

Expanded View Figures

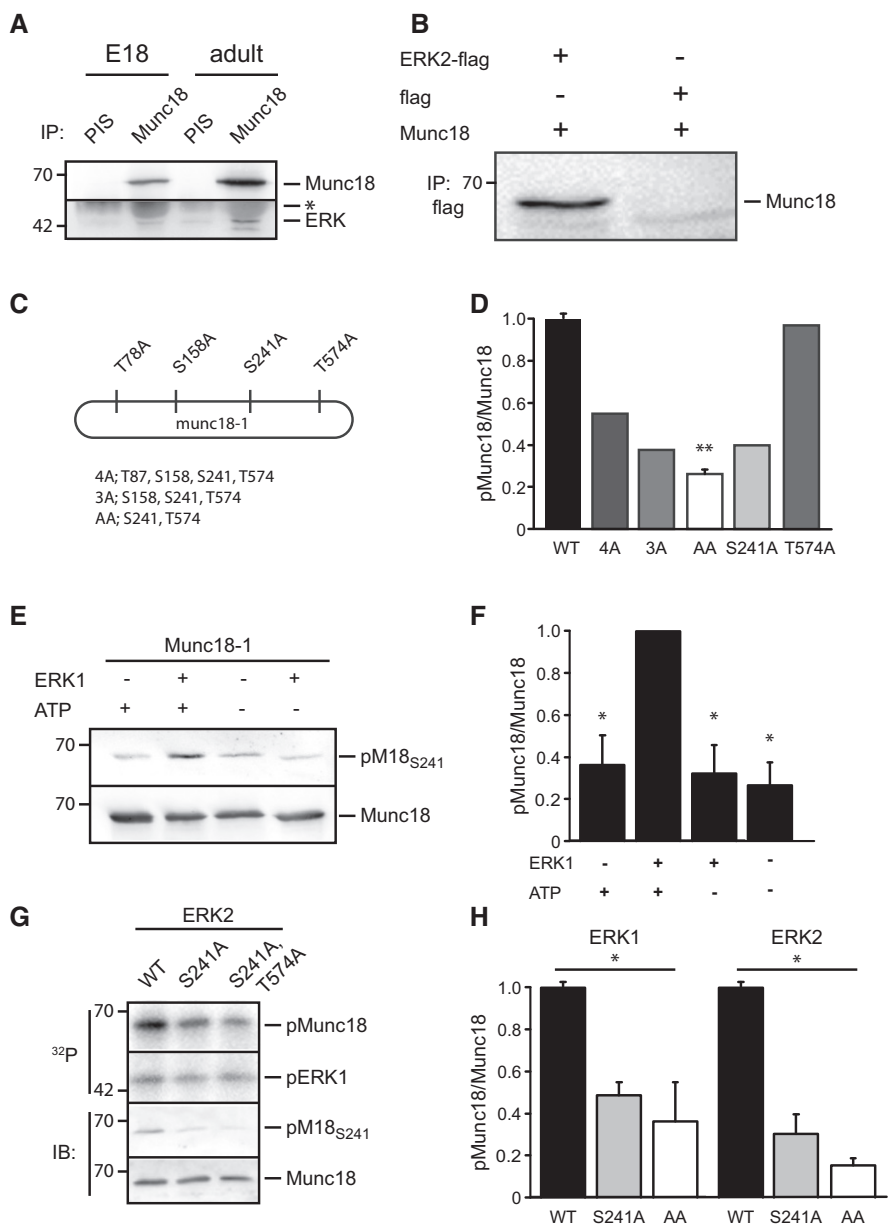


Figure EV1. ERK1/2 binds and phosphorylates Munc18-1.

Related to Figure 1.

A, B Binding between Munc18-1 and ERK in brain and in HEK293T cells. (A) Co-immunoprecipitation (IP) of ERK and Munc18-1 from embryonic (E18) or adult brain lysate with Munc18-1 antibody, but not with pre-immune serum (PIS). The immunoglobulin heavy chain is visible (*) above the ERK band. (B) Co-immunoprecipitation from HEK293T cells transfected with Munc18-1 and ERK2-flag or empty-flag (as negative control) with flag antibody. Western blot is stained for Munc18-1.

C Diagram of Munc18-1 showing 4 potential ERK phosphorylation sites (P-X-S/T-P). These sites were mutated to alanine yielding 4A (T87A, S158A, S241A, and T574A), 3A (S158A, S241A, and T574A), AA (S241A and T574A), T574A and S241A.

