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1	Robustness of STDP to spike timing jitter
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15 ABSTRACT

16 In Hebbian plasticity, neural circuits adjust their synaptic weights depending on patterned firing. Spike-timing-dependent plasticity (STDP), a synaptic Hebbian learning rule, relies on 17 18 the order and timing of the paired activities in pre- and postsynaptic neurons. Classically, in 19 ex vivo experiments, STDP is assessed with deterministic (constant) spike timings and time 20 intervals between successive pairings, thus exhibiting a regularity that differs from biological 21 variability. Hence, STDP emergence from noisy inputs as occurring in *in vivo*-like firing 22 remains unresolved. Here, we used noisy STDP pairings where the spike timing and/or 23 interval between pairings were jittered. We