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# Phosphorylation of Ser1166 on GluN2B by PKA Is Critical to Synaptic NMDA Receptor Function and Ca<sup>2+</sup> Signaling in Spines

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The NMDA-type glutamate receptor (NMDAR) is essential for synaptogenesis, synaptic plasticity, and higher cognitive function. Emerging evidence indicates that NMDAR Ca<sup>2+</sup> permeability is under the control of cAMP/protein kinase A (PKA) signaling. Whereas the functional impact of PKA on NMDAR-dependent Ca<sup>2+</sup> signaling is well established, the molecular target remains unknown. Here we identify serine residue 1166 (Ser1166) in the carboxy-terminal tail of the NMDAR subunit GluN2B to be a direct molecular and functional target of PKA phosphorylation critical to NMDAR-dependent Ca<sup>2+</sup> permeation and Ca<sup>2+</sup> signaling in spines. Activation of  $\beta$ -adrenergic and D<sub>1</sub>/D<sub>5</sub>-dopamine receptors induces Ser1166 phosphorylation. Loss of this single phosphorylation site abolishes PKA-dependent potentiation of NMDAR Ca<sup>2+</sup> permeation, synaptic currents, and Ca<sup>2+</sup> rises in dendritic spines. We further show that adverse experience in the form of forced swim, but not exposure to fox urine, elicits striking phosphorylation of Ser1166 *in vivo*, indicating differential impact of different forms of stress. Our data identify a novel molecular and functional target of PKA essential to NMDAR-mediated Ca<sup>2+</sup> signaling at synapses and regulated by the emotional response to stress.

## Introduction

NMDARs are glutamate-gated ion channels and are enriched at excitatory synapses, where they are strategically positioned to play a crucial role in regulation of synaptic function (Cull-Candy and Leszkiewicz, 2004; Lau and Zukin, 2007; Traynelis et al., 2010). A unique feature of NMDARs is their high permeability to Ca<sup>2+</sup>. Ca<sup>2+</sup> influx through NMDARs is essential for synaptogenesis, plasticity of neural circuitry, and higher cognitive functions, such as learning and memory. NMDARs are tetrameric assemblies of GluN1 and GluN2 subunits, which cotranslationally assemble in the endoplasmic reticulum to form functional chan-

nels. In adult hippocampus and neocortex, GluN2A and 2B are the primary GluN2 subunits expressed at synapses (Gray et al., 2011). The precise subunit composition governs the physiological and pharmacological properties of NMDARs. GluN2B-containing receptors exhibit slower decay kinetics (Yashiro and Philpot, 2008), carry more Ca<sup>2+</sup> per unit charge (Sobczyk et al., 2005), and preferentially tether to the synaptic plasticity protein CaMKII (Barria and Malinow, 2005; Halt et al., 2012). In addition, GluN2B expression can enhance the magnitude of LTP, promote hippocampal-based learning, plasticity-associated spine growth, and dendritic patterning critical to information processing (Tang et al., 1999; von Engelhardt et al., 2008).

Synaptic NMDARs localize to the postsynaptic density (PSD) where they are structurally organized and spatially restricted in a large macromolecular signaling complex comprised of scaffolding and adaptor proteins, which physically link receptors to kinases, phosphatases, and other downstream signaling proteins (Sheng and Hoogenraad, 2007). Protein kinase A (PKA) is a major kinase known to regulate NMDAR trafficking and gating. At synaptic sites, PKA is kept in close proximity to NMDARs by association of the PKA regulatory subunit with A-kinase anchoring proteins (AKAPs), such as AKAP9 and AKAP5 (Sanderson and Dell'Acqua, 2011).

Emerging evidence reveals that PKA signaling represents a fundamental mechanism by which NMDAR-mediated Ca<sup>2+</sup> influx is modulated in neurons. Direct activation of PKA promotes NMDAR Ca<sup>2+</sup> permeability, Ca<sup>2+</sup> signaling in dendritic spines, and induction of LTP at Schaffer collateral to CA1 pyramidal cell

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The authors declare no competing financial interests.

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synapses (Skeberdis et al., 2006). Consistent with this, extracellular signals that modulate cAMP bidirectionally regulate  $Ca^{2+}$  permeation through NMDARs and  $Ca^{2+}$  transients in spines in a PKA-dependent manner. Whereas GABA<sub>B</sub> and dopamine D<sub>2</sub> receptors negatively couple to the cAMP/PKA cascade and inhibit NMDAR-mediated  $Ca^{2+}$  signaling in spines, 2A-type adenosine receptors positively engage PKA signaling, enhance NMDAR  $Ca^{2+}$  permeation, and  $Ca^{2+}$  signaling in spines (Chalifoux and Carter, 2010; Higley and Sabatini, 2010). In addition, activation of  $\beta$ -adrenergic receptors enhances NMDA currents (Raman et al., 1996), and activation of D<sub>1</sub>/D<sub>5</sub> dopamine receptors facilitates induction of LTP at CA1 synapses in a PKA-dependent manner (Otmakhova and Lisman, 1996).

Whereas the functional impact of PKA on NMDAR-dependent  $Ca^{2+}$  signaling is well characterized, the molecular target remains unknown. Using a combination of site-directed mutagenesis, electrophysiology, two-photon microscopy, glutamate uncaging, and  $Ca^{2+}$  imaging, we identify Ser1166 in the carboxy-terminal tail of GluN2B as a novel molecular target of PKA phosphorylation critical to  $Ca^{2+}$  signaling in dendritic spines and regulated by the emotional response to stress.

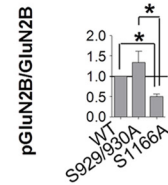
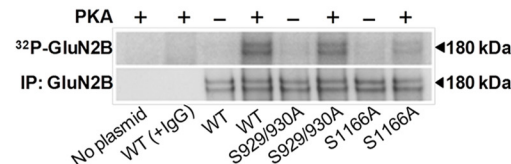
## Materials and Methods

**Animals.** All animal procedures were in accordance with guidelines approved by the Albert Einstein, Harvard University, and University of California Institutional Animal Care and Use Committee and federal guidelines.

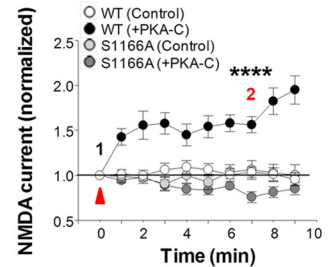
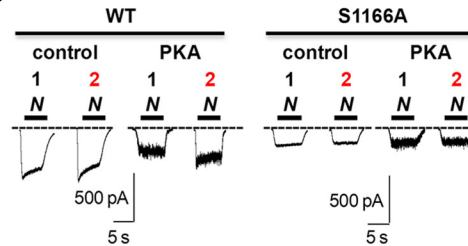
**Constructs.** GluN1–1a and GluN2B(WT) in pRK5 were a kind gift from M.D. Ehlers (Duke University, Durham, North Carolina). The S929/930A and S1166A GluN2B mutants were generated using the QuikChange XL Site-Directed Mutagenesis Kit (Stratagene) according to the manufacturer's instructions and verified by DNA sequencing. The following primers from 5' to 3' were used for GluN2B(S929/930A): forward, GACTTCATCCGCCGAGAGGCCAGCCGTCTACGACATCTCTG, and reverse, CAGAGATGTCGTAGACGGCTGCTCTCGGCGGATGAAGTC. The following primers from 5' to 3' were used for GluN2B(S1166A): forward, GACTTCAAGCGAGATGCGGTCTAGTGGAGGTG, and reverse, CACTCCACTGACCCGATCTCGCTTGAAGTC.

**In vitro phosphorylation assay.** To assess PKA-dependent phosphorylation of NMDARs, HEK293 cells were transfected with WT and mutant GluN2B with Lipofectamine 2000 (Invitrogen) as described previously (Lin et al., 2004). Forty-eight to 72 h after transfection, lysates were immunoprecipitated with an antibody to GluN2B (Millipore), and immunocomplexes were incubated with PKA-C (Sigma). Samples were run on 3–8% Tris-acetate gels (Invitrogen) for subsequent autoradiography and Western blotting (adapted from Leonard and Hell, 1997).

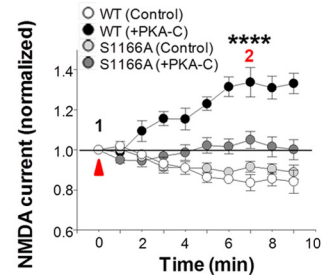
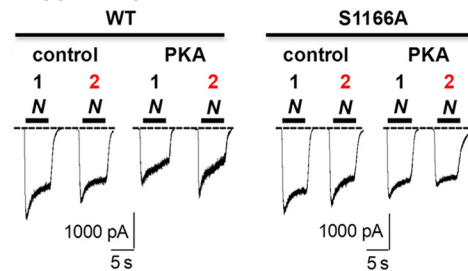
### a HEK293 cells