D Quantification of radioactive phospho-Munc18-1 intensity in different Munc18-1 mutants from Fig 1B. Phosphorylation was measured as the amount of ³²P incorporation in Munc18-1 normalized to total Munc18. Phosphorylation of Munc18_{WT} was set as 100% (Munc18_{WT}: 100 ± 3%, n = 3; Munc18_{4A}: 55%, n = 1; Munc18_{3A}: 38%, n = 1; Munc18_{AA}: 24 ± 2%, n = 3, **P < 0.01; Munc18_{S241A}: 36%, n = 1; Munc18_{T574A}: 100%, n = 1), one-way ANOVA.

E–H Test of phospho-specific Munc18_{S241} antibody. (E) *In vitro* ERK kinase assay, HEK293T cells were transfected with biotinylated Munc18-1. Protein complexes were precipitated with streptavidin-coated magnetic beads and subjected to *in vitro* ERK kinase assay with recombinant ERK1. Western blots were stained with a phospho-specific antibody directed against pS241 (pM18_{S241}) and Munc18-1 antibody. (F) Quantification of (E).

Phosphorylation was measured as ratio of pM18_{S241} over total Munc18. Condition ERK1 plus ATP was set as 100% (n = 3, *P < 0.05), one-way ANOVA. (G) *In vitro* ERK kinase assay, HEK293T cells were transfected with M18_{WT}, M18_{S241A,T574A}, and M18_{S241A}. Munc18-1 variants were immune-precipitated and subjected to *in vitro* kinase assay with recombinant ERK2. Radioactive phosphoblot and immunoblot of *in vitro* ERK kinase assay are stained with pM18_{S241} and Munc18-1 antibody. (H) Quantification of phosphorylation as pM18_{S241} intensity in (G) normalized to total Munc18. Phosphorylation of M18_{WT} was set to 100% (n = 3 for ERK1 and ERK2 *in vitro* assays, *P < 0.05), one-way ANOVA.

Data information: Data are expressed as mean ± SEM.

