postsynaptic activity to enhanced presynaptic release

Dendritic translation initiated by both 4TBS- and 8TBS-evoked Ca\(^{2+}\) signals is likely to result in the synthesis of many plasticity-related proteins, one or more of which appears to be required for the presynaptic enhancement. Considering the dependence of presynaptic expression on NO signaling (Johnstone and Raymond, 2011), it is possible that translation of NO-related signaling proteins, such as NOS, could be important. NOS expression is known to be upregulated under certain stress conditions (De Oliveira et al., 2000; Yao et al., 2007), but has yet to be demonstrated during hippocampal synaptic plasticity. Furthermore, NOS mRNA has not been detected in CA1 dendrites and to date has only been observed in developing olfactory (Gibson and Nighorn, 2000) and optic (Koriyama et al., 2009) axons. Additionally, most reports suggest that NO exerts its influence only during induction of LTP (Schuman and Madison, 1991), therefore it seems likely that NO signaling will be upstream of the postsynaptic translation. Indeed, existing evidence suggests that NO is an important regulator of mRNA stability, acting to either stabilize or destabilize certain genes (Bouton and Demple, 2000; Ma et al., 2004). Under these conditions NO may play a critical

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FIGURE 6 | Inhibition of postsynaptic translation with gelonin abolishes the enhanced Pr associated with LTP2 and LTP3. (A) PPR data in slices stimulated with 4TBS and 8TBS (at time point indicated by arrow) with gelonin in the patch pipette internal solution. (B) Bar graph showing mean PPR during the first 10 min of recording (Baseline) and at 80 min and 160 min post-4TBS. PPR remained unchanged for the duration of the recording period. (C) Bar graph as for (B), except for LTP3 data induced with 8TBS. PPR did not change over the entire recording period.

role in regulating turnover of mRNA transcripts that are required to trigger changes in the kinetics of release.

These transcripts will likely encode proteins that are capable of signaling to the presynaptic cell, for example cell–cell adhesion molecules which are known to be crucial for some forms of LTP (Dityatev et al., 2008) and are important for regulating exocytosis (Saghatelyan et al., 2004; Dityatev et al., 2008). Alternatively, dendritic translation of proteins that act as a second retrograde signal might be important. Given the existing data supporting a role for BDNF in presynaptic expression of TBS-induced LTP (Zakharenko et al., 2003), the preponderance of BDNF transcripts in hippocampal dendrites (Tongiorgi et al., 1997; An et al., 2008) and the observation that translation-dependent forms of LTP require BDNF (Bramham and Messaoudi, 2005) it is likely that postsynaptic translation of BDNF is a key process. Two recent studies provide further insight into the putative steps in the cascade. Henry et al. (2012) document the activity of the translation promoter mTOR complex 1 (mTORC1) in driving an activity-dependent, retrograde signaling cascade in hippocampal neurons that promotes enhanced neurotransmitter release from closely apposed presynaptic terminals. They further identify BDNF as a key factor acting downstream of mTORC1 activation in establishing enhanced presynaptic function. Complementary to this, Penney and colleagues (2012) found that overexpression of TOR at the Drosophila neuromuscular junction is sufficient to trigger a strong retrograde increase in neurotransmitter release. These studies, in combination with the data we present here, suggest that postsynaptic activation of mTORC1 and subsequent retrograde action of BDNF is a likely pathway by which neurotransmitter release is enhanced during persistent forms of LTP.

EXPERIMENTAL CONSIDERATIONS

There are a small number of considerations that must be taken into account when interpreting the data presented here. Firstly, the results do not exclude a role for presynaptic translation in the expression of LTP2 and LTP3. It is possible that translation in the postsynaptic cell may trigger changes in the presynaptic terminal leading to further presynaptic translation of key proteins.

Secondly, it should be noted that gelonin has previously only been used as an inhibitor of synaptic translation in Xenopus (Zhang and Poo, 2002) and Aplysia (Sherff and Carew, 2004; Villareal et al., 2007) neuronal preparations, therefore, we cannot discount the possibility that it produces its inhibitory effects on LTP via different means in rodent hippocampus. It is also possible that gelonin may exert a local toxic effect which inhibits LTP induction in a manner that is independent of translation. Other non-toxic effects of gelonin might also be important. For example, gelonin may alter circuit excitability by interfering with sodium currents in the somatic compartment and/or reducing the amplitude or number of back-propagating action potentials during the induction stimulation. We consider these possibilities unlikely, however, given the qualitatively similar results we
achieved with the membrane-permeable translation inhibitor anisomycin (Johnstone and Raymond, 2011) and the null effect of gelonin on LTP1 (Figure 1F) and normal synaptic transmission (Figure 1B).

Thirdly, the reduced magnitude of initial LTP2 and LTP3 that occurs with gelonin in the pipette (Figures 1B,C) merits a brief discussion. A possible explanation for this result is that gelonin might be acting upon products of so-called “immediate early genes.” For example, c-fos is known to be activated within approximately 5 min of stimulation (Flavell and Greenberg, 2008), and could conceivably have an effect on EPSC size. Other activity-regulated genes that directly regulate circuit connectivity may also play an important role in this early phase of gene expression. For example candidate plasticity gene 15 (Cpg15) is known to be regulated by activity and induces rapid AMPA receptor insertion, as well as initiating other changes that increase synaptic efficacy (Nedivi et al., 1998; Cantallops et al., 2000).

A final aspect of the experimental design that is important to consider is the rapid manner in which gelonin exerts its inhibition on LTP. Because of the “wash-out” effect associated with the degree and persistence of potentiation at a synapse, allowing only afferent input that induces sufficient postsynaptic Ca2+ to recruit changes in presynaptic release. It has been known for some time that protein synthesis is required for LTP to persist beyond a few hours. Our findings suggest that part of this requirement is related to the recruitment of a presynaptic expression mechanism that itself is strongly correlated with LTP persistence.

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