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Disruption of NMDA receptor function prevents normal experience-dependent homeostatic synaptic plasticity in mouse primary visual cortex

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Abbreviated title: NMDARs in homeostatic synaptic plasticity of V1

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43 **Abstract**

44 Homeostatic regulation of synaptic strength allows for maintenance of neural activity
45 within a dynamic range for proper circuit function. There are largely two distinct modes of
46 synaptic plasticity that allow for homeostatic adaptation of cortical circuits: synaptic scaling and
47 sliding threshold (BCM theory). Previous findings suggest that the induction of synaptic scaling
48 is not prevented by blocking NMDA receptors (NMDARs) while sliding threshold model posits
49 that the synaptic modification threshold of LTP and LTD readjusts with activity hence, the
50 outcome of synaptic plasticity is NMDAR-dependent. While synaptic scaling and sliding
51 threshold have been considered two distinct mechanisms, there are indications from recent
52 studies that these two modes of homeostatic plasticity may interact or that they may operate
53 under two distinct activity regimes. Here we report using both sexes of mouse that acute genetic
54 knockout of the obligatory subunit of NMDAR or acute pharmacological block of NMDAR
55 prevents experience-dependent homeostatic regulation of AMPAR-mediated miniature
56 excitatory postsynaptic currents (mEPSCs) in layer 2/3 of visual cortex. This was not due to
57 gross changes in postsynaptic neuronal activity with inhibiting NMDAR function as determine
58 by c-Fos expression and 2 photon Ca^{2+} imaging in awake mice. Our results suggest that
59 experience-dependent homeostatic regulation of intact cortical circuits is mediated by NMDAR-
60 dependent plasticity mechanisms, which supports a sliding threshold model of homeostatic
61 adaptation.

62

63 **Significance statement**

64 Prolonged changes in sensory experience lead to homeostatic adaptation of excitatory synaptic
65 strength in sensory cortices. Both sliding threshold and synaptic scaling models can account for
66 the observed homeostatic synaptic plasticity. Here we report that visual experience-dependent
67 homeostatic plasticity of excitatory synapses observed in superficial layers of visual cortex is
68 dependent on NMDA receptor function. In particular, both strengthening of synapses induced by
69 visual deprivation and the subsequent weakening by reinstatement of visual experience were
70 prevented in the absence of functional NMDA receptors. Our results suggest that sensory
71 experience-dependent homeostatic adaptation depends on NMDA receptors, which supports the
72 sliding threshold model of plasticity and input-specific homeostatic control observed *in vivo*.

73

74 **Introduction**

75 Neuronal circuits constantly undergo changes through development, experience, and
 76 learning that allow for adaptation to different environments or internal states. Correlation based
 77 synaptic plasticity mechanisms such as long-term potentiation (LTP) and long-term depression
 78 (LTD) are largely responsible for storing such information. However, LTP and LTD have innate
 79 positive feedback loop that requires additional homeostatic mechanisms to allow stability of
 80 neural circuits undergoing plasticity. Several models of homeostatic plasticity can achieve this
 81 function, including synaptic scaling (Turrigiano, 2008) and the sliding threshold model (Cooper
 82 and Bear, 2012).

83 According to synaptic scaling, prolonged reduction in neuronal activity leads to an
 84 upscaling of the strength of excitatory synapses, while a period of enhanced activity results in a
 85 downscaling (Turrigiano et al., 1998). Synaptic scaling was initially proposed to occur globally
 86 across the majority of synapses in a multiplicative manner to preserve relative differences in
 87 synaptic weight (Turrigiano et al., 1998). Initial demonstrations of synaptic scaling were done by
 88 observing changes in the amplitude of miniature excitatory postsynaptic currents (mEPSCs) in
 89 cultured neurons upon pharmacological manipulation of neural activity, and its induction is
 90 largely independent of NMDA receptor (NMDAR) activation (O'Brien et al., 1998; Turrigiano et
 91 al., 1998). Similar homeostatic changes in mEPSCs, which has been interpreted as synaptic
 92 scaling, can be induced in pyramidal neurons of rodent primary visual cortex (V1) by dark
 93 exposure (DE) (Goel et al., 2006; Goel and Lee, 2007; He et al., 2012), intraocular TTX injection
 94 (Desai et al., 2002), enucleation (He et al., 2012), and retinal lesions (Keck et al., 2013). Similar
 95 to what is observed in cultured neurons, synaptic scaling in V1 is largely mediated by the
 96 insertion or removal of AMPA receptors (AMPA) (Goel et al., 2006; Goel et al., 2011).

97 Sliding threshold model states that prolonged periods of altered activity result in the
 98 modification of the threshold for LTP and LTD induction. The synaptic modification threshold
 99 shifts bidirectionally depending on the history of neuronal activity: an extended period of low
 100 activity slides the threshold to favor LTP, while high activity shifts it to favor LTD (Abraham
 101 and Bear, 1996; Cooper and Bear, 2012). Sliding threshold has been demonstrated in rodent V1,
 102 in which dark-rearing (DR) or DE leads to a lower threshold for LTP induction (Kirkwood et al.,
 103 1996; Philpot et al., 2003; Guo et al., 2012). Synaptic modification threshold “slides” by changes
 104 in either the induction mechanisms of LTP/LTD, such as alterations in NMDAR function
 105 (Quinlan et al., 1999; Philpot et al., 2003) and inhibition (Steele and Mauk, 1999), or the
 106 expression mechanisms of LTP/LTD, such as changes in AMPAR phosphorylation (Huang et al.,
 107 2012).

108 Although seemingly different, both synaptic scaling and sliding threshold allow
 109 homeostatic adaptation of synapses to maintain neuronal activity within a physiologically
 110 relevant, yet stable dynamic range. Previous investigations have shown that both changes in
 111 mEPSC amplitude and changes in the modification threshold happen *in vivo* in V1. They have,
 112 however, failed to address how these two processes interact, if at all, and to which extent. Recent
 113 evidence suggests that different levels of activity change may trigger synaptic scaling versus
 114 sliding threshold mode of homeostatic plasticity in V1 (Bridi et al., 2018). Based on the reported
 115 differences in the requirement of NMDAR activation for synaptic scaling and sliding threshold
 116 models, we examined the role of NMDARs on visual experience-dependent changes in mEPSCs
 117 of L2/3 neurons of mouse V1, which have been interpreted as synaptic scaling (Desai et al.,
 118 2002; Goel et al., 2006; Goel and Lee, 2007; He et al., 2012; Keck et al., 2013). We reasoned
 119 that if the observed scaling of mEPSCs with visual experience is a consequence of LTP/LTD due

120 to the sliding threshold, then these changes would depend on NMDAR activation. Using cell-
 121 type specific knockout mice or an antagonist of NMDARs, we found evidence supporting a key
 122 role of NMDARs in mediating experience-dependent homeostatic synaptic plasticity in V1.

123

124 **Materials and Methods**

125 *Visual experience manipulation*

126 All animal handling and manipulations were approved by the Institutional Animal Care
 127 and Use Committee (IACUC) at Johns Hopkins University and followed the guidelines
 128 established by the National Institutes of Health (NIH). Male and female NR1^{flox} mice
 129 (<https://www.jax.org/strain/005246>; RRID: IMSR_JAX:005246) were raised under a 12 hours
 130 light/dark cycle until postnatal day 25-35 (P25-P35). At this point a group of mice was placed in
 131 24-hour dark conditions for 2 days (2 days dark exposure, DE). Animals in the dark were cared
 132 for by using infrared vision goggles. A group of DE mice were taken out of the dark and re-
 133 exposed to light for 2 hours (2 hours light-exposed, LE). Age matched control animals were
 134 continuously raised in the normal 12 hours light/dark cycle (Ctl).