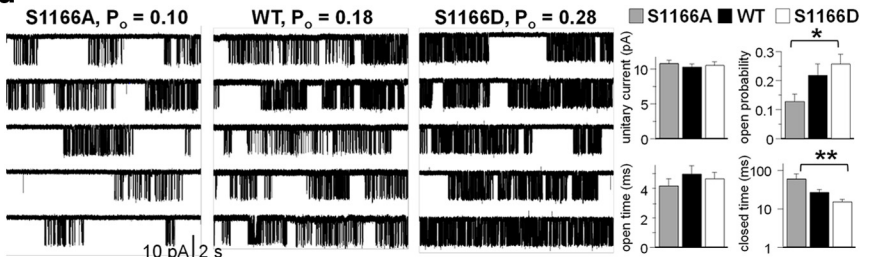
### b HEK293 cells



### c Hippocampal neurons



### d



**Figure 1.** Ser1166 in GluN2B is a direct target of PKA phosphorylation critical to PKA potentiation. **a**, HEK293 cells expressing WT or mutant GluN2B subunits were immunoprecipitated and incubated with PKA-C in the presence of  $\gamma$ -<sup>32</sup>P-ATP. Top, Autoradiogram depicting incorporation of  $\gamma$ -<sup>32</sup>P into GluN2B. Bottom, Western blot probed with an antibody to GluN2B. Right, Summary data. Values for mutant receptors were normalized to values for WT receptors. **b**, PKA potentiation of NMDA-evoked currents is abolished in HEK293 cells expressing GluN1 with GluN2B(S1166A); bar above sample records represents duration of NMDA + glycine application (*M*). Left, Sample traces at 0 (1) and 7 (2) min. Right, Summary time course data of the peak NMDAR current, defined as the initial maximum current. Potentiation is defined as the ratio of the peak current recorded from the same cell at 7 min versus 0 min. Values for WT and mutant receptors in the presence (PKA-C) and absence (Control) of PKA-C were normalized to values for WT receptors in the absence of PKA-C (control). WT + PKA-C, *n* = 9 cells; S1166A + PKA-C, *n* = 11 cells; WT control, *n* = 7 cells; S1166A control, *n* = 6 cells from 4 to 9 independent experiments. **c**, PKA potentiation of NMDA-evoked currents is abolished in hippocampal neurons expressing GluN2B(S1166A). Left, Representative traces. Right, Summary time course data as in **b**. WT + PKA-C, *n* = 5 cells; S1166A + PKA-C, *n* = 8 cells; WT control, *n* = 5 cells; S1166A control, *n* = 5 cells from 3 or 4 independent experiments. **d**, Far left, Representative traces for individual GluN1/GluN2B(S1166A) (left), WT (center), and S1166D (right) channels. Low gating activity of S1166A relative to that of WT is restored with the phosphomimetic mutation, S1166D. Single-channel activity was recorded with cell-attached pipettes as inward currents through one S1166A, WT, or S1166D channel expressed in a HEK293 cell. Far right, Summary of single-channel parameters for S1166A (gray, *n* = 14), WT (black, *n* = 19), and S1166D (white, *n* = 16). The apparent differences in *P*<sub>o</sub> and closed time values for either the S1166A or S1166D mutant versus WT were not significant, consistent with variability in the basal phosphorylation state of WT GluN1/2B receptors. \**p* < 0.05. \*\**p* < 0.01. \*\*\*\**p* < 0.0001. Here and in all figures, error bars indicate SEM.

**Whole-cell patch-clamp recording.** Primary cultures of hippocampal neurons were generated from embryonic day (E)18 Sprague Dawley rats and transfected at 5–7 days *in vitro* (DIV) with WT or mutant GluN2B and GFP by means of calcium phosphate as described previously (Jiang et al., 2004). To minimize NMDAR-induced toxicity, cells were maintained in the presence of the NMDAR antagonist 2R-amino-5-phosphonovaleric

**Table 1. Summary of statistics for Figures 1–3**

Figure	N	Mean ± SEM	p	Statistical test
1a	4 experiments	WT: 1.000 ± 0.0 S929/930A: 1.306 ± 0.284 S1166A: 0.4616 ± 0.0802	*, 0.0218	One-way ANOVA with Dunn's post-test
1b	WT = 9 cells S1166A = 11 cells WT (control) = 7 cells S1166A (control) = 6 cells	WT: 1.584 ± 0.0763 S1166A: 0.9028 ± 0.0234 WT (control): 1.025 ± 0.0126 S1166A (control): 0.9935 ± 0.0152	****, < 0.0001	One-way ANOVA with Tukey's post-test (WT vs S1166A, WT vs WT (control), and WT vs S1166A (control))
1c	WT = 5 cells S1166A = 8 cells WT (control) = 5 cells S1166A (control) = 5 cells	WT: 1.073 ± 0.013 S1166A: 0.8984 ± 0.0145 WT (control): 0.8991 ± 0.0220 S1166A (control): 0.9249 ± 0.0132	****, < 0.0001	One-way ANOVA with Tukey's post-test (WT vs S1166A, WT vs WT (control), and WT vs S1166A (control))
1d	WT = 20 cells  S1166A = 14 cells  S1166D = 16 cells	WT: 1, 10.4 ± 0.3; P <sub>o</sub> , 0.22 ± 0.04; MOT, 5.0 ± 0.5; MCT, 27.6 ± 4.5 S1166A: 1, 10.9 ± 0.4; P <sub>o</sub> , 0.13 ± 0.02; MOT, 4.2 ± 0.5; MCT, 61.2 ± 19.3 S1166D: 1, 10.6 ± 0.5; P <sub>o</sub> , 0.26 ± 0.03; MOT, 4.7 ± 0.4; MCT, 15.7 ± 2.1	P <sub>o</sub> *, 0.0317 MCT:**, 0.0097	One-way ANOVA with Tukey's post-test (P <sub>o</sub> , S1166A vs S1166D; MCT, S1166A vs S1166D)
2c	WT = 6 cells S1166A = 6 cells	WT: ΔE <sub>rev</sub> = -6.79 mV S1166A: ΔE <sub>rev</sub> = 0.46 mV	*, 0.0207	Two-tailed paired t test for WT vs S1166A
2f	WT = 7 cells S1166A = 5 cells	WT % inhibition: 59.14 ± 7.20% S1166A % inhibition: 20.20 ± 12.58%	*, 0.0165 **, 0.0030	Two-tailed unpaired t test for % inhibition by H-89 of WT vs S1166A; one-way ANOVA with Dunn's post-test (WT control vs H-89)
3e	WT (control) = 23 cells WT (ISO) = 26 cells S1166A (control) = 22 cells S1166A (ISO) = 22 cells	WT (control): 17.03 ± 1.58 WT (ISO): 29.34 ± 2.21 S1166A (control): 18.09 ± 1.94 S1166A (ISO): 20.25 ± 1.95	*, < 0.001	Two-tailed unpaired t test for WT (control) vs WT (ISO)
3f	WT (control) = 24 cells WT (ISO) = 27 cells S1166A (control) = 22 cells S1166A (ISO) = 22 cells	WT (control): -9.677 ± 1.048 WT (ISO): -16.69 ± 1.58 S1166A (control): -11.15 ± 1.05 S1166A (ISO): -10.50 ± 1.86	*, < 0.001	Two-tailed unpaired t test for WT (control) vs WT (ISO)

\*p &lt; 0.05; \*\*p &lt; 0.01; \*\*\*\*p &lt; 0.0001.

acid (AP5) (Tocris Bioscience) from the time of transfection until recording. Whole-cell patch-clamp recordings were obtained from hippocampal neurons at 11–15 DIV as described previously (Skeberdis et al., 2006). The external solution contained the following (in mM): 145 NaCl, 2.5 KCl, 2 CaCl<sub>2</sub>, 10 glucose, 10 HEPES buffer, pH 7.4, 0.5 μM TTX (Ascent Scientific) to block action potentials and 200 nM Zn<sup>2+</sup> to preferentially inhibit GluN2A-containing NMDARs (Paoletti et al., 1997; Rodenas-Ruano et al., 2012). Glass electrodes (2–4 MΩ) were filled with internal solution containing the following (in mM): 140 CsMeSO<sub>3</sub>, 1 EGTA, 0.1 CaCl<sub>2</sub>, 0.5 MgCl<sub>2</sub>, 10 HEPES, pH 7.3 with CsOH, 4 Mg-ATP, 0.4 Na<sub>2</sub>-GTP, and 5 phosphocreatine. Voltage-clamp recordings were performed at a holding potential of -60 mV using an Axopatch 200A amplifier; data were filtered at 2 kHz and digitized at 10 kHz. NMDA currents were elicited by application of NMDA (100 μM + 10 μM glycine) in the presence or absence of PKA-C (Calbiochem, Millipore) (250 nM; see Fig. 1; or in Fig. 2, the PKA inhibitor H-89 (10 μM) (Sigma). Rapid agonist application was achieved by stimulating a single cell repeatedly with agonist under computer control from a multibarrel array fed by gravity, which achieved changes in solutions in <100 ms (Lin et al., 2004). Given the speed of this system, the peak as measured may represent an underestimate of the actual peak. Nevertheless, the comparison of the peak current before and after PKA application (and between WT and S1166 mutant) is likely to be valid. The height of the solution column did not change significantly over 10 applications, thereby eliminating any systematic change in the perfusion rate. Time course data represent averaged values of the peak phase of NMDA-elicited currents at each time point for 5–12 cells per condition; values for cells infused with PKA-C were normalized to the corresponding value for control cells infused with recording solution. Data analysis was performed with Clampfit software.

Recording from HEK293 cells was performed at 1–2 d after transfection as described for neurons with the following exceptions: The external solution consisted of the following (in mM): 145 NaCl, 5.4 KCl, 10 HEPES, 11 glucose,

and 2 CaCl<sub>2</sub>, pH 7.4. The internal solution was as follows (in mM): 145 KCl, 5.5 EGTA, 4 Mg-ATP, and 10 HEPES, pH 7.25. Currents were elicited by application of NMDA (300 μM + 10 μM glycine). PKA-C was infused to a final concentration of 100 nM. Cells were washed for 1 min to allow recovery to basal levels before the next stimulation.