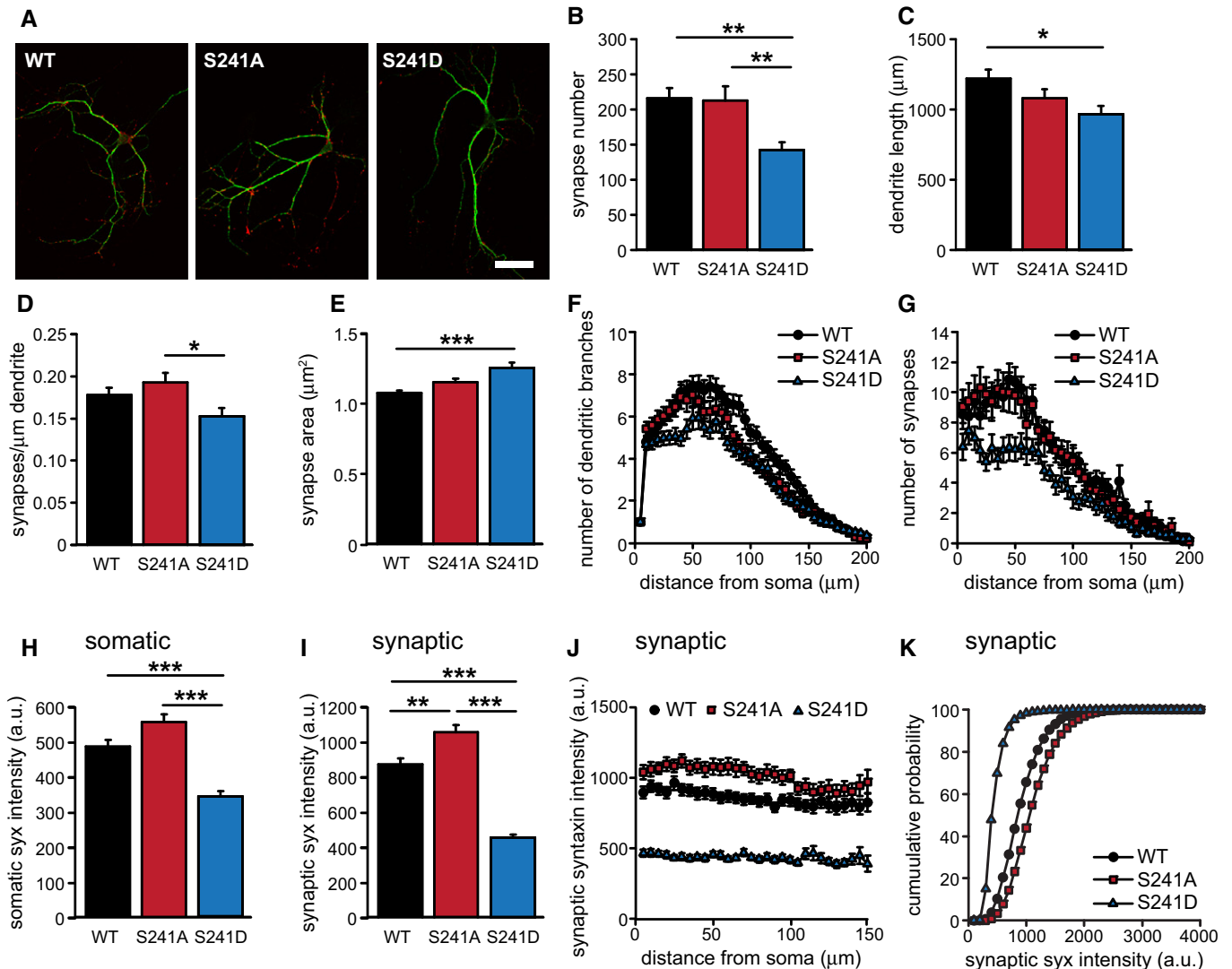


Figure EV2. Neuronal morphology of rescued Munc18-1 null mutant cells.

Related to Figure 4. Hippocampal autaptic neurons of Munc18-1 null mutant mice were rescued with lentiviruses expressing M18_{WT}, M18_{S241A}, or M18_{S241D} and stained for dendritic MAP2, synapse marker VAMP2, and syntaxin.

A Typical examples. MAP2 (green), VAMP2 (red). Scale bar represents 50 μm .

B Mean synapse number per cell (M18_{WT}: 216 ± 15 , $n = 51$; M18_{S241A}: 214 ± 21 , $n = 46$; M18_{S241D}: 143 ± 11 , $n = 40$; $**P < 0.01$, one-way ANOVA).

C Mean dendrite length per cell (M18_{WT}: 1226 ± 61 μm , $n = 51$; M18_{S241A}: 1084 ± 65 μm , $n = 46$; M18_{S241D}: 971 ± 59 μm , $n = 40$; $*P < 0.05$, one-way ANOVA).

D Number of synapses per μm dendrite length (M18_{WT}: 0.179 ± 0.008 , $n = 51$; M18_{S241A}: 0.193 ± 0.012 , $n = 46$; M18_{S241D}: 0.153 ± 0.010 , $n = 40$; $*P < 0.05$, one-way ANOVA).

E Mean synapse area (M18_{WT}: 1.08 ± 0.02 μm^2 , $n = 51$; M18_{S241A}: 1.16 ± 0.03 μm^2 , $n = 46$; M18_{S241D}: 1.26 ± 0.04 μm^2 , $n = 40$; $***P < 0.001$, one-way ANOVA).

F Number of dendritic branches as a function of radial distance from the soma quantified by Sholl analysis.

G Number of synapses as a function of radial distance from the soma.

H Mean somatic syntaxin intensity (M18_{WT}: 485 ± 19 , $n = 49$; M18_{S241A}: 554 ± 23 , $n = 46$; M18_{S241D}: 343 ± 14 , $n = 39$; $***P < 0.001$, one-way ANOVA).

I Mean synaptic syntaxin intensity (M18_{WT}: 875 ± 35 , $n = 51$; M18_{S241A}: $1,058 \pm 42$, $n = 46$; M18_{S241D}: 458 ± 19 , $n = 40$; $**P < 0.01$; $***P < 0.001$, one-way ANOVA).

J Mean synaptic syntaxin intensity as a function of radial distance from the soma.

K Cumulative probability plot of synaptic syntaxin intensity.

Data information: Data are expressed as mean \pm SEM.