135

136 *Targeted viral transfection*

137 Male and female NR1^{flox} mice between P23-P27 were bilaterally injected with an adeno-
 138 associated viral vector expressing Cre-GFP under the control of CaMKII promoter
 139 (AAV9.CaMKII.HI.eGFP-Cre.WPRE.SV40; Penn Vector Core, University of Pennsylvania,
 140 Cat#: AV-9-PV2521) in V1. Layer 2/3 of V1 was targeted by using the following stereotaxic
 141 coordinates relative to Bregma: posterior -3.6mm, lateral 1.5mm, and depth -0.3mm. Mice
 142 recovered on a heated pad until movement, eating and drinking behaviors were evident. Animals

143 were returned to the mouse colony after recovery and remained under 12 hours light/dark
 144 conditions until experimental use. Viral expression and knockout of NR1 gene was confirmed
 145 experimentally 6-7 days after transfection as determined by significantly reduced NMDAR
 146 currents (Fig. 1). Manipulation of visual experience therefore commenced 1 week (6-7d) after
 147 viral injections. Control mice underwent the same procedure, but instead were injected with a
 148 GFP-expressing adeno-associated virus (AAV9.CaMKII0.4.eGFP.WPRE.rBG; Penn Vector
 149 Core, University of Pennsylvania, Cat#: AV-9_PV1917).

150

151 *In vivo CPP application*

152 For control normal-reared and DE groups, NMDAR antagonist D-4-[(2*E*)-3-Phosphono-
 153 2-propenyl]-2-piperazinecarboxylic acid (d-CPP; Tocris, Minneapolis, MN; Cat#: 1265) was
 154 delivered intraventricularly for 2 days via Alzet osmotic minipumps (DURECT Corporation,
 155 Cupertino, CA; Cat#: 1007D) coupled to a brain infusion cannula (Alzet, DURECT
 156 Corporation, Cupertino, CA; Cat# 8851). Control groups were infused with saline instead of d-
 157 CPP. In brief, all osmotic minipumps (0.5 μ L/hr) were backfilled with either d-CPP (10 μ M) or
 158 saline solution and primed by incubating in saline solution at 37°C for at least 5 hours before
 159 implantation. Implantation surgery was done under constant administration of 1.5-2%
 160 isoflurane/oxygen mix under aseptic conditions. Anesthesia level and vital signs were monitored
 161 during the surgery. Craniotomy was performed for cannula insertion by drilling a small hole in
 162 the skull at stereotaxic coordinates -0.22 mm posterior, 1 mm lateral from Bregma to target the
 163 lateral cerebral ventricle using a dental drill and a sterilized 0.5 mm drill-bit. The neck was
 164 aseptically cleaned, a small cut was made at the base and blunt forceps were used to separate the
 165 fascia, then an osmotic mini-pump was inserted subcutaneously. The cannula was guided to the

166 drilled hole on the skull and secured in place with dental cement (TEETS denture material;
 167 Patterson dental, MN; Cat#: 223-3773). Following the surgery, mice were recovered on a 30°C
 168 heat pad, and returned to the home cage where drinking water was supplemented with 0.07
 169 mg/mL carprofen (Sigma-Aldrich 33975; CAS#: 53716-49-7) (Ingrao et al., 2013). For DE
 170 group, mice were allowed to recover for at least 12 hours before placed inside a darkroom for 2
 171 days. For LE groups, d-CPP (10 mg/kg) or saline were delivered intraperitoneally (200 µl
 172 volume) in the dark room 10 min before light exposure.

173

174 ***Primary visual cortex slice preparation***

175 Mice between P25-P35 were deeply anesthetized with isoflurane gas in a chamber placed
 176 in a chemical fume hood. Anesthesia was delivered to dark exposed animals in a light-tight
 177 chamber. After confirming the absence of pinch or righting reflex, mice were decapitated and the
 178 brain was immediately placed in ice-cold dissection buffer containing the following (in mM):
 179 212.7 sucrose, 10 dextrose, 3 MgCl₂, 1 CaCl₂, 2.6 KCl, 1.23 NaH₂PO₄•H₂O, and 26 NaHCO₃,
 180 which was bubbled with 95% O₂/5% CO₂ gas. Blocks containing V1 were rapidly isolated and
 181 sectioned coronally into 300 µm thick slices, while submerged in ice-cold dissection buffer,
 182 using a vibratory tissue slicer (PELCO easiSlicer, Ted Pella; product#: 11000). Slices were
 183 transferred to a submersion holding chamber filled with artificial cerebrospinal fluid containing
 184 (in mM): 124 NaCl, 5 KCl, 1.25 NaH₂PO₄•H₂O, 26 NaHCO₃, 10 dextrose, 2.5 CaCl₂, and 1.5
 185 MgCl₂, bubbled with 95% O₂/5% CO₂. The slices recovered for 1 hour at room temperature
 186 before electrophysiological recordings started.

187

188 ***Electrophysiological recordings***

189 Slices were transferred to a submersion-type recording chamber and perfused with oxygenated
 190 ACSF (bubbled 95% O₂/5% CO₂ at 32 ± 2°C) at a rate of 2mL/min. The chamber was mounted
 191 on a fixed stage under an upright microscope (E600 FN; Nikon, Tokyo, Japan) with oblique
 192 infrared illumination. Pyramidal neurons in L2/3 of V1 were visually identified and patched
 193 using a glass pipette with a tip resistance between 3 and 5 MΩ, which was filled with internal
 194 solution containing (in mM): 120 CsOH, 120 Gluconic acid, 10 phosphocreatine, 0.5 GTP, 4
 195 ATP, 8 KCl, 1 EGTA, 10 HEPES and 5 QX-314. An Axon patch-clamp amplifier (Multiclamp
 196 700B, Molecular Devices) was used for voltage-clamp recordings and data was acquired through
 197 Igor Pro software (WaveMetrics, <http://www.wavemetrics.com/products/igorpro/igorpro.htm>;
 198 RRID:SCR_000325). Only data from cells with input resistance (R_i) > 150 MΩ and series
 199 resistance (R_s) < 25 MΩ were analyzed.

200 NMDAR/AMPA ratio Glutamatergic currents were recorded in response to electric stimulation
 201 delivered through a bipolar glass electrode placed in V1 L4 or L2/3. Recordings were done in the
 202 presence of 20 μM bicuculline methiodide (Enzo Life Sciences, product #: BML-EA149-0050)
 203 in the ACSF. The stimulation intensity was adjusted so that a single-peak response was produced
 204 with an onset latency of 2-3ms. The AMPA receptor component was taken as the average peak
 205 amplitude of responses recorded at V_h=-70 mV. The NMDA receptor component was taken as
 206 the average amplitude of responses recorded at V_h=+40 mV 70 ms after onset. Responses were
 207 recorded every 10 s and a minimum of 10 responses were averaged for each component.