**Single-channel recording and analyses.** Currents were recorded and analyzed with the cell-attached patch-clamp technique as described previously (Kussius et al., 2009). The pipette solution contained the following (in mM): 150 NaCl, 2.5 KCl, 1 EDTA, and 10 HEPES and NMDAR agonists glutamate (1 mM) and glycine (0.1 mM). The pipette voltage was clamped at +100 mV (membrane potential ~-120 mV), and inward sodium currents were recorded and amplified (Axopatch 200B), low pass filtered (10 kHz), and sampled (40 kHz, NIDAQ board) into digital files. Only records originating from one-channel patches were retained for analyses. The selected records were idealized with the SKM algorithm in QuB software to calculate average unitary current amplitude, open probability, and durations for open and closed intervals for each record. Statistical significance was evaluated by ANOVA, followed by Tukey's *post hoc* analysis.

**Ca<sup>2+</sup> imaging in HEK293 cells.** To image NMDA-evoked Ca<sup>2+</sup> signals, cells were loaded with the Ca<sup>2+</sup> indicator Fura-2 AM (Invitrogen) in external recording solution or DMEM (1 h at 37°C). NMDA-evoked Ca<sup>2+</sup> rises were elicited by application of NMDA (100 μM + 10 μM glycine) 2 min before, during, and 10 min after application of the PKA inhibitor H-89 (10 μM). Cells were washed for 10–15 min to allow Ca<sup>2+</sup> levels to return to baseline before each stimulation with NMDA and glycine. The ratio of the Ca<sup>2+</sup> signal at 340 to 380 nm was calculated using MetaFluor software.

**Slice culture preparation.** Organotypic hippocampal slice cultures were prepared from 6- to 7-d-old mice as described previously (Carter and Sabatini, 2004). Two days later, DNA constructs were biolistically transfected with a Helios Gene Gun. Bullets were loaded with 50 μg of WT

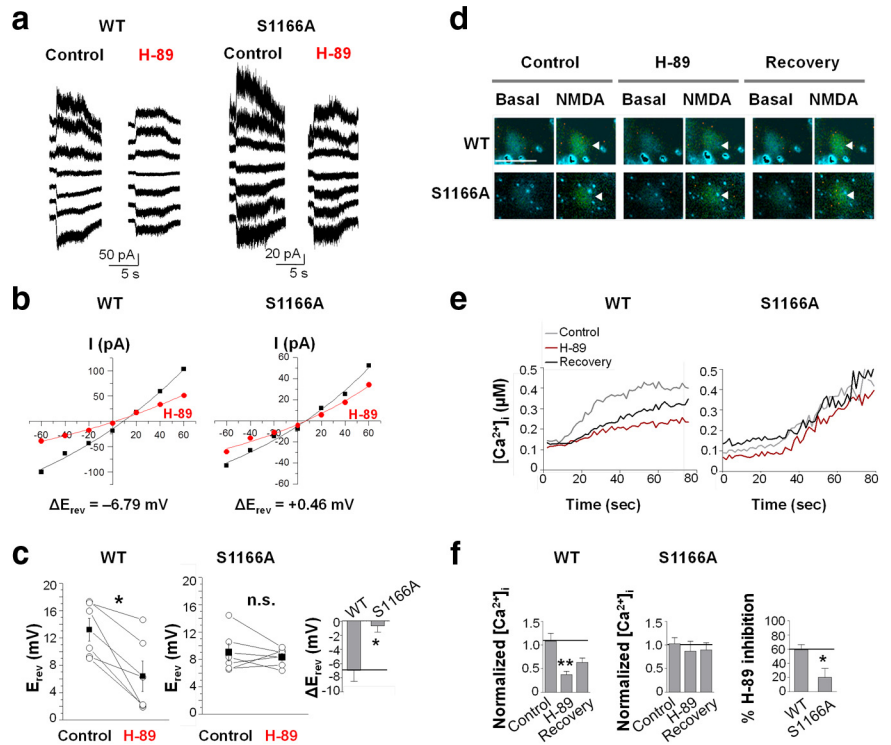
GluN2B or GluN2B(S1166A) and tdTomato (30  $\mu$ g) to assess efficiency of transfection and identify transfected cells.

**Two-photon laser glutamate uncaging.** For two-photon laser uncaging experiments, synaptic  $Ca^{2+}$  transients were measured and uncaging-evoked EPSCs (uEPSCs) were simultaneously recorded at individual spines from tdTomato<sup>+</sup> CA1 neurons expressing wild-type (WT) or mutant GluN2B(S1166A) before and after glutamate photolysis in organotypic hippocampal slice cultures at 10–14 d after transfection. Photolysis of MNI glutamate was performed by focal illumination with a 0.5 ms pulse of light at 720 nm at  $\sim 0.5$   $\mu$ m from the target spine head. In all conditions,  $Mg^{2+}$ -free ACSF (in mM) 127 NaCl, 2.5 KCl, 25  $NaHCO_3$ , 1.25  $NaH_2PO_4$ , 2  $CaCl_2$ , and 25 glucose included 20  $\mu$ M NBQX to block AMPAR-mediated currents, 1  $\mu$ M TTX to block action potentials, and 2.5 mM MNI glutamate to elicit laser-induced glutamate release. Glass electrodes (2–3.5  $M\Omega$ ) were filled with internal solution containing the following (in mM): 130 CsMeSO<sub>4</sub>, 10 HEPES, 1.8  $MgCl_2$ , 4  $Na_2ATP$ , 0.3  $NaGTP$ , and 8 sodium creatine phosphate, 10  $CsCl_2$ , 300  $\mu$ M Fluo-5F, 20  $\mu$ M AlexaFluor-594, pH 7.3. Voltage-clamp recordings were performed at a holding potential of  $-70$  mV using an Axopatch 200A amplifier; data were filtered at 2 kHz and digitized at 10 kHz. Intracellular  $Ca^{2+}$  imaging and glutamate uncaging were performed on a custom microscope, as described previously (Carter and Sabatini, 2004). Neurons were filled via the patch electrode for 15 min before imaging.

For isoproterenol-treated cells, 10  $\mu$ M isoproterenol was added 5 min after break-in. Fluo-5F (green) and AlexaFluor-594 (red) were excited using 840 nm light to monitor  $Ca^{2+}$  signals and spine morphology, respectively. To measure  $Ca^{2+}$  transients in spines, green and red fluorescence was acquired during 500 Hz line scans across a spine and a neighboring (control) spine. Reference frame scans were taken between acquisitions to correct for spatial drift.  $Ca^{2+}$  signals were quantified as increases in green fluorescence from baseline normalized to the average red fluorescence ( $\Delta G/R$ ) and represented as percentage of the  $G/R$  ratio measured in saturating  $Ca^{2+}$  ( $G_{sat}/R$ ).

Imaging and physiology data were acquired using a National Instruments data acquisition boards and custom software written in MATLAB as described previously (Pologruto et al., 2003). Off-line analysis was performed using custom routines written in MATLAB and Igor Pro. For each analyzed dendritic spine, the amplitude of NMDAR-mediated uEPSCs was calculated by averaging all uEPSC traces and determining the peak in the 40 ms time window after uncaging. Only spines from cells with  $R_s < 25$   $M\Omega$  were considered for analysis. For the same spine population, measurements of  $\Delta G/G_{sat}$  were calculated by averaging all  $Ca^{2+}$  transients for each spine and subsequently determining the peak of the average trace by averaging the 20–50 ms after uncaging time window. All graphs and data are expressed as mean  $\pm$  SEM, and significance was assessed using unpaired, two-tailed Student's *t* test.

**Slice treatments.** Acute hippocampal slices were prepared as described previously (Lu et al., 2007) and treated with vehicle ( $H_2O$  or EtOH), 1  $\mu$ M isoproterenol, 1  $\mu$ M isoproterenol + propranolol for 5 min, 10  $\mu$ M SKF81297, or 10  $\mu$ M SKF81297 + 5  $\mu$ M SCH23390 (0.5–15 min). After treatment, slices were shock frozen and extracted with 1% deoxycholate (DOC) in buffer containing the following (in mM): 50 Tris pH 8.5, 10 EDTA, 10 EGTA, and 135 NaCl, and cleared by ultracentrifugation (Leonard and Hell, 1997). All buffers included the following protease and