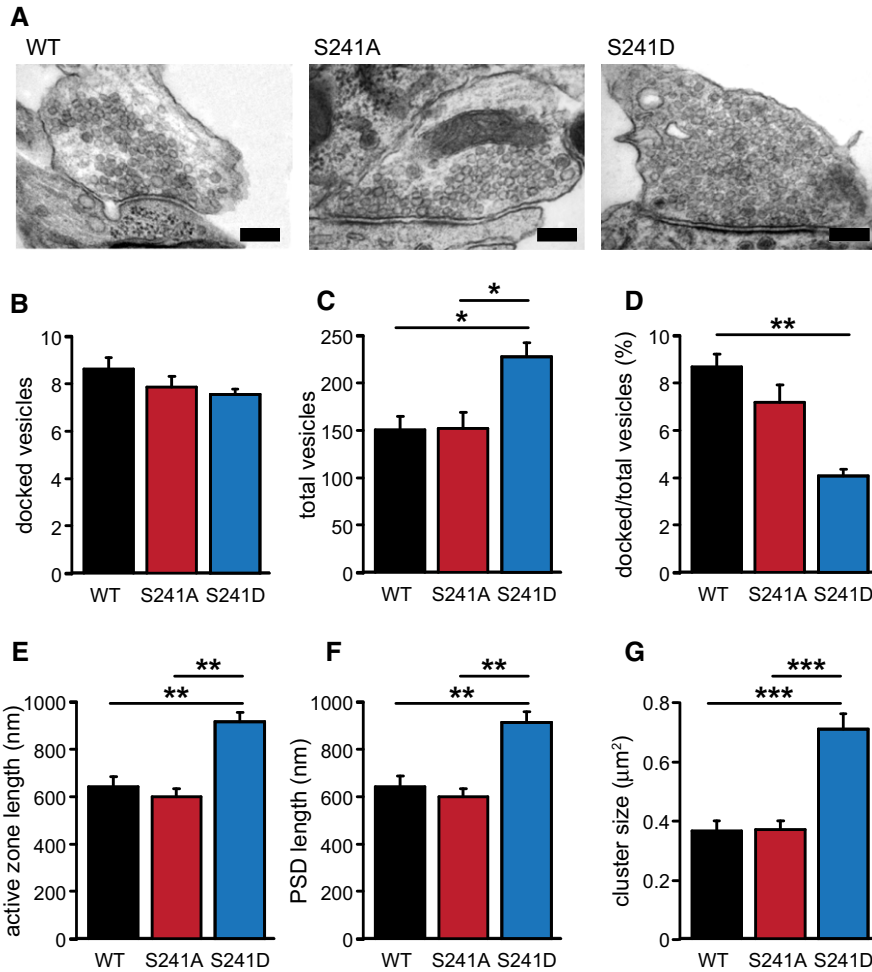


Figure EV3. Ultra-structural morphology of rescued Munc18-1 null mutant cells.

Related to Figure 4. Electron microscopy analysis of hippocampal autaptic neurons of Munc18-1 null mutant mice rescued with lentiviruses expressing M18_{WT}, M18_{S241A}, or M18_{S241D}. *N*_{cultures} = 3, M18_{WT}: *n*_{cells} = 6, *n*_{synapses} = 138; M18_{S241A}: *n*_{cells} = 5, *n*_{synapses} = 146; M18_{S241D}: *n*_{cells} = 3, *n*_{synapses} = 77. Synapses were averaged per cell. Graphs represent cell averages.

- A Typical examples. Scale bar represents 200 nm.
- B Mean number of docked vesicles (M18_{WT}: 8.64 ± 0.48; M18_{S241A}: 7.89 ± 0.44; M18_{S241D}: 7.57 ± 0.22, n.s., one-way ANOVA). *n* = 3.
- C Mean number of total vesicles (M18_{WT}: 151.99 ± 14.28; M18_{S241A}: 152.65 ± 16.54; M18_{S241D}: 228.16 ± 15.86; **P* < 0.05, one-way ANOVA). *n* = 3.
- D Percentage of docked vesicles out of total number of vesicles (M18_{WT}: 8.69 ± 0.55; M18_{S241A}: 7.20 ± 0.74; M18_{S241D}: 4.09 ± 0.28; ***P* < 0.01, one-way ANOVA). *n* = 3.
- E Mean active zone length (M18_{WT}: 643.12 ± 43.84 nm; M18_{S241A}: 601.62 ± 33.45 nm; M18_{S241D}: 918.24 ± 39.47 nm; ***P* < 0.01, one-way ANOVA). *n* = 3.
- F Mean postsynaptic density (PSD) length (M18_{WT}: 644.70 ± 43.74 nm; M18_{S241A}: 603.05 ± 33.49 nm; M18_{S241D}: 914.09 ± 47.52 nm; ***P* < 0.01, one-way ANOVA). *n* = 3.
- G Mean cluster size (M18_{WT}: 2.97 ± 0.15 μm²; M18_{S241A}: 3.00 ± 0.11 μm²; M18_{S241D}: 4.40 ± 0.23 μm²; ****P* < 0.001, one-way ANOVA). *n* = 3.