208 miniature EPSCs AMPA receptor-mediated miniature excitatory postsynaptic currents
 209 (mEPSCs) were isolated by recording with 1 μM tetrodotoxin citrate (TTX; Abcam; product#:
 210 ab120055), 20 μM bicuculline methiodide, and 100 μM DL-2-amino-5 phosphonopentanoic acid
 211 (DL-APV; Sigma-Aldrich A5282; CAS#: 76326-31-3) in the ACSF. Events were recorded at

212 Vh=-80 mV for a minimum of 4 minutes initiated 1-2 minutes after cell break-in. The recorded
 213 data were digitized at 2 kHz by a data acquisition board (National Instruments), acquired with
 214 Igor Pro software and analyzed using the MiniAnalysis program (Synaptosoft,
 215 <http://www.synaptosoft.com/MiniAnalysis/>; RRID:SCR_002184). The detection threshold for
 216 mEPSCs was set to 3 times the root mean square (RMS) noise and events with a rise time > 3 ms
 217 were excluded from analysis. Events within bursts (more than 2 events, inter-event-interval < 10
 218 ms) were excluded from the measurement of amplitudes. The average of total isolated events
 219 (200-220) was used to calculate the decay time constant for each neuron. Cells were discarded if
 220 Ri or Rs changed more than 15% during the duration of the recording.

221

222 ***Biocytin processing***

223 Slices used for electrophysiological recordings were immediately fixed in 10% formalin
 224 (Sigma-Aldrich, HT5014; MDL: MFCD00003274) solution overnight at 4°C. Slices were rinsed
 225 0.01M phosphate buffered saline (PBS) at room temperature and permeabilized in 2% Triton X-
 226 100 (Fisher) in PBS for 1 h. Slices were then incubated in 1:2000 solution of avidin-Texas Red
 227 conjugate (Life Technologies; A820) in 1% Triton X (in PBS) overnight. After incubation, slices
 228 were washed in PBS, mounted on glass slides, and coverslipped with Prolong Gold Anti-fade
 229 (Fisher; product #: P36930) mounting medium. Images were taken using a Zeiss LSM 510
 230 META confocal microscope.

231

232 ***Immunohistochemistry***

233 NR1^{flox} mice were deeply anesthetized with isoflurane vapors in a closed chamber placed
 234 in a fume hood. 2dDE animals were anesthetized in a light-tight chamber. Animals were

235 perfused transcardially with PBS followed by 10% formalin solution. The brains were then
 236 extracted and kept in 10% formalin overnight. V1 was isolated and sectioned coronally in 40 μ m
 237 thick slices. Free floating slices containing V1 were incubated with 1% sodium borohydride
 238 (Fisher; Cat#: S-678-10) for 15 minutes at room temperature and then washed with PBS. The
 239 same slices were blocked for 2 hours in a solution containing 3% goat serum (Sigma-Aldrich;
 240 product #: 9023) and 0.3% Triton-X in PBS or mixture of 10% normal goat serum (Jackson
 241 Immunoresearch; product code#: 005-000-121), 5% normal donkey serum (Jackson
 242 Immunoresearch; product code#: 017-000-121) and 0.5% Triton-X in PBS. Cortical slices were
 243 then incubated with antibodies against c-Fos and Neuronal nuclei protein (NeuN) in the blocking
 244 buffer overnight. Slices were rinsed and then incubated for 2 hours with fluorescently labeled
 245 secondary antibodies. Slices were then washed with PBS, incubated with DAPI and mounted on
 246 glass slides with Prolong Gold Anti-fade medium. The antibody concentrations were as follows:
 247 1:20,000 rabbit anti-cFos (Calbiochem, Cat#PC38; RRID:AB_2106755) or 1:500 rabbit anti-c-
 248 Fos (Cell Signaling Technology, Cat#2250S; RRID:AB_10692514), 1:200 mouse anti-NeuN
 249 (Millipore MAB377; RRID: AB_2298772), 1:200 donkey anti-mouse Alexa Fluor 488 (Thermo
 250 Fisher; RRID: AB_141607), 1:200 goat anti-rabbit Alexa Fluor 633 (Thermo Fisher; RRID:
 251 AB_2535731), goat anti-mouse Alexa Fluor 555 (Thermo Fisher; RRID: AB_2535844). Slices
 252 were imaged using either Zeiss LSM 700 or 800 confocal microscope with a step size of 0.5 μ m.
 253 All images were analyzed using Volocity software.

254 *Two Photon Ca^{2+} imaging in awake head fixed mice*

255 Male and female Emx1-Cre x Ai96 mice (<https://www.jax.org/strain/005628> RRID:
 256 MGI:2684615, <https://www.jax.org/strain/024106>) underwent head plate and cranial window

258 implantation surgery at p23-p25. The head plate and cranial windows, consisting of one 5 mm
 259 and two 3 mm round coverslips, were based on an existing design (Goldey et al., 2014). After 3-
 260 4 days of recovery, the mice were habituated to head-fixation in a body tube for 4-5 days.
 261 Habituated mice readily accepted sucrose and did not react aversively to visual stimuli. 2-photon
 262 calcium imaging was performed at P33-P35, using a custom-built microscope based on a Janelia
 263 Farm design (<https://wiki.janelia.org/wiki/display/shareddesigns/MIMMS>). GCaMP6s
 264 expressing neurons were imaged through a 16x 0.8 NA Nikon using a Chameleon Ultra 2 laser
 265 (Coherent) at 940 nm. Due to relatively weak fluorescence of the Ai96 line, imaging depth was
 266 limited to 150-220 μm with laser power set to 75-100 mW. Images were acquired at 30 Hz using
 267 Scanimage 2018 (Pologruto et al., 2003) and analyzed using custom scripts written in Matlab
 268 (Mathworks).

269 Visual stimuli and data acquisition During the imaging session, mouse was head-fixed in the
 270 body tube with visual stimuli displayed on a monitor centered 25 cm from the contralateral eye
 271 (60 Hz, mean luminance 30 cd/m^2). The stimuli were moving sinusoidal gratings at 8
 272 orientations (45° increments), with spatial frequency 0.05 cycles/degree, temporal frequency 3
 273 cycles/second. The gratings were shown for 3 s followed by 6 s interstimulus interval and were
 274 organized into blocks of 10 stimuli, with each block containing all the gratings and two
 275 additional blank stimuli in random order. After 12 repetitions of each stimulus (lasting 18
 276 minutes), the mouse was taken out of the body tube and given an i.p. injection of either saline or
 277 10 mg/kg CPP (Tocris, Cat#: 01773). A second round of imaging was then done 30-60 minutes
 278 after the injection in the same region. Depth was manually adjusted to match pattern of cell
 279 bodies from first round of imaging.

280

281 Data processing and analysis After cross-correlation based frame alignment, ROIs were
 282 manually drawn around visible cell bodies and cell body masks were calculated in semi-
 283 automated manner using correlation with a seed pixel. Neuropil signal was estimated from pixels
 284 within ROIs that were at least 2 pixels away from the mask boundary. Fluorescent traces for both
 285 mask signal and neuropil were filtered by 0.5 second running average to reduce noise.
 286 Fluorescent baseline F_0 was calculated as the running 10th percentile over 1800 frames (~1
 287 minute) of the mask signal. Fluorescence signal from cell bodies was then calculated as $\Delta F/F_0$
 288 $=((F_{\text{mask}} - 0.7 * F_{\text{neuropil}}) - F_0) / F_0$. Visual responses were calculated as the mean $\Delta F/F_0$ over the 3 s
 289 duration of the stimulus minus mean $\Delta F/F_0$ over 1 s preceding the visual stimulus. Spontaneous
 290 activity was calculated as the mean $\Delta F/F_0$ over 7 s window following the onset of blank stimuli.
 291 To determine whether the neuron was visually responsive, the difference between visual
 292 response and blank stimulus response was compared to distribution of such differences for
 293 scrambled data (1000 re-samplings). The neuron was considered visually responsive if the
 294 difference was higher than at least 950 (95%) of differences from the scrambled distribution.
 295 Note that this excludes neurons that are suppressed by visual stimuli or show off-response to
 296 visual stimuli. A similar procedure was used to determine whether the neuron had significant
 297 orientation selectivity. The orientation selectivity index $OSI = (R_{\text{pref}} - R_{\text{ortho}}) / (R_{\text{pref}} + R_{\text{ortho}})$ was
 298 calculated and then compared to distribution of OSI calculated from scrambled data. If the OSI
 299 was higher than 95% of OSIs from the scrambled distribution, the neuron was considered
 300 significantly orientation selective.