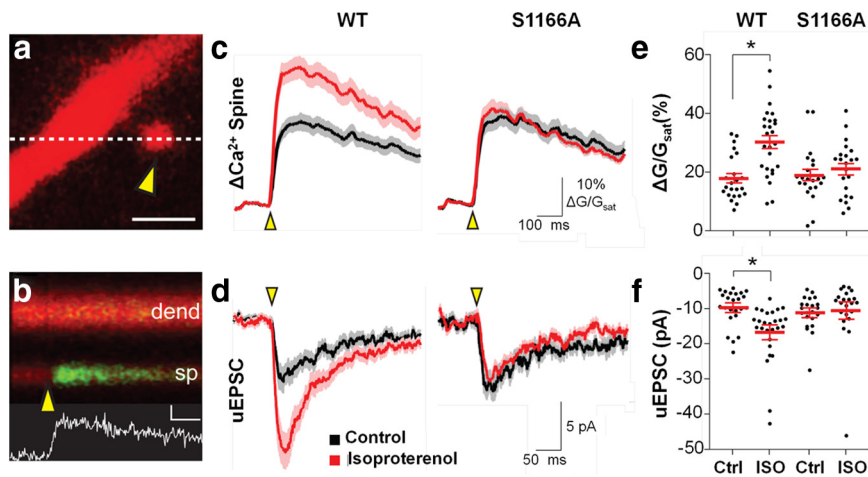


**Figure 2.** Ser1166 is critical for modulation of NMDAR  $Ca^{2+}$  permeability and NMDA-evoked  $Ca^{2+}$  rises by PKA. Representative traces (**a**) and current–voltage relationship (**b**) for NMDA-elicited currents recorded from HEK293 cells expressing GluN1 with WT and mutant GluN2B(S1166A) in the absence and presence of H-89. H-89 induced a negative shift in reversal potential in cells expressing WT ( $\Delta E_{rev} = -6.79$  mV), but not GluN2B(S1166A) ( $\Delta E_{rev} = 0.46$  mV). **c**, Summary data depicting reversal potential ( $E_{rev}$ ); WT,  $n = 6$  cells; S1166A,  $n = 6$  cells from 3 independent experiments.  $*p = 0.0207$ ; n.s., Not significant. **d–f**, PKA potentiation of NMDA-evoked rises in  $Ca^{2+}$  is diminished in HEK293 cells expressing GluN1 with GluN2B(S1166A). **d**, Representative images of basal and NMDA-evoked  $Ca^{2+}$  rises before (control), during (H-89), and after washout of H-89 (recovery) for cells expressing GluN1 with WT or S1166A receptors. **e**, Time course data of cells in **d**. **f**, Left, Summary data for NMDA-evoked  $Ca^{2+}$  rises. Right, Summary data expressed as percentage inhibition by H-89; WT,  $n = 7$  cells; S1166A,  $n = 5$  cells from 5 independent experiments.  $*p = 0.0165$ .  $**p < 0.01$ . Scale bar, 30  $\mu$ m.

phosphatase inhibitors: 1.5  $\mu$ M pepstatin A, 2.1  $\mu$ M leupeptin, 0.3  $\mu$ M aprotinin, 0.2 mM PMSF, 1 mM *p*-nitrophenylphosphate, 25 mM NaPPi, 25 mM NaF, and 2  $\mu$ M microcystin-LR. Immunoprecipitation for GluN2B and GluA1 was as described previously (Lu et al., 2007). Samples were run on 7.5% polyacrylamide gels followed by quantitative immunoblotting with antibodies against pSer1166-GluN2B (PhosphoSolutions), GluN2B, pSer897-GluN1, GluN1, pSer845-GluA1, and GluA1 as described previously (Lu et al., 2007). Chemiluminescent immunosignals were digitized by scanning films and quantified with Adobe Photoshop.

**Fox urine assay.** Postnatal day (P30) rats were placed in a cage containing a 50 ml tube with kimwipes soaked in fox urine (Wildlife Research Supplies) or 0.9% saline for 5 min, and immediately anesthetized with isoflurane and decapitated as described previously (Liu et al., 2010). Hippocampi were isolated and processed for immunoprecipitation with GluN2B and GluA1 antibodies and Western blot analysis as above.

**Injections and forced swim assay.** P24–P32 rats were subjected to forced swim in a 4 L beaker chamber filled to 3 L at a temperature of 16–22°C for 5 min. The chamber was cleaned after every one or two rats. One hour prior to forced swim, rats were anesthetized with isoflurane, stereotaxically injected into the hippocampal CA1 with the PKA inhibitor Rp-8-Br-cAMPS (25 nmol, 4  $\mu$ l) (Enzo Life Sciences) or with saline. The coordinates for injection into the hippocampus were  $-3$  anterior/posterior,  $-2$  medial/lateral, and  $-4$  dorsal/ventral from bregma. Immediately after forced swim, rats were anesthetized with isoflurane, decapitated, and the hippocampi transferred to ice-cold PBS containing protease and phosphatase inhibitor cocktails as above. Hippocampi were microdissected around the injection site, flash-frozen in liquid  $N_2$ , and stored at  $-80^\circ C$ . For Western blot analysis, hippocampi were processed as described above.



**Figure 3.** Ser1166 of GluN2B is essential for  $\beta$ -adrenergic-induced  $\text{Ca}^{2+}$  rises in spines. **a**, Representative 2PLSM image of a dendritic spine from a CA1 hippocampal neuron filled with the red fluorophore AlexaFluor-594 and the green  $\text{Ca}^{2+}$ -sensitive indicator Fluo-5F. The yellow arrowhead indicates the location of a 500  $\mu\text{s}$  pulse of 720 nm laser light used to focally uncage glutamate near the spine head. White dashed line indicates the region used for line scan analysis of fluorescence. Scale bar, 1  $\mu\text{m}$ . **b**, Red and green fluorescence in the spine head (sp) and neighboring dendrite (dend) measured in a line scan over the region indicated by the dashed line in **a**. Inset scale bars, 100 ms and 5%  $\Delta\text{G}/\text{G}_{\text{sat}}$ . Yellow arrowhead indicates time of uncaging. The increase in green fluorescence indicates increased intracellular  $\text{Ca}^{2+}$ , and the corresponding spine  $\text{Ca}^{2+}$  transient (white trace) is below. **c**, Average trace of uncaging-evoked  $\Delta\text{Ca}^{2+}_{\text{spine}}$  produced by single uncaging stimuli in neurons expressing GluN2B(WT) (left) or GluN2B(S1166A) (right) under control conditions (black) or after isoproterenol incubation (red). Mean (solid lines) and mean  $\pm$  SEM (shaded regions) are shown. **d**, Average trace of uEPSCs produced by single uncaging stimuli in neurons expressing GluN2B(WT) (left) or GluN2B(S1166A) (right) under control conditions (black) or after isoproterenol incubation (red). **e**, Population data for peak  $\Delta\text{Ca}^{2+}_{\text{spine}}$  and **(f)** uEPSC elicited by single uncaging stimuli in neurons expressing GluN2B(WT) or GluN2B(S1166A) under control conditions (Ctrl) or after isoproterenol incubation. Red bars represent the mean of distribution  $\pm$  SEM. \* $p < 0.001$ .

**Surface biotinylation assay.** To assess surface expression of GluN2B WT and mutant constructs, HEK293 cells were transfected with GluN1 and WT or mutant GluN2B constructs with Lipofectamine 2000 as above. Forty-eight to 72 h after transfection, total and surface extracts were isolated and processed for Western blots as described previously (Lin et al., 2004). Membranes were probed with antibodies directed to GluN1, GluN2B, and  $\beta$ -actin (loading control) and bands quantified as above.

**Statistical analyses.** Data are represented as mean  $\pm$  SEM. All statistical analyses were performed using Prism software (GraphPad). Significance was defined as  $p < 0.05$ .

## Results

### PKA phosphorylates GluN2B at Ser1166

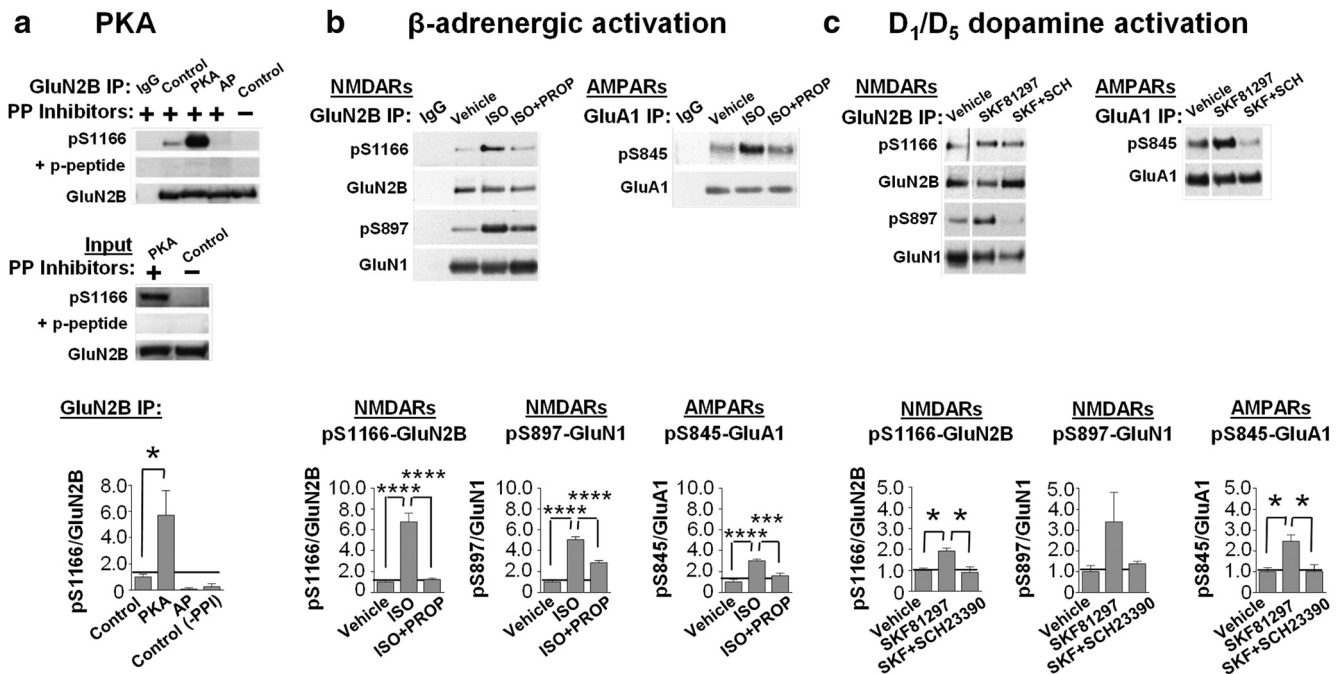
To identify the molecular target of PKA relevant to regulation of NMDAR  $\text{Ca}^{2+}$  permeability and  $\text{Ca}^{2+}$  signaling in spines, we used NetPhosK and scansite.mit.edu software set at medium stringency algorithms to identify putative PKA consensus sequences in the GluN2B carboxy-tail. The GluN2B carboxy-terminal tail sequence harbors three sites that conform to a PKA consensus phosphorylation motif (K/R-K/R-X(X)-S/T): Ser929, Ser930, and Ser1166. We used site-directed mutagenesis to mutate these residues singly or together to the nonphosphorylatable residue alanine. To examine the ability of PKA to phosphorylate GluN2B at these sites, we expressed GluN2B(WT), GluN2B(S929/930A), or GluN2B(S1166A) in HEK293 cells and immunoprecipitated cell lysates with an antibody directed to GluN2B. Immunocomplexes were incubated with the recombinant, constitutively active catalytic subunit of PKA (PKA-C) or vehicle in the presence of  $\gamma$ - $^{32}\text{P}$ -labeled ATP, and samples were subjected to autoradiography and Western blotting (Fig. 1a). In HEK293 cells, GluN2B subunits migrate as doublets, corresponding to the glycosylated and nonglycosylated forms of the receptor (Petralia et al., 1994). Background phosphorylation (defined as phosphorylation in the absence of PKA-C) of WT and mutant receptor subunits was negligible (Fig. 1a). PKA-C in-