Data information: Data are expressed as mean ± SEM.

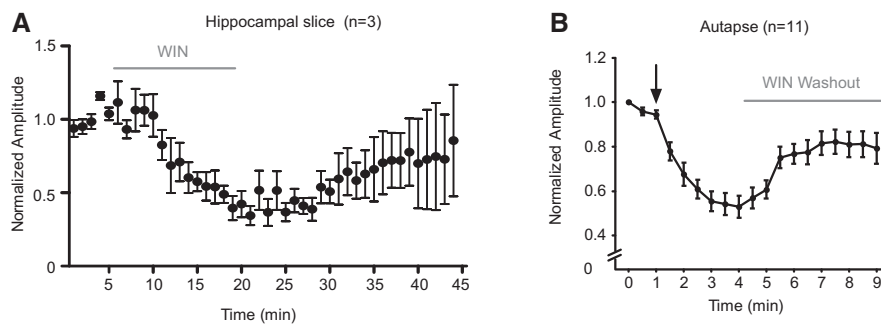


Figure EV4. WIN55,212-2-induced depression of transmission is reversible.

Related to Figure 7.

- A Normalized EPSC amplitudes in hippocampal CA1 neurons during and after a 15-min application of 5 μM WIN55,212-2 (gray bar) shows the gradual recovery of EPSCs after WIN washout (*n* = 3 slices).
- B Normalized EPSC amplitudes in wild-type hippocampal autapses during and after 30-s application of 1 μM WIN55,212-2 shows recovery during WIN washout (gray bar) (*n* = 11 neurons).

Data information: Data are expressed as mean ± SEM.