302 *Experimental Design and Statistical Analysis*

303 Data are presented as mean \pm SEM. All statistical analyses were done using Prism 7.0
 304 (GraphPad Software; RRID: SCR_002798). One-factor analysis of variance (ANOVA) was used
 305 to compare multiple groups followed by a Tukey's multiple comparison *post hoc* test. Unpaired
 306 Student's t-tests were used for two group comparisons. The Kolmogorov-Smirnov (K-S) test was
 307 used to compare cumulative probabilities. A p-value < 0.05 was used as a measure of
 308 significance in t-tests, ANOVAs, and Tukey's or Bonferroni *post hoc* analyses. For K-S tests, p-
 309 values < 0.01 were used as a measure of significance. Asterisk (*) is used in both text and figures
 310 to denote statistical significance.

311

312 Results

313 *Neuron-specific NMDAR KO*

314 To test whether NMDAR is necessary for experience-dependent homeostatic synaptic
 315 plasticity in V1, we aimed to specifically knockout (KO) NMDARs in principal neurons. This
 316 was achieved by targeted injection of an adeno-associated viral construct expressing Cre-GFP
 317 under the control of the CaMKII promoter (AAV9.CaMKII.Cre-eGFP; Cre-GFP condition) into
 318 V1 L2/3 of NR1^{flox} transgenic mice (Tsien et al., 1996). In this scheme, the expression of Cre
 319 recombinase leads to excision of the *Grin1* gene, which encodes the obligatory NMDAR NR1
 320 (GluN1) subunit. To control for effects only due to viral transfection, a second group of NR1^{flox}
 321 transgenics were injected with a GFP-expressing viral construct (AAV9.CaMKII.GFP; GFP-only
 322 condition). We verified that viral transfection efficiency was similar for both constructs by
 323 quantifying the percentage of GFP-positive cells relative to the total number of neurons in a
 324 given tissue section of V1 L2/3 (Fig. 1A). To determine whether Cre-GFP condition leads to
 325 effective knockout of NMDARs, we used whole-cell voltage clamp to measure

326 NMDAR/AMPA ratios after viral injections. We corroborated the specificity of the functional
 327 NMDAR knockout by also measuring NMDAR/AMPA ratios from non-GFP expressing
 328 neurons that were neighbors to the knockout cells (neighbors) (Fig. 1B, C). We found a
 329 significant decrease in NMDAR currents for NMDAR knockout cells (NMDAR KO; Cre-GFP)
 330 7 days after viral injection (Fig. 1C). These results confirm both the specificity and the
 331 effectiveness of the virally mediated NMDAR KO used in this study.

332 ***NMDAR KO abolishes experience-dependent homeostatic changes in synaptic strength***

333 Next we examined whether NMDARs play a role in homeostatic synaptic plasticity by
 334 measuring changes in the strength of excitatory synapses on V1 L2/3 pyramidal neurons
 335 following manipulations to visual experience in the presence or absence of NMDARs.
 336 Homeostatic synaptic plasticity has been characterized both *in vitro* and *in vivo* as an increase in
 337 synaptic strength after prolonged periods of decreased neuronal activity and a decrease in
 338 synaptic strength after periods of increased activity (O'Brien et al., 1998; Turrigiano et al., 1998;
 339 Desai et al., 2002; Goel and Lee, 2007). Previous studies have established that 2 days of visual
 340 deprivation in the form of dark exposure (DE) increases the strength of excitatory synaptic
 341 transmission, as measured by mEPSC amplitude, which is rapidly reversed by reinstating visual
 342 experience for a short period (light exposure, LE) (Goel and Lee, 2007; Gao et al., 2010).

343 In GFP-only control neurons, the average mEPSC amplitude was significantly increased
 344 after 2-days of DE and returned to normal-reared control (Ctl) values after 2-hours of LE (Fig.
 345 2A). Changes in the average mEPSC amplitude was also evident in the distribution of mEPSC
 346 amplitudes plotted in cumulative probability graphs (Fig. 2A), where the distribution of mEPSC
 347 amplitudes of control and DE group were significantly different while those between control and
 348

349 LE did not show statistical significance. Alterations in visual experience had no significant effect
350 on the average frequency of mEPSCs of GFP-only condition neurons (Fig. 2A). These results are
351 consistent with previous studies showing that homeostatic synaptic plasticity in V1 L2/3 mainly
352 manifests as postsynaptic change in AMPA receptors (Goel et al., 2006; Goel et al., 2011; He et
353 al., 2012).

354 In contrast, mEPSCs recorded from NMDAR KO neurons (Cre-GFP condition) lacked
355 regulation by changes in visual experience. There was no significant change in either the average
356 amplitude or frequency of mEPSCs across control, DE or LE groups (Fig. 2B). Moreover, we did
357 not observe any significant difference in mEPSC amplitude distribution across the 3 groups, as
358 shown by overlapping cumulative probability graphs (Fig. 2B). There was no significant
359 difference in basal mEPSC amplitude between normal-reared GFP-only controls and NMDAR
360 KO neurons (Student's t-test: $t=1.416$, $p=0.1713$). However, we noted a significant increase in
361 baseline frequency of mEPSC in control normal-reared NMDAR KO neurons when compared to
362 normal-reared GFP-only neurons (GFP-only: 2.5 ± 0.15 Hz, $n=9$; NR1 KO: 4.9 ± 0.57 Hz, $n=14$;
363 unpaired Student's t-test: $t=3.258$, $*p=0.0038$). This is similar to an observation made in a
364 previous study measuring mEPSCs from NMDAR KO neurons in CA1 (Adesnik et al., 2008),
365 which implicated NMDARs in regulating the number of functional synapses. Unexpectedly, data
366 from neighbor neurons, which did not express Cre-GFP and have intact NMDAR current (Fig.
367 1C), also failed to modulate the average mEPSC amplitude with changes in visual experience
368 (Fig. 2C). While there was no significant change in the average mEPSC amplitude across groups,
369 there was a statistically significant increase in the distribution of mEPSC amplitudes of LE group
370 as seen in the cumulative probability graph (Fig. 2C). In addition, unlike the NMDAR KO
371 neurons or GFP-only condition, these neighbor neurons showed significant increase in mEPSC

frequency after LE relative to control conditions (Fig. 2C). At this point, we cannot explain the phenotype of neighbor neurons except that NMDAR KO may not simply have a cell autonomous regulation of homeostatic plasticity in V1 L2/3 neurons. In any case, our results indicate that knocking out NMDARs prevents experience-dependent homeostatic synaptic plasticity and therefore support a necessary role of NMDARs in this process.