duced striking phosphorylation of WT and mutant GluN2B(S929/930A) receptors (Fig. 1a; Table 1). In contrast, PKA-induced phosphorylation of mutant GluN2B(S1166A) receptors was greatly diminished but not abolished (Fig. 1a; Table 1). The small residual signal may reflect additional sites of PKA phosphorylation on GluN2B, at least under the conditions of this assay. WT and mutant receptors exhibited comparable levels of total GluN2B abundance, assessed by Western blot analysis (Fig. 1a), and surface expression, assessed by surface biotinylation assays (data not illustrated). These results indicate that Ser1166 of GluN2B is a main site of PKA phosphorylation.

### PKA potentiation of NMDA-evoked currents is abolished in cells expressing GluN2B(S1166A)

To address whether Ser1166 is a functional target of PKA, we undertook several experimental approaches. First, we examined the impact of Ser1166 on PKA-dependent potentiation of NMDA-elicited currents (300  $\mu\text{M}$  with 10  $\mu\text{M}$  glycine) recorded from HEK293 cells expressing GluN1 together with WT or mutant GluN2B(S1166A) receptors and GFP to enable visualization of transfected cells (Fig. 1b). HEK293 cells are relatively small, geometrically simple, and more nearly isopotential than are neurons and enable examination of NMDAR function in a system expressing a homogeneous population of NMDARs (see Materials and Methods). In these experiments, the same cell was stimulated repeatedly with agonist (once per minute; 5 s per application) for a total of 10 applications under computer control before and after action of PKA-C. PKA-C was delivered directly into the cell via the patch pipette at the time of break-in ( $t = 0$  min). Potentiation was defined as the ratio of peak currents recorded from the same cell at two different time points (7 vs 0 min). In cells expressing WT NMDARs, PKA-C potentiated the peak phase of NMDA-elicited currents by  $\sim 1.5$ -fold (Fig. 1b; Table 1). In contrast, in cells expressing GluN1/GluN2B(S1166A) receptors, the peak to steady-state ratio of the basal current was markedly reduced, and PKA potentiation was abolished (Fig. 1b; Table 1). The peak phase of NMDA-evoked currents recorded from cells expressing WT or mutant receptors and treated with vehicle was not detectably altered within the time course of the experiment.

Next, we addressed whether Ser1166 is a molecular target of PKA relevant to potentiation of NMDA currents by PKA in primary cultures of hippocampal neurons, a more physiologically relevant system to assess NMDAR function. NMDARs and the machinery required for proper NMDAR targeting and function, such as NMDAR-interacting proteins and AKAPs, are highly expressed in neurons and enriched in dendritic spines (Lau and Zukin, 2007; Sanderson and Dell'Acqua, 2011). Toward this end, we transfected hippocampal neurons at 5–7 DIV with WT or mutant GluN2B(S1166A) together with GFP and performed recording from GFP<sup>+</sup> neurons at 11–15 DIV. Thirty minutes before recording, we applied the adenylyl cyclase inhibitor SQ22536 (100  $\mu\text{M}$ ) to deplete basal levels of cAMP and reduce endogenous PKA



**Figure 4.** Physiologically relevant stimuli that activate cAMP/PKA signaling promote Ser1166 phosphorylation. **a**, PKA-C promotes phosphorylation of GluN2B at Ser1166 after immunoprecipitation from brain lysates in the presence or absence of phosphatase (PP) inhibitors as indicated (+/–). Top, GluN2B immunocomplexes were incubated with no enzyme (control), PKA-C, or alkaline phosphatase (AP). Membranes were probed with antibodies to pSer1166-GluN2B in the absence or presence of phosphopeptide and total GluN2B to detect IP levels. Middle, Input samples (lysates). Bottom, Summary data showing pSer1166-GluN2B normalized to total GluN2B and then to control (no PKA-C);  $n = 3$ .  $*p = 0.0138$  for control versus PKA-C. **b**, **c**,  $\beta$ -Adrenergic and  $D_1/D_5$  dopamine receptor agonists promote phosphorylation of Ser1166 in forebrain slices. **b**, Top, Slices treated with vehicle or isoproterenol in the absence or presence of the  $\beta$ -adrenergic antagonist propranolol. Membranes were probed with antibodies to pSer1166-GluN2B, pSer897-GluN1, and pSer845-GluA1 to assess phosphorylation or to GluN2B, GluN1, or GluA1 to determine total subunit levels. Bottom, Quantification of phosphorylation levels of Ser1166, Ser897, and Ser845 normalized to total IP levels of GluN2B, GluN1, and GluA1, respectively, and then normalized to vehicle. **c**, Slices were treated with vehicle, the  $D_1/D_5$  dopamine receptor agonist SKF81297, or with SKF81297 in the presence of the  $D_1/D_5$  dopamine receptor antagonist SCH23390. GluN2B and GluA1 were immunoprecipitated and processed for immunoblotting as above. Images are from the same blot and exposure, but non-relevant lanes were removed. Bottom, Summary data as in **b**.  $*p < 0.05$ .  $***p < 0.001$ .  $****p < 0.0001$ .

activity. In neurons expressing WT or mutant NMDARs and treated with vehicle, NMDA-evoked responses exhibited rundown (Fig. 1c), consistent with previous findings (Skeberdis et al., 2006). Direct delivery of PKA-C via the patch pipette potentiated the peak phase of NMDA-elicited currents by ~1.3-fold (Fig. 1c; Table 1). In neurons expressing WT GluN1 with mutant GluN2B(S1166A), PKA potentiation was essentially abolished (Fig. 1c; Table 1), indicating the ability of overexpression of mutant GluN2B to outcompete endogenous GluN2B and GluN2A for assembly with GluN1 and to dominate NMDA-elicited currents.

The results thus far indicate a role for Ser1166 in potentiation of NMDA currents by PKA, but they do not address the mechanism by which this occurs at the single-channel level. Toward this end, we recorded activity in patches containing only a single channel from HEK293 cells expressing GluN1 together with WT or mutant GluN2B receptors. The nonphosphorylatable mutant, S1166A, exhibited an open probability that was markedly reduced ( $P_o$ ,  $0.13 \pm 0.02$ ,  $n = 14$ ) relative to that of the phosphomimetic mutation S1166D ( $0.26 \pm 0.03$ ,  $n = 16$ ,  $p = 0.0317$ ; Fig. 1d; Table 1). The WT receptor exhibited an open probability ( $0.22 \pm 0.04$ ,  $n = 20$ ) that was intermediate in value but did not differ statistically from that of S1166A or S1166D ( $p > 0.05$ , ANOVA) (Fig. 1d; Table 1), consistent with the substantial variability in GluN2B channel activity noted previously (Amico-Ruvio and Popescu, 2010; Borschel et al., 2012; Maki et al., 2012) and with the concept that WT GluN2B-containing NMDARs are a combination of both phosphorylated and nonphosphorylated receptor species. The large difference in activity between the

S1166A and the S1166D mutants is largely attributable to a change in mean closed durations, as neither mean open duration nor unitary current amplitude was altered (Fig. 1d; Table 1). As with  $P_o$  values, mean closed durations of WT NMDARs ( $28 \pm 5$  ms) were intermediate in value but did not differ significantly between that of S1166A ( $61 \pm 19$  ms) and S1166D ( $16 \pm 2$ ) mutant receptors ( $p > 0.05$ , ANOVA).