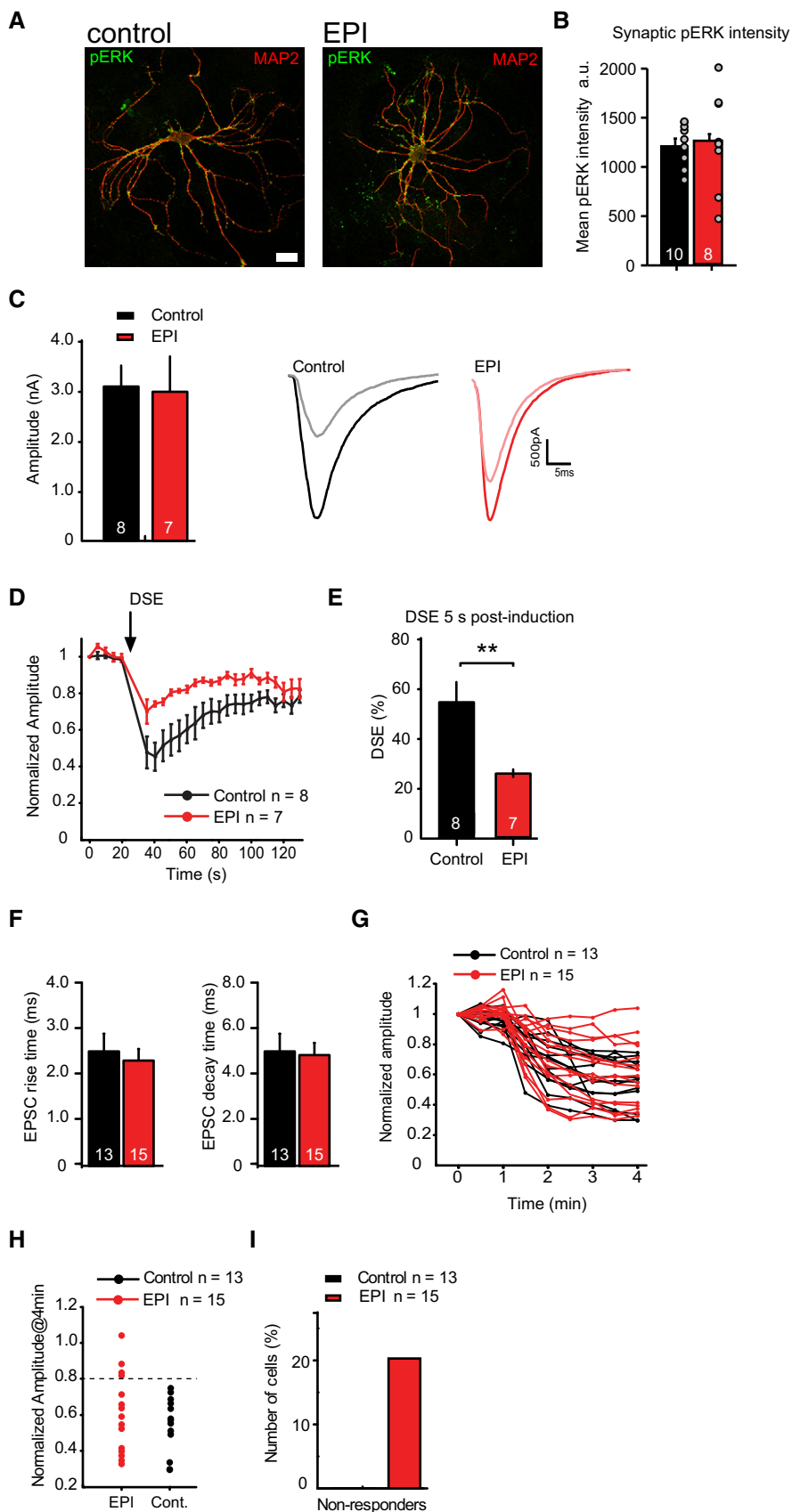


Figure EV5. ERK pathway inhibition decreases depolarization-induced suppression of excitation (DSE) and WIN5,212-2-induced depression in a subset of cultured neurons.

Related to Figure 7.

A Typical examples of isolated neurons pretreated with ERK inhibitors (EPI) or DMSO (control) for 3 h. Dendritic marker MAP2 (red), phospho-ERK antibody (pERK, green). Scale bar represents 10 μ m.

B Semi-quantitative analysis of phospho-ERK intensities at synapses (identified by VAMP2, not shown) shows that pretreatment with ERK inhibitors does not result in reduction of phospho-ERK levels in all neurons (control $n = 10$ neurons, EPI $n = 8$ neurons). In these neurons, ERK appears to remain in a phosphorylated state much longer than in hippocampal slices (compare with Fig 3C).

C–E DSE duration and amplitude is decreased in hippocampal autaptic neurons upon ERK pathway inhibition compared to control. **(C)** Left: Average EPSC amplitude in wild-type autaptic neurons before a 10-s depolarizing pulse in control and ERK pathway inhibitors (EPI: 20 μ M PD98059 and 10 μ M U0126, applied 3 h before recordings) pretreated neurons is similar (control: 3.1 ± 0.4 nA, $n = 8$; EPI: 3.0 ± 0.7 nA, $n = 7$). Right: Examples of average EPSC traces for control (black and gray) and EPI (red and pink)-treated neurons prior to and after DSE, respectively. **(D)** Time course of normalized EPSC amplitudes recorded before and after a 10-s depolarizing pulse to 0 mV (DSE, indicated by the arrow) in wild-type autaptic neurons in the absence (control) or presence of ERK pathway inhibitors (EPI) (control $n = 8$, EPI $n = 7$ from 2 independent experiments). **(E)** DSE effect size in control and EPI-treated autapses 5 s after a 10-s depolarizing pulse to 0 mV (control $n = 8$, EPI $n = 7$, $**P < 0.01$, Mann–Whitney U -test).

F–H ERK pathway inhibitors block effect of CB1 agonist WIN5,212-2 in a subset of hippocampal autaptic neurons. **(F)** EPSC rise and decay times of control (black bars) and EPI (red bars)-treated neurons prior to WIN5,212-2 application are similar. **(G)** Effect of 1 μ M WIN application in individual cells after 4 min (start of application is at $t = 1$ min) in the absence (control, $n = 13$) and presence of ERK pathway inhibitors (EPI: 20 μ M PD98059 and 10 μ M U0126, applied 3 h before recordings, $n = 15$). **(H)** Normalized EPSC amplitudes 4 min after WIN application in control and EPI neurons. The dashed line represents cutoff used to distinguish non-responders (EPSC rundown < 20%) and responders (EPSC rundown > 20%) in **(G)**.

I Ratio of non-responding versus responding neurons to 1 μ M WIN application shows that in 20% of EPI-treated neurons, EPSC during WIN application remained above 80% of EPSC before WIN stimulation. In control cells, WIN application resulted in > 20% reduction of EPSCs in all neurons tested.

Data information: Data are expressed as mean \pm SEM.