NMDAR function is required for experience-dependent homeostatic synaptic plasticity

Our data so far suggest that NMDAR KO prevents experience-dependent homeostatic synaptic plasticity. One caveat of our results from virally mediated NMDAR KO experiments is that NMDAR KO cells lacked upscaling by DE, hence we could not confirm whether NMDAR is required also for downscaling of mEPSCs when DE mice are re-exposed to light. In order to test this, a more acute manipulation of NMDAR function is needed hence we used pharmacology to acutely inhibit NMDAR function just during the LE. We blocked NMDAR function pharmacologically by administration of D-4-[(2*E*)-3-Phosphono-2-propenyl]-2-piperazinecarboxylic acid (d-CPP), which is a selective and competitive antagonist of NMDAR (Lehmann et al., 1987). Mice pertaining to the LE group were placed in the dark room for two days without any drug to allow normal scaling up process by DE, and then received d-CPP via an intraperitoneal injection (i.p., 10 mg/kg) 10 minutes before light re-exposure. To control for i.p. injection, a group of mice received the same volume of saline injection instead of d-CPP before light re-exposure. This design allows for specific testing of the role of NMDAR in LE-induced downscaling, which was not possible to determine in NMDAR KO condition where DE-induced upscaling was absent. In order to determine the effect of LE, we also had a group of control normal-reared and DE mice receiving saline. Because DE induced upscaling requires 2 days,

395 saline was administered into the lateral ventricle using an osmotic mini-pump coupled to a
396 cannula to allow 2 days of saline infusion during DE. Control normal-reared also received 2 days
397 of saline via osmotic mini-pump to control for any effect of osmotic mini-pump surgery. We
398 found that d-CPP injection right before LE prevented the normal downscaling of average
399 mEPSC amplitude (Fig. 3A), which was also evident when comparing the cumulative probability
400 of mEPSCs recorded from saline LE versus d-CPP LE groups (Fig. 3B). There was no
401 significant difference in average mEPSC frequency across groups (Fig. 3C). These results
402 suggest that NMDAR function is also required for downscaling mEPSCs in LE condition.

403 We also attempted to see whether blocking NMDAR function with d-CPP blocks scaling
404 up of mEPSCs with DE as observed in NMDAR KO neurons. This was to determine whether the
405 failure to scale up mEPSCs in NMDAR KO neurons is due to missing the NMDAR protein
406 itself, which is known to have a structural role in organizing downstream signaling molecules at
407 synapses via its intracellular domain (Sprengel et al., 1998; Kohr et al., 2003), or absence of
408 NMDAR function. For the purpose of this experiment d-CPP was perfused for 2 days into the
409 lateral ventricle through an osmotic minipump coupled to a cannula for Ctl normal-reared or DE
410 animals. Unexpectedly, we found that 2 days of d-CPP infusion in control normal-reared mice
411 significantly increased the amplitude of mEPSCs when compared to mice receiving the same
412 duration of saline (Fig. 3D). This presents difficulty in interpreting our results from d-CPP
413 treated DE mice (average mEPSC amplitude = 11.52 ± 0.45 pA, n=13). Despite the lack of
414 conclusive data from d-CPP treated DE group, our result from LE group suggests that functional
415 NMDARs are required for rapid downscaling mEPSCs with visual experience. Taken together
416 with our NMDAR KO data, these results support the previous conclusion that functional
417 NMDARs are required to undergo proper experience-dependent synaptic scaling.

418

419 *NMDAR disruption does not alter overall postsynaptic activity*

420 A potential concern regarding the lack of homeostatic adaptation observed with NMDAR
 421 function block is that it may reflect an inability of the network to modulate activity levels with
 422 visual experience. Changes in postsynaptic activity levels are thought to drive homeostatic
 423 synaptic plasticity (Ibata et al., 2008; Goold and Nicoll, 2010)(but see (Fong et al., 2015)).
 424 Therefore, we investigated whether disrupting NMDAR function altered the overall activity of
 425 V1 L2/3 neurons. To do this, we used the expression of the immediate early gene c-Fos as a
 426 proxy for neuronal activation under different conditions (Hoffman et al., 1993; Joo et al., 2016).

427 We first examined how global block of NMDAR function by d-CPP may have altered V1
 428 L2/3 neurons, hence compared c-Fos expression in V1 L2/3 neurons under Ctl, DE and LE
 429 conditions in saline and d-CPP infused mice (Fig. 4A-C). As in our mEPSC recording studies
 430 (Fig. 3), Ctl and DE group received saline or d-CPP via osmotic pump for 2 days, while LE
 431 group was placed in a darkroom for 2 days to allow normal DE-induced up-scaling and only
 432 received saline or d-CPP via i.p. injection 10 min before light re-exposure. V1 slices were co-
 433 stained for cFos and NeuN (neuronal marker), and the fraction of c-Fos positive neurons in L2/3
 434 were quantified by dividing the number of c-Fos positive cells by the number of NeuN positive
 435 cells. We found that the fraction of c-Fos positive neurons decreased with DE and returned to
 436 control levels with LE in saline infused mice, which is consistent with a reduction in activity
 437 levels with visual deprivation. We observed the same pattern of regulation in c-Fos positive
 438 neuronal fraction in mice infused with d-CPP, which suggests that a global block of NMDAR
 439 function in the whole brain does not grossly alter the level of neuronal activation leading to c-Fos
 440 expression in V1 L2/3.

441 To test if selective KO of NMDAR in V1 neurons also preserves neuronal activity across
 442 visual manipulations, we compared the fraction of c-Fos positive neurons following viral KO of
 443 NMDARs by expressing Cre-GFP or control-GFP in NR1^{fllox} mice (Fig. 4D-F). Here we only
 444 quantified the fraction of c-Fos positive cells among neurons that were co-labeled with GFP and
 445 NeuN, hence the quantification corresponds to activity of GFP or Cre-GFP transfected neurons.
 446 In V1 sections from control GFP expressing mice, we did not observe a clear down regulation of
 447 c-Fos expression in control GFP expressing neurons with DE, but there was significant increase
 448 in c-Fos expression with LE. In V1 sections from Cre-GFP expressing mice (NMDAR KO), we
 449 found that DE decreased c-Fos expression which returned to normal levels with LE similar to
 450 what we saw in saline infused animals. Despite the caveat that we cannot explain the lack of a
 451 significant decrease in c-Fos expression with DE in control GFP expressing neurons, our data
 452 nonetheless suggest that KO of NMDAR in neurons do not grossly alter their activity levels
 453 across different visual manipulations. Collectively, our results suggest that the disruption in
 454 homeostatic regulation of excitatory synapses in the absence of NMDAR function is not likely
 455 due to major alterations in the overall activity of V1 L2/3 neurons.

456 Assessing neural activity with immediate early gene expression, such as c-Fos, has innate
 457 limitation in that they can only monitor whether the neural activity is beyond the threshold for
 458 immediate early gene induction, but cannot determine fine scale changes in neural activity. In
 459 order to more directly measure the effect of NMDAR blockade on V1 L2/3 neuronal activity, we
 460 performed *in vivo* calcium imaging from awake mice before and after CPP injection (Fig. 5A).
 461 Individual neurons sometimes displayed differences in response to moving sinusoidal gratings
 462 pre and post CPP injection (Fig. 5B). However, the distribution of these differences was not
 463 significantly different from that for control mice receiving saline for spontaneous activity

464 measured during blank screen presentation (Fig. 5C left panel) or activity measured during
465 presentation of visual stimuli (Fig. 5C middle panel). This also held true if we restricted analysis
466 only to neurons that were significantly visually activated (Fig. 5C right panel). NMDA spikes
467 have previously been implicated in stimulus selectivity of layer 4 neurons in the barrel cortex
468 (Lavzin et al., 2012), so we considered the possibility that CPP might affect orientation tuning of
469 the neurons (Fig. 5D). The response at preferred orientation did not change significantly in mice
470 receiving saline versus CPP (Fig. 5D left panel). Likewise, the changes in orientation selectivity
471 index were not significant (Fig 5D right panel), although there was a trend that CPP may have
472 affected OSI in a small subpopulation of L2/3 neurons. Overall, our results indicate that blocking
473 NMDARs with CPP is unlikely to grossly affect responses of V1 L2/3 neurons in a way that
474 could account for the observed effects on plasticity.

475

476 Discussion

477 The mechanisms of homeostatic adaptation to changes in sensory experience has been
478 largely explained by two distinct models: sliding threshold and synaptic scaling. While these two
479 models are considered different, there is emerging body of recent work suggesting that these two
480 modes of homeostatic plasticity may share similarities and may interact with each other to
481 regulate synaptic strength (see (Fox and Stryker, 2017; Keck et al., 2017) for recent discussions).
482 Furthermore, there is evidence that sliding threshold or synaptic scaling mode of homeostatic
483 plasticity may be employed depending on the regime of activity changes *in vivo* (Bridi et al.,
484 2018). Based on the consensus of the field that synaptic scaling is largely independent of
485 NMDAR activation, here we tested the role of NMDARs in homeostatic synaptic plasticity
486 induced *in vivo* by changes in visual experience. By disrupting NMDAR function through cell

specific knock-out or pharmacology, we showed that the principal neurons within L2/3 of V1 require functional NMDARs to undergo proper visual experience-dependent homeostatic adaptation of excitatory synaptic strength. Disruption of normal experience-dependent homeostatic plasticity by removing NMDAR function was not due to gross changes in the activity of V1 L2/3 neurons. Our results support a role for NMDAR-dependent mechanisms in homeostatic synaptic plasticity induced *in vivo*.

We observed that NMDAR function is required for proper scaling of mEPSCs in V1 L2/3 neurons following a few days of DE or brief re-exposure to light (LE). While previous studies have shown that there is co-regulation of NMDAR together with AMPAR under inactivity conditions (Watt et al., 2000) and changes in NMDAR function after visual deprivation (Quinlan et al., 1999; Philpot et al., 2003; Guo et al., 2012), homeostatic synaptic scaling is largely thought of as occurring independent of NMDAR activity (Turrigiano et al., 1998; Turrigiano, 2008) and is absent in several genetic models with preserved LTP/LTD (Stellwagen and Malenka, 2006; Hu et al., 2010). It has previously been proposed that a switch in NMDAR function slides the synaptic modification threshold for LTP/LTD induction after DE (Quinlan et al., 1999; Philpot et al., 2003; Guo et al., 2012). Our results suggest a possibility that scaling up of mEPSCs could be a consequence of sliding down of synaptic modification threshold. Lowered synaptic modification threshold by DE would promote LTP across a large population of synapses, and hence manifest as global scaling up of excitatory synapses. This would imply that the amount of activity in V1 under DE condition is sufficient to act on the lowered threshold to induce NMDAR-dependent LTP. This contradicts *in vitro* studies done in cultured neurons, where prolonged blockade of action potentials was able to scale up excitatory synapses. Our data suggest that this may not be the case *in vivo*, where we surmise there may be sufficient activity in

510 the deprived cortex that can activate NMDARs to potentiate synaptic strength across a large
 511 number of synapses following a reduction in LTP threshold. Indeed, we recently reported that
 512 homeostatic upscaling of mEPSCs in V1 L2/3 neurons with DE is dependent on spontaneous
 513 activity and upregulation of GluN2B (Bridi et al., 2018). GluN2B containing NMDARs have a
 514 longer current duration (Monyer et al., 1994), which may enable better integration of lower
 515 frequency activity as occurs during spontaneous firing. Our current data add to this by showing
 516 that NMDAR function in V1 L2/3 neurons is necessary for upscaling with DE using cell-type
 517 specific genetic KO, and also demonstrate that NMDAR function is necessary for scaling down
 518 of mEPSCs with LE.

519 In addition to disrupting normal visual experience-dependent regulation of mEPSCs, we
 520 unexpectedly observed that blocking NMDAR function with d-CPP for 2 days significantly
 521 scales up mEPSCs under control normal-reared conditions. This suggests that there may be on-
 522 going activation of NMDARs by normal visual experience that actively reduces mEPSCs. This is
 523 reminiscent of what we observed in Arc KOs, which also show larger basal mEPSCs in V1 L2/3
 524 neurons under normal-reared conditions and lack visual experience-dependent regulation of
 525 mEPSCs (Gao et al., 2010). While d-CPP scaled up mEPSCs under basal conditions, we did not
 526 observe an increase in mEPSC amplitude in the NMDAR KO neurons (Fig. 2). It seems unlikely
 527 that the absence of basal “upscaling” in NMDAR KO was simply due to the incomplete removal
 528 of pre-existing NMDARs during the 6-7day Cre expression because the NMDAR current is
 529 mostly absent at that time point (Fig. 1). It is possible that the difference may have resulted from
 530 d-CPP having a more global effect while NMDAR KO was done in a cell-type specific manner
 531 in V1 L2/3. We cannot rule out an alternative possibility that complete blockade of NMDAR
 532 triggers mEPSCs to potentiate, which would be consistent with some of the *in vitro* studies

533 showing NMDAR blockade facilitates scaling up of synapses (Sutton et al., 2006; Aoto et al.,
534 2008) by blocking spontaneous Ca^{2+} transients mediated by NMDARs (Reese and Kavalali,
535 2015). Despite such caveats, the fact that both methods detect failures in experience-dependent
536 homeostatic regulation suggests that it is the function of NMDARs that is critical for proper
537 regulation of AMPAR-mEPSCs by changes in visual experience.

538 Despite initial studies showing induction of synaptic scaling as being largely independent
539 of NMDARs, there is some evidence from previous studies *in vitro* demonstrating interaction
540 between NMDAR function and synaptic scaling. For example, blocking NMDARs has been
541 shown to accelerate synaptic upscaling in cultured neurons (Sutton et al., 2006) and decreasing
542 NMDAR calcium permeability has been shown to downscale AMPAR currents (Pawlak et al.,
543 2005). Other potential mechanism relating NMDAR activation with scaling involve
544 “unsilencing” of synapses after activity blockade *in vitro* that promotes further LTP induction
545 (Arendt et al., 2013). These findings suggest that NMDAR activity can also have profound
546 influence on synaptic scaling mechanisms in addition to sliding the threshold for LTP/LTD.

547 Synaptic scaling has been largely considered a cell-autonomous process that is triggered
548 by readout of postsynaptic spikes, or more precisely postsynaptic depolarization (Ibata et al.,
549 2008; Goold and Nicoll, 2010). For example, optogenetic activation of postsynaptic neurons was
550 shown to be sufficient to drive down-scaling of excitatory synapses (Goold and Nicoll, 2010),
551 and blocking somatic spikes was sufficient to scale up synapses (Ibata et al., 2008). However, a
552 recent study suggested that synaptic scaling is not dependent on postsynaptic spike rate *per se*,
553 but due to changes in glutamatergic inputs (Fong et al., 2015). This suggests that the level of
554 activation of glutamate receptors could ultimately be the condition monitored by neurons in
555 order to trigger homeostatic adaptation. Our data is consistent with the latter, where

556 glutamatergic transmission level may be detected by the activation of NMDARs to induce
557 changes in AMPAR-mediated mEPSCs.

558 Our data provide *in vivo* evidence for a homeostatic mechanism that requires NMDAR
559 activation in order to cope with changes in visual experience. These results suggest either a role
560 for NMDAR function in *in vivo* synaptic scaling or that homeostatic scaling of synapses is a
561 manifestation of Hebbian forms of plasticity triggered by lowered or increased synaptic
562 modification threshold according to changes in visual experience. If it is the latter, a major
563 implication is that *in vivo* homeostasis could be implemented in an input specific manner.
564 Indeed, DE has been shown to increase the synaptic strength of lateral intracortical inputs to V1
565 L2/3 neurons without affecting those originating from L4 (Petrus et al., 2015), which could be
566 the basis for non-multiplicative synaptic scaling observed in adult V1 (Goel and Lee, 2007). A
567 recent study reported that homeostatic scaling of dendritic spines in V1 following monocular
568 enucleation occurs in a dendritic branch specific manner (Barnes et al., 2017), which is
569 consistent with input specific regulation. Input-specific homeostatic synaptic plasticity has also
570 been observed in the hippocampus where inactivity scales up excitatory synaptic transmission in
571 feedforward synapses while decreasing mEPSC frequency in recurrent synapses within CA3
572 (Kim and Tsien, 2008). Collectively, these observations suggest that input-specific homeostatic
573 adaptation is likely a general phenomenon across different brain circuits that receive input from
574 several sources. Synapse-specific homeostatic plasticity has been observed in reduced
575 preparations where activity was selectively manipulated at individual synapses via genetic
576 methods of silencing specific synapses (Hou et al., 2008; Lee et al., 2010; Beique et al., 2011).
577 One such study demonstrated that NMDAR subunit composition can be selectively altered at
578 individual synapses (Lee et al., 2010), which allows for synapse-specific adjustment of sliding

579 threshold. Such input-specific homeostatic control will allow cortical neurons that participate in
580 multiple functional circuits to adapt effectively to changes in select inputs to provide stability
581 without compromising the function of other synapses. Our results suggest that one way to
582 achieve such input-specific homeostatic control is via NMDAR-dependent plasticity
583 mechanisms.

584

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712 **Figure Legends**

713 **Figure 1. Neuron-specific NMDAR knockout**

714 **(A)** GFP and Cre-GFP expressing viruses result in similar transfection efficiency for V1 neurons

715 (mean percentage of transfected neurons: GFP-only = $67.7 \pm 4.6\%$, Cre-GFP = $60.8 \pm 3.9\%$;

716 unpaired t-test: $t=1.634$, $p=0.1079$; number of slices quantified reported, 6 mice per condition).

717 **(B)** Confocal image of biocytin filled NMDAR KO (green arrow, expressing Cre-GFP in the

718 nucleus) and neighbor (orange arrow) neurons in V1 L2/3.

719 **(C)** Left: Comparison of average NMDAR/AMPA ratio for each condition (mean

720 NMDAR/AMPA ratio for GFP-only 0.25 ± 0.04 , NR1 KO 0.05 ± 0.007 , neighbor 0.22 ± 0.07 ;

721 one-way ANOVA: $F(2,13)=5.502$, $*p=0.0186$; Tukey's multiple comparison *post hoc*: GFP vs

722 NR1 KO $*p=0.02$, GFP vs neighbor $p=0.93$, neighbor vs NR1 KO $p=0.05$; number of cells

723 reported). Right: Example traces of NMDAR (measured at +40 mV holding potential) and

724 AMPAR (measured at -70 mV holding potential) mediated currents in GFP-only, NMDAR KO

725 and neighbor neurons. Traces were normalized to match the amplitude of AMPAR current.

726

727 **Figure 2. NMDAR knockout neurons lack homeostatic regulation of mEPSCs**

728 Top row: Average mEPSC traces from normal-reared (Ctl), 2 days DE, and 2 hours LE. Second

729 row: Bar graph comparison of average mEPSC amplitude from Ctl, DE and LE groups. Average

730 mEPSC amplitude for each cell is plotted in gray circle. Third row: Cumulative probability

731 graphs of mEPSC amplitudes from Ctl (gray solid line), DE (black solid line, and LE (black

732 dashed line). Fourth row: Example mEPSC recording traces from Ctl (top), DE (middle) and LE

733 (bottom). Bottom row: Comparison of average mEPSC frequency across Ctl, DE and LE groups.

734 Average mEPSC frequency for each cell is plotted in gray circle.

735 **(A)** Results from neurons transfected with GFP-only virus. There was a significant increase in
 736 average mEPSC amplitude with DE which is reversed by LE (mean amplitude: Ctl = $11.38 \pm$
 737 0.51 pA, DE = 13.29 ± 0.55 pA, LE = 10.59 ± 0.49 pA; one-way ANOVA: $F(2,26)=7.213$,
 738 $*p=0.0032$; Tukey's multiple comparison *post-hoc*: Ctl vs DE $*p=0.040$, Ctl vs LE: $p=0.574$, DE
 739 vs LE $*p=0.003$). DE significantly shifted the cumulative probability curve of mEPSC
 740 amplitudes to the right with DE (Kolmogorov-Smirnov test: $*p<0.0001$), which then returned to
 741 Ctl distribution with LE (Kolmogorov-Smirnov test: $p=0.0138$). There was no significant change
 742 in mEPSCs frequency across groups (mean frequency: Ctl = 2.43 ± 0.14 Hz, DE = 2.92 ± 0.28
 743 Hz, LE = 3.30 ± 0.53 Hz; one-way ANOVA: $F(2,26)=1.196$, $p=0.3192$).

744 **(B)** Results from Cre-GFP transfected neurons. NMDAR KO neurons failed to undergo
 745 significant changes in average mEPSC amplitude (Ctl = 12.85 ± 0.76 pA, DE = 12.55 ± 0.95 pA,
 746 LE = 11.96 ± 0.86 pA; one-way ANOVA: $F(2,35)=0.2863$, $p=0.7528$). Despite no difference in
 747 average mEPSC amplitude, cumulative probability curves of mEPSC amplitudes from Cre-GFP
 748 expressing NMDAR KO neurons show a slight shift towards higher amplitudes in Ctl group
 749 relative to DE (Kolmogorov-Smirnov test: $p=0.0005$) and LE (Kolmogorov-Smirnov test:
 750 $p=0.0005$), but there was no significant difference between DE and LE (Kolmogorov-Smirnov
 751 test: $p=0.1441$). There was no significant change in the average mEPSC frequency across groups
 752 (Ctl = 4.92 ± 0.57 Hz, DE = 4.49 ± 0.48 Hz, LE = 4.93 ± 0.58 Hz; one-way ANOVA:
 753 $F(2,35)=0.1947$, $p=0.8240$).

754 **(C)** Results from neighbor neurons that did not express Cre-GFP. There was no significant
 755 change in the average mEPSCs amplitude across groups (Ctl = 11.71 ± 0.88 pA, DE = $11.47 \pm$
 756 0.36 pA, LE = 13.86 ± 0.97 pA; one-way ANOVA: $F(2,25)=2.465$, $p=0.1054$). While there was
 757 no significant difference in cumulative probability of mEPSC amplitudes between Ctl and DE

values (Kolmogorov-Smirnov test: $p=0.0168$), there was a significant shift towards larger values for LE when compared to DE (Kolmogorov-Smirnov test: $*p < 0.0001$) or Ctl (Kolmogorov-Smirnov test: $*p < 0.0001$). There was a significant change in the average mEPSC frequency after LE relative to control conditions (Ctl = 3.28 ± 0.55 Hz, DE = 5.15 ± 0.62 Hz, LE = 6.57 ± 0.71 Hz; one-way ANOVA $*p=0.0034$, $F(2,25)=7.182$; Tukey's multiple comparison *post hoc*: Ctl vs DE $p=0.091$, Ctl vs LE $*p=0.003$, DE vs LE: $p=0.291$).