#### Ser1166 is critical to the PKA-dependent shift in NMDAR reversal potential

The results thus far indicate that Ser1166 of GluN2B is critical to PKA potentiation of NMDA-elicited currents, but they do not address a direct role for Ser1166 in regulation of NMDAR  $Ca^{2+}$  permeability by PKA. Toward this end, we undertook two experimental approaches. First, we examined current–voltage relations for HEK293 cells expressing GluN1 together with WT or mutant GluN2B(S1166A) receptors in the absence and presence of H-89. If, indeed, PKA facilitates  $Ca^{2+}$  permeation through NMDARs, inhibitors of PKA should cause a negative shift in the reversal potential ( $E_{rev}$ ) of the NMDAR current–voltage relationship ( $I/V$  curve) (Skeberdis et al., 2006). HEK293 cells are well suited for these experiments because of the relatively small contribution of voltage-gated conductance. NMDA-elicited currents were recorded at holding potentials ranging from  $-60$  to  $60$  mV, in 20 mV steps. In cells expressing WT NMDARs, the  $I/V$  curve was slightly outwardly rectifying and the NMDA reversal potential ( $E_{rev}$ ) was  $\sim 10$  mV (Fig. 2a,b), the known reversal potential for NMDARs. The noise in the responses reflects the stochastic gat-

**Table 2. Summary of statistics for Figures 4–6**

Figure	N	Mean ± SEM	p	Statistical test
4a	2–5 slices	Control: 0.2732 ± 0.0585 PKA: 1.564 ± 0.513 AP: 0.02005 ± 0.02396 PPI: 0.06882 ± 0.06091	*, 0.0138	One-way ANOVA with Tukey's post-test (control vs PKA)
4b	6–10 slices	S1166: vehicle, 0.2978 ± 0.04566; ISO, 2.019 ± 0.244; ISO + PROP, 0.3674 ± 0.03934 S897: vehicle, 0.2044 ± 0.03897; ISO, 1.034 ± 0.0576; ISO + PROP, 0.5836 ± 0.04153 S845: vehicle, 0.3929 ± 0.09082; ISO, 1.190 ± 0.0627; ISO + PROP, 0.6275 ± 0.09449	****, < 0.0001 ****, < 0.0001 ****, < 0.0001	One-way ANOVA with Tukey's post-test (vehicle vs ISO and ISO vs ISO + PROP)
4c	2–5 slices	S1166: vehicle, 0.4482 ± 0.04804; SKF, 0.8647 ± 0.05879; SKF + SCH, 0.4041 ± 0.1217 S897: vehicle, 0.1839 ± 0.05562; SKF, 0.6249 ± 0.2622; SKF + SCH, 0.2558 ± 0.0205 S845: vehicle, 0.4306 ± 0.08369; SKF, 1.063 ± 0.131; SKF + SCH, 0.4389 ± 0.1382	***, < 0.001 *, < 0.05 *, < 0.05	One-way ANOVA with Tukey's post-test (vehicle vs SKF81297)
5a	4 animals	S1166: control, 0.7940 ± 0.1259; Fox U, 0.7689 ± 0.0638 S897: control, 1.000 ± 0.117; Fox U, 1.067 ± 0.074 S845: control, 0.3718 ± 0.1036; Fox U: 0.6740 ± 0.1125	** , 0.0030	Two-tailed paired t test for GluA1 S845 control vs Fox U
5b	5 animals	S1166: control, 0.1226 ± 0.0198; FS, 1.073 ± 0.175 S897: control, 0.2884 ± 0.0358; FS, 1.208 ± 0.107 S845: control, 0.7569 ± 0.0874; FS, 1.415 ± 0.052	***, 0.0006 ****, < 0.0001 ***, 0.0002	Two-tailed unpaired t test for control vs FS
6b,c	5 animals	S1166: Sal control, 0.8118 ± 0.1251; Sal FS, 1.543 ± 0.238; Rp-cAMPS control, 0.4344 ± 0.1228, Rp-cAMPS FS, 0.8291 ± 0.1975 S897: Sal control, 1.147 ± 0.309; Sal FS, 1.508 ± 0.366; Rp-cAMPS control, 0.3963 ± 0.1025; Rp-cAMPS FS, 0.6003 ± 0.0837 S845: Sal control, 0.5753 ± 0.0382; Sal FS, 0.8542 ± 0.0903; Rp-cAMPS control, 0.2804 ± 0.0389 Rp-cAMPS FS, 0.4563 ± 0.0500	*, < 0.05 *, < 0.05 ***, < 0.001	One-way ANOVA with Tukey's post-test (S1166 and S845 Sal control vs FS; Sal FS vs Rp-cAMPS FS)

\**p* < 0.05; \*\**p* < 0.01; \*\*\**p* < 0.001; \*\*\*\**p* < 0.0001.

ing of single NMDA channels. Noise levels increased during NMDA application and were greatly reduced at potentials near the NMDA  $\Delta E_{rev}$ . Moreover, the direction of the NMDA-elicited current reversed at the known reversal potential for NMDARs, consistent with an NMDAR-mediated response. Application of H-89 triggered a marked reduction in slope conductance and shifted the  $E_{rev}$  of NMDA currents in the negative direction ( $\Delta E_{rev} = -6.79$  mV; Fig. 2*a–c*; Table 1), indicative of a decrease in the ratio of  $Ca^{2+}$  to monovalent cation permeability ( $P_{Ca^{2+}}/P_{monovalent}$ ) in corroboration of our earlier work (Skeberdis et al., 2006). In cells expressing GluN2B(S1166A), H-89 produced a much smaller reduction in the slope conductance and did not detectably alter the reversal potential ( $\Delta E_{rev} = 0.46$  mV; Fig. 2*a–c*; Table 1). We obtained similar results when we fit only the points closer to the  $E_{rev}$  at  $-20$ ,  $0$ , and  $20$  mV. These findings document that loss of a single residue abolishes the impact of PKA on NMDAR  $Ca^{2+}$  permeability.

#### Ser1166 of GluN2B is critical to PKA-dependent potentiation of NMDA-evoked $Ca^{2+}$ rises

Second, we examined the impact of Ser1166 on PKA-dependent NMDAR-mediated  $Ca^{2+}$  rises. Toward this end, we imaged NMDA-evoked  $Ca^{2+}$  rises in HEK293 cells expressing GluN1 together with WT or mutant GluN2B(S1166A) receptors. Cells were loaded with the  $Ca^{2+}$  indicator Fura-2 AM; and after a brief

(15–20 s) stable baseline,  $Ca^{2+}$  transients were elicited by a single application (60 s) of NMDA (100  $\mu$ M with 10  $\mu$ M glycine), and images were taken before (control), during (H-89), and after washout (recovery) of H-89. Application of NMDA and glycine elicited a rise in  $Ca^{2+}$  (Fig. 2*d,e*). In cells expressing WT NMDARs, H-89 markedly inhibited NMDA-evoked  $Ca^{2+}$  rises (Fig. 2*d–f*; Table 1). After washout of the inhibitor, NMDA-evoked  $Ca^{2+}$  rises exhibited substantial recovery. In cells expressing GluN2B(S1166A), inhibition by H-89 was greatly diminished (Fig. 2*d–f*; Table 1). Collectively, these findings strongly suggest that Ser1166 is important to regulation of NMDAR  $Ca^{2+}$  signaling by PKA.

#### Ser1166 is required for regulation of NMDAR-mediated $Ca^{2+}$ transients in spines

The results thus far indicate that Ser1166 is critical to regulation of NMDA-evoked  $Ca^{2+}$  signaling but do not distinguish between synaptic and extrasynaptic  $Ca^{2+}$  signaling.  $Ca^{2+}$  transients in the dendrites of CA1 pyramidal neurons induced by uncaging of glutamate can be restricted to a single spine and are mainly NMDAR-mediated (Yuste et al., 1999; Bloodgood and Sabatini, 2009). Moreover, spine  $Ca^{2+}$  is under the control of cAMP/PKA signaling (Skeberdis et al., 2006; Chalifoux and Carter, 2010; Higley and Sabatini, 2010). Whereas the functional impact of PKA on NMDARs is well established, the molecular target remains unknown. To investigate whether Ser1166 is critical to regulation



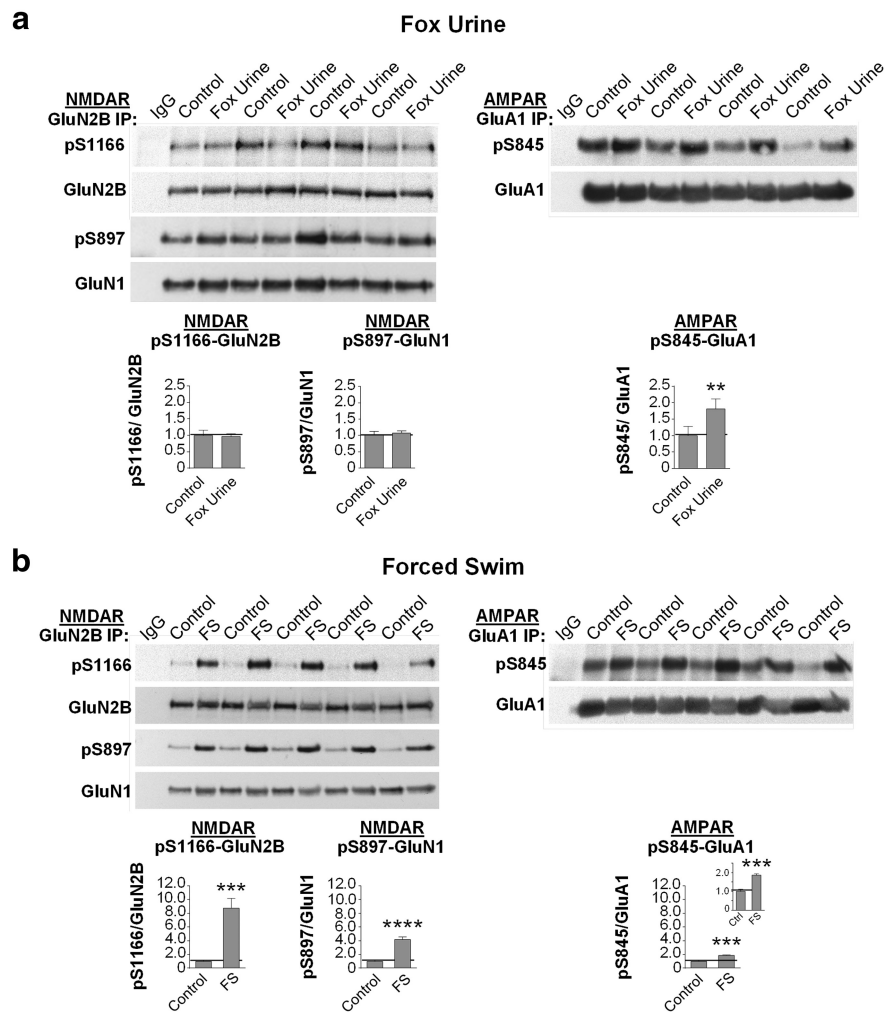
of  $Ca^{2+}$  rises in dendritic spines by PKA, we biolistically overexpressed WT or mutant GluN2B(S1166A) in organotypic hippocampal slices. At 10–14 d after transfection, we activated glutamatergic synapses by focal glutamate uncaging and simultaneously measured  $Ca^{2+}$  transients by two-photon microscopy and recorded uEPSCs at individual spines before and after application of isoproterenol, a well-known activator of  $\beta$ -adrenergic receptors and upstream activator of PKA (Fig. 3*a,b*).

In slices overexpressing WT GluN2B, application of isoproterenol (10  $\mu$ M, 5 min) induced a marked increase in the pharmacologically isolated, NMDA component of uEPSC amplitude and  $Ca^{2+}$  transients in individual dendritic spines (Fig. 3*c–f*; Table 1). In contrast, in slices overexpressing GluN2B(S1166A), isoproterenol did not detectably alter either the uEPSC amplitude or  $Ca^{2+}$  transients (Fig. 3*c–f*; Table 1). Thus, the loss of Ser1166 is sufficient to abolish the impact of  $\beta$ -adrenergic receptor (and hence PKA) activation on NMDAR-mediated  $Ca^{2+}$  transients and currents in individual spines. The finding that isoproterenol increases not only NMDAR-mediated  $Ca^{2+}$  transients, but also NMDA uEPSCs, is consistent with our earlier observation that, in the case of GluN2B-containing NMDARs, PKA modulates not only NMDAR  $Ca^{2+}$  signaling but also a substantial fraction of NMDA-evoked currents (Skeberdis et al., 2006). Collectively, these data indicate that Ser1166 is critical to  $\beta$ -adrenergic receptor-dependent potentiation of NMDAR-mediated  $Ca^{2+}$  signaling and uEPSCs in individual spines. Moreover, these findings, together with our findings in Figure 1*c*, demonstrate that GluN2B(S1166A) serves as a dominant-negative mutant in hippocampal neurons.

#### Activation of $\beta$ -adrenergic and $D_1/D_5$ dopamine receptors promote phosphorylation of Ser1166

Our findings thus far indicate that Ser1166 of GluN2B is a molecular and functional target of PKA, but they do not address whether physiological stimuli that engage and activate PKA signaling promote phosphorylation of Ser1166. Toward this end, we generated a phospho-specific antibody directed to Ser1166 of GluN2B. To validate the antibody, we performed several control experiments. First, we documented the ability of PKA-C to phosphorylate GluN2B at Ser1166 in GST constructs (data not illustrated), as well as immunocomplexes isolated from brain lysates (Fig. 4*a*; Table 2) and in whole-brain lysates (Fig. 4*a*). Second, we showed that preincubation with the phosphopeptide (p-peptide) blocked the phospho-Ser1166 signal, indicating specificity of the antibody (Fig. 4*a*). Third, we showed that omission of phosphatase inhibitors to permit endogenous phosphatase activity or addition of alkaline phosphatase abolishes the phospho-Ser1166 signal (Fig. 4*a*; Table 2).

The  $\beta$ -adrenergic (Hall, 2004; Dai et al., 2009) and  $D_1/D_5$  dopa-



**Figure 5.** Phosphorylation of GluN2B at Ser1166 is induced by forced swim, but not exposure to fox urine, *in vivo*. **a**, Representative Western blots (top) and summary data (bottom) for hippocampal lysates from rats exposed to fox urine for 5 min. Western blots were probed with antibodies to pSer1166-GluN2B, pSer897-GluN1, and pSer845-GluA1 and then with antibodies to total GluN2B, GluN1, and GluA1. Exposure to fox urine increased phosphorylation of GluA1 at Ser845, with little or no effect on phosphorylation levels of GluN2B at Ser1166 or GluN1 at Ser897.  $n = 4$  animals per treatment group,  $^{**}p = 0.0030$ . **b**, Representative Western blots (top) and summary data (bottom) for hippocampal lysates from rats exposed to a single episode of forced swim for 5 min. Western blots were probed with antibodies as above for exposure to fox urine. Forced swim induced striking phosphorylation of GluN2B at Ser1166, GluN1 at Ser897, and GluA1 at Ser845. Values for phosphorylated protein were normalized to the corresponding value for total protein and then to the corresponding value for control animals.  $n = 5$  animals per treatment group.  $^{***}p = 0.0006$  (S1166 control vs FS).  $^{***}p = 0.0002$  (S845 control vs FS).  $^{****}p < 0.0001$ .

minergic (Beaulieu and Gainetdinov, 2011) signaling cascades are well-known pathways that engage and activate cAMP/PKA signaling in neurons. Application of isoproterenol (1  $\mu$ M, 5 min) to forebrain slices induced robust phosphorylation of the NMDAR subunit GluN2B at Ser1166 (Fig. 4*b*; Table 2), which was reversed by the  $\beta$ -adrenergic receptor antagonist propranolol (Fig. 4*b*; Table 2). Isoproterenol also promoted marked phosphorylation of the NMDAR subunit GluN1 at Ser897 and AMPA receptor (AMPA) subunit GluA1 at Ser845, two well-established PKA targets (Roche et al., 1996; Tingley et al., 1997), in a propranolol-sensitive manner (Fig. 4*b*; Table 2). Similarly, application of the  $D_1/D_5$  dopamine receptor agonist SKF81297 (10  $\mu$ M, 15 min) induced prominent phosphorylation of GluN2B at Ser1166 (Fig. 4*c*; Table 2), which was reversed by the  $D_1/D_5$  receptor antagonist SCH23390 (Fig. 4*c*; Table 2). SKF81297 also induced prominent phosphorylation of GluN1 at Ser897 and GluA1 at Ser845 (Fig. 4*c*; Table 2). These results demonstrate that Ser1166 on GluN2B is phos-

phosphorylated in brain slices in response to physiologically relevant stimuli that activate PKA.

### Phosphorylation of Ser1166 is induced by the emotional response to stress *in vivo*

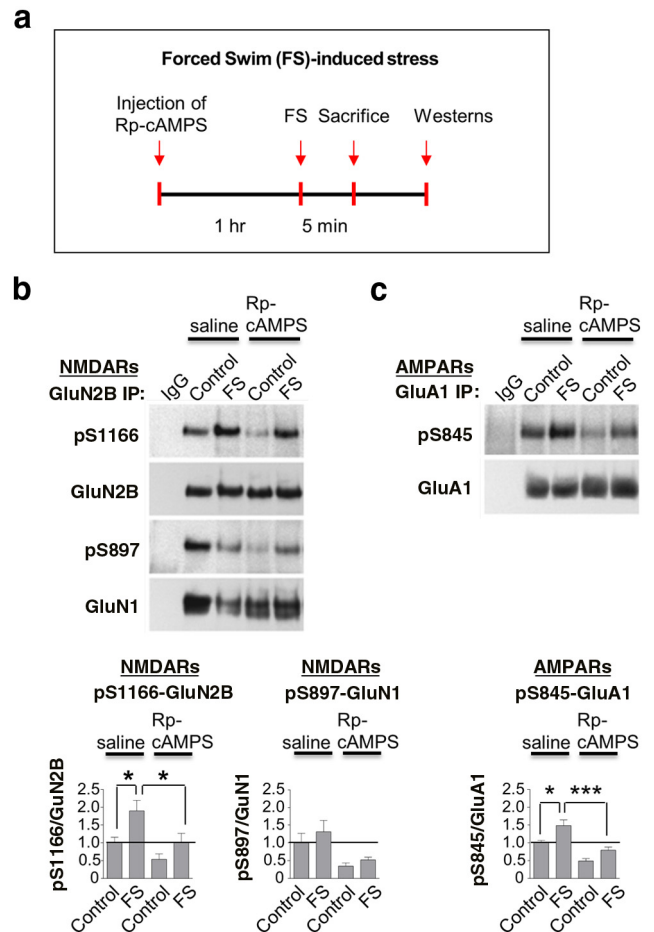
The results thus far indicate that phosphorylation of GluN2B at Ser1166 is critical to modulation of NMDAR function by PKA, but do not address experience-driven regulation of NMDAR phosphorylation at this site. Adverse experience in the form of stress promotes the release of norepinephrine and  $\beta$ -adrenergic signaling (McGaugh and Roozendaal, 2002; Padgett and Glaser, 2003). We examined whether adverse experience promotes phosphorylation of GluN2B at Ser1166 in the hippocampus. A single, brief exposure to fox urine stimulates release of norepinephrine and induces freezing behavior, indicative of fear (Hu et al., 2007; Liu et al., 2010). Exposure of animals to fox urine (5 min) increased phosphorylation of GluA1 at Ser845 in the hippocampal CA1, in corroboration of others (Hu et al., 2007) (Fig. 5*a*; Table 2), with little or no effect on phosphorylation of the NMDAR subunits GluN2B at Ser1166 and GluN1 at Ser897 (Fig. 5*a*; Table 2). These findings indicate that phosphorylation of NMDARs and AMPARs in response to emotional stress is under the control of different intracellular signaling pathways *in vivo*.