Figure 3. Lack of visual experience-dependent scaling down of mEPSCs with NMDAR blockade

(A) Comparison of average mEPSC amplitude changes in saline treated groups and LE group that was treated with d-CPP to block NMDAR activity. Left: Average mEPSC traces from saline treated normal-reared (Ctl), 2 days DE, and 2 hours LE compared to d-CPP treated LE group. Right: Bar graph comparison of average mEPSC amplitude from saline treated Ctl, DE and LE groups compared to d-CPP treated LE group (Saline Ctl = 10.55 ± 0.42 pA, Saline DE = 12.14 ± 0.35 pA, Saline LE = 10.69 ± 0.55 pA, CPP LE = 12.35 ± 0.73 pA; ANOVA: $F(3,35) = 3.139$, $*p = 0.0375$; Bonferroni's Multiple Comparison test: Saline LE vs CPP LE $*p < 0.05$). Average mEPSC amplitude for each cell is plotted as gray circle.

(B) Cumulative probability of mEPSC amplitudes from saline treated LE group (black dashed line) and d-CPP treated LE group (black solid line) show statistically significant difference (Kolmogorov-Smirnov test: $*p < 0.0001$)

(C) There was no significant difference in average mEPSC frequency across groups (Saline Ctl = 4.62 ± 0.85 Hz, Saline DE = 5.41 ± 0.64 Hz, Saline LE = 4.05 ± 0.61 Hz, CPP LE = 5.62 ± 0.82

780 Hz; ANOVA: $F(3,35) = 0.9418$, $p = 0.4309$). Average mEPSC frequency for each cell is plotted
781 as gray circle.

782 **(D)** Treatment of d-CPP for 2 days in control normal-reared mice significantly increased the
783 average mEPSC amplitude compared to saline treated controls. Left: Average mEPSC traces.
784 Middle: Comparison of average mEPSC amplitude (saline Ctl: 10.55 ± 0.42 pA, $n=9$; d-CPP Ctl:
785 13.19 ± 0.52 pA, $n=10$; unpaired Student's t -test $t(17)=3.88$, $*p=0.0012$). Right: Comparison of
786 average mEPSC frequency (saline Ctl: 4.62 ± 0.85 Hz, $n=9$; d-CPP Ctl: 5.59 ± 0.73 Hz, $n=12$;
787 unpaired Student's t -test $t(19)=0.85$, $p=0.4031$).
788

789 **Figure 4. Comparison of neuronal activity measured with c-Fos expression**

790 **(A-C)** Comparison of c-Fos expression in V1 L2/3 neurons from Ctl, DE, and LE mice treated
791 with saline or d-CPP. Representative confocal images of V1 L2/3 from Ctl, DE, and LE mice
792 receiving saline **(A)** or d-CPP **(B)** infusion. Sections were stained with NeuN (green) and c-Fos
793 (magenta). Scale bars: 36 μ m. **(C)** Comparison of quantified fraction of c-Fos positive neurons,
794 which were calculated as (number of c-Fos positive neurons)/(number of NeuN positive neurons).
795 For saline group, DE significantly decreased the fraction of c-Fos positive neurons, which
796 increased to Ctl values with LE (Ctl = 0.62 ± 0.03 , DE = 0.14 ± 0.02 , LE = 0.63 ± 0.04 ; ANOVA:
797 $F(2,33) = 103.24$, $*p < 0.0001$; Tukey's multiple comparison *post hoc*: Ctl vs DE $*p < 0.0001$,
798 Ctl vs LE $p = 0.9931$, DE vs LE $*p < 0.0001$). For d-CPP group, DE significantly decreased the
799 fraction of c-Fos positive neurons, which increased to Ctl values with LE (Ctl = 0.54 ± 0.32 , DE
800 = 0.13 ± 0.02 , LE = 0.58 ± 0.02 ; ANOVA: $F(2,32) = 93.5424$, $*p < 0.0001$; Tukey's multiple
801 comparison *post hoc*: Ctl vs DE $*p < 0.0001$, Ctl vs LE $p=0.4234$, DE vs LE $*p < 0.0001$).

802 **(D-F)** Comparison of c-Fos expression in GFP transfected V1 L2/3 neurons from Ctl, DE or LE
 803 mice. Representative confocal images of V1 L2/3 from NR1-flox mice that received viral
 804 transfection of GFP-only **(D)** or Cre-GFP (NMDAR KO) **(E)**. Sections were stained with NeuN
 805 (blue) and c-Fos (red). GFP expression from either GFP-only or Cre-GFP is shown in green.
 806 Scale bars: 36 μm . **(F)** Comparison of the fraction of c-Fos positive GFP neurons, which were
 807 calculated as (number of c-Fos and GFP positive neurons)/(number of total GFP positive
 808 neurons). For GFP-only group, there was a significant increase in the fraction of c-Fos positive
 809 GFP neurons following LE compared to DE (Ctl = 0.35 ± 0.04 , DE = 0.25 ± 0.04 , LE = $0.54 \pm$
 810 0.09 ; ANOVA: $F(2,15)=6.35$, $*p=0.0100$; Tukey's multiple comparison test: Ctl vs DE $p=0.5654$,
 811 Ctl vs LE $p=0.1503$, DE vs LE $**p=0.0077$). In NMDAR KO (Cre-GFP) group, there was a
 812 significant decrease in the fraction of c-Fos positive Cre-GFP neurons in DE, which increased
 813 back to Ctl levels with LE (Ctl = 0.49 ± 0.03 , DE = 0.31 ± 0.06 , LE = 0.56 ± 0.08 ; ANOVA:
 814 $F(2,16)=5.52$, $*p=0.0149$; Tukey's multiple comparison test: Ctl vs DE $*p=0.04$, Ctl vs LE
 815 $=0.6629$, DE vs LE $*p=0.0212$).

816
 817 **Figure 5. Blocking NMDARs with d-CPP injection does not acutely affect visually evoked**
 818 **responses in V1**

819 **(A)** Example of imaged region in an Emx1-Ai96 mouse before (left) and after (right)
 820 intraperitoneal CPP injection. Both images are maximum projections of GCaMP6s fluorescence.
 821 **(B)** Visually evoked responses from ROIs shown in (A) before (gray) and after (red) CPP
 822 injection. The dashed line denotes onset of the visual stimulus and its height corresponds to 0.5
 823 $\Delta F/F_0$. Each trace is 11 seconds long.

824 **(C)** Distribution of visual response change after application of saline (blue) versus CPP (red).
825 (Left panel) Comparison of changes in spontaneous activity of all neurons before and after saline
826 or CPP injection. Spontaneous activity was measured during a 7 s window when a blank screen
827 was presented. N=107 neurons across 7 mice for saline and N=81 neurons across 5 mice for CPP.
828 Kolmogorov-Smirnov test, $P=0.448$. (Middle panel) Response change for all activated neurons,
829 regardless of whether the level of activation is significant. The same neurons as in the left panel.
830 Kolmogorov-Smirnov test, $P=0.087$. (Right panel) Response change only for significantly
831 visually activated neurons. N=36 neurons across 7 mice for saline and N=36 neurons across 6
832 mice for CPP. Kolmogorov-Smirnov test, $P=0.46$.
833 **(D)** (Left panel) Change of response at preferred orientation after application of saline (blue)
834 versus CPP (red). Unpaired t-test, $P=0.24$. (Right panel) Change of orientation selectivity after
835 application of saline (blue) versus CPP (red). Unpaired t-test, $P=0.069$. Only neurons with
836 significant orientation tuning were included in this analysis (N=30 neurons for saline conditions
837 and N=28 neurons for CPP condition).

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