We next examined the impact of forced swim, another well-established paradigm of emotional as well as mild physical stress (Anisman et al., 2001), on NMDAR phosphorylation in the hippocampus. A single, brief exposure to forced swim stimulates release of norepinephrine (Yoshitake et al., 2004) and increases the magnitude of LTP at synapses onto dentate gyrus granule cells (Korz and Frey, 2003). Exposure of animals to forced swim (5 min) induced striking phosphorylation of GluN2B at Ser1166 and (to a lesser extent) phosphorylation of GluN1 at Ser897 (Fig. 5*b*; Table 2). Forced swim also induced a relatively modest phosphorylation of GluA1 at Ser845. The finding that adverse experience in the form of forced swim, but not exposure to fox urine, facilitates phosphorylation of GluN2B at Ser1166 suggests that different forms and/or intensities of stress can induce differential phosphorylation of PKA targets in a receptor- and site-specific manner.

To identify the signaling pathway that mediates stress-induced phosphorylation of GluN2B at Ser1166, we injected rats with the PKA inhibitor Rp-8-Br-cAMPS or saline, directly into the hippocampal CA1 at 1 h prior to forced swim and microdissected the small region of CA1 directly around the injection site (Fig. 6*a*). Forced swim induced an approximately twofold increase in phosphorylation of GluN2B at Ser1166 and a modest increase in phosphorylation of GluN1 at Ser897. Rp-8-Br-cAMPS markedly reduced phosphorylation of GluN2B at Ser1166 (and, to a lesser extent, GluN1 at Ser897) under both control and forced swim conditions (Fig. 6*b*; Table 2). These findings are consistent with the concept that both basal and experience-dependent phosphorylation of Ser1166, at least in part, is mediated by cAMP/PKA signaling. Rp-8-Br-cAMPS also reduced basal and forced swim-induced phosphorylation of GluA1 at Ser845 (Fig. 6*c*; Table 2).

### Discussion

The present study identifies Ser1166 in the NMDAR subunit GluN2B as a novel molecular and functional target of PKA phosphorylation critical to potentiation of NMDAR  $Ca^{2+}$  signaling at synapses and regulated by the emotional response to stress. We show that physiologically relevant stimuli that engage and activate cAMP/PKA signaling, and are important to the acquisition



**Figure 6.** Forced swim (FS)-induced phosphorylation of Ser1166 is PKA-dependent. *a*, Schematic of experimental paradigm for injections and FS. *b*, *c*, Top, Representative Western blots. Bottom, Summary data for lysates of hippocampal CA1 from rats injected with Rp-8-Br-cAMPS or saline 1 h before a single episode of FS for 5 min. Western blots were probed with antibodies to pSer1166-GluN2B, pSer897-GluN1, pSer845-GluA1, and total GluN2B, GluN1, and GluA1, respectively. Rp-8-Br-cAMPS markedly reduced FS-induced phosphorylation of GluN2B at Ser1166 and GluA1 at Ser845, with little or no effect on pSer897-GluN1. Values for phosphorylated protein were normalized to the corresponding value for total protein and then to the corresponding value for control animals.  $n = 5$  animals per treatment group. \* $p < 0.05$ . \*\*\* $p < 0.001$ .

and consolidation of memory (McGaugh and Roozendaal, 2002; Hu et al., 2007), regulate phosphorylation of NMDARs and  $Ca^{2+}$  signaling in spines. Accordingly, activation of  $\beta$ -adrenergic and  $D_1/D_5$  dopamine receptors stimulates phosphorylation of GluN2B at Ser1166 in brain slices. Moreover, activation of  $\beta$ -adrenergic receptors potentiates  $Ca^{2+}$  transients and uncaging-evoked synaptic currents in individual spines of neurons expressing WT, but not mutant S1166A, NMDARs. Thus, loss of this single phosphorylation site is sufficient to abolish  $\beta$ -adrenergic-dependent potentiation of NMDA currents and  $Ca^{2+}$  transients, linking phosphorylation of NMDARs to  $Ca^{2+}$  signaling in spines. This is significant in that central noradrenergic signaling mediates emotional arousal and the emotional response to stress (McGaugh and Roozendaal, 2002; Padgett and Glaser, 2003). We further show that adverse experience in the form of forced swim, but not exposure to fox urine, elicits striking phosphorylation of Ser1166 in a PKA-dependent manner *in vivo*, revealing experience-dependent phosphorylation of this site in a physiologically relevant context. These findings indicate that different stress

paradigms differentially regulate NMDAR phosphorylation and, by inference,  $\text{Ca}^{2+}$  permeation and  $\text{Ca}^{2+}$  signaling.

cAMP/PKA signaling represents a pivotal mechanism by which NMDAR-mediated  $\text{Ca}^{2+}$  signaling is modulated in neurons under physiological and pathological conditions. PKA promotes NMDAR  $\text{Ca}^{2+}$  permeability,  $\text{Ca}^{2+}$  rises in spines, and induction of LTP at CA1 synapses (Skeberdis et al., 2006). Moreover, extracellular signals that modulate cAMP bidirectionally regulate  $\text{Ca}^{2+}$  permeation through NMDARs and spine  $\text{Ca}^{2+}$  in a PKA-dependent manner (Chalifoux and Carter, 2010; Higley and Sabatini, 2010). Although these studies establish a functional role for PKA, they do not address the molecular target. In the present study, we reveal a previously unrecognized mechanism by which activation of major signaling pathways, known to couple positively to cAMP/PKA and play a critical role in learning and memory, can alter  $\text{Ca}^{2+}$  signaling in activated dendritic spines. Although not addressed by the present manuscript, phosphorylation of the carboxy-terminal tails of channels is known to affect channel properties, including ion selectivity. Moreover, phosphorylation events that regulate channel activity typically occur in either the N-terminal or C-terminal domains of the channel and not in the pore itself (Levitan and Kaczmarek, 2002). By analogy, phosphorylation of Ser1166 may induce a delocalized conformational change. Alternatively, S1166 might form part of a recognition motif for an as yet unidentified protein, which in some capacity regulates the selectivity pore within the NMDA channel.

A novel finding of the present study is that emotional stress in the form of a single exposure to forced swim, but not exposure to fox urine, elicits striking phosphorylation of Ser1166 in a PKA-dependent manner *in vivo*, revealing experience-driven regulation of phosphorylation at this site. Notably, these effects persist as late as P32, the oldest age examined, consistent with the concept that regulation of NMDAR  $\text{Ca}^{2+}$  permeation and signaling in spines (and GluN2B expression) are fundamental mechanisms relevant throughout the life span of the animal. Although exposure to fox urine promotes PKA-dependent phosphorylation of GluA1 at Ser845 (Hu et al., 2007) and induces transcription of the AMPAR subunit GluA2 in cerebellar stellate cells (Liu et al., 2010), it does not alter the phosphorylation status of NMDARs. At hippocampal CA1 synapses, phosphorylation of GluA1 at Ser845 facilitates synaptic delivery of GluA1-containing AMPARs and lowers the threshold for LTP (Oh et al., 2006; Hu et al., 2007); and at cerebellar synapses, transcription of GluA2 drives a switch in AMPAR phenotype and  $\text{Ca}^{2+}$  signaling (Liu et al., 2010). Whereas phosphorylation of GluN2B at Ser1166 potentiates NMDAR function and  $\text{Ca}^{2+}$  signaling in dendritic spines, with little or no effect on NMDAR trafficking (Skeberdis et al., 2006), phosphorylation of GluN1 at Ser897 promotes forward trafficking of NMDARs from the endoplasmic reticulum to the cell surface (Scott et al., 2003). These findings reinforce the notion that different forms of stress can promote PKA-dependent phosphorylation of different glutamate receptors and/or receptor-associated proteins at different sites. Although both exposure to fox urine and a single episode of forced swim elevate noradrenergic signaling, they differ in their characteristics and severity. Whereas exposure to fox urine, an olfactory stimulus, is a psychological form of stress, forced swim stress has both psychological and physical components (Anisman et al., 2001). Forced swim also promotes surface expression of GluN2B and enhances NMDAR EPSCs at synapses of the prefrontal cortex (Yuen et al., 2009).

In conclusion, our data provide evidence for a mechanism by which physiologically relevant stimuli, important to emotional arousal and learning and memory, regulate NMDAR  $\text{Ca}^{2+}$  perme-

ation and  $\text{Ca}^{2+}$  signaling in spines. Whereas other protein kinases that phosphorylate NMDARs, such as Src (Yu et al., 1997), Fyn (Prybylowski et al., 2005), PKC (Lu et al., 1999; Lan et al., 2001), and casein kinase 2 (Sanz-Clemente et al., 2010), regulate NMDAR trafficking and gating in response to neuronal activity, PKA is unique in that it regulates NMDAR-mediated  $\text{Ca}^{2+}$  permeation. We identify Ser1166 on GluN2B as a novel molecular and functional target of PKA essential to  $\text{Ca}^{2+}$  signaling in spines and regulated by the emotional response to stress. Importantly, different forms of adverse experience differentially regulate phosphorylation of this site. These findings add a new dimension to our understanding of how experience influences synaptic proteins and function. Given that NMDARs and PKA are expressed widely throughout the brain, phosphorylation of Ser1166 represents a potentially powerful mechanism to adjust synaptic function. NMDARs are critical to many forms of synaptic plasticity and to higher cognitive functions, such as learning and memory, and their dysregulation is implicated in neuropsychiatric disorders, such as schizophrenia, stroke, Parkinson's disease, and Huntington's disease. Thus, our findings have important ramifications for NMDARs not only under normal, but also pathological, conditions.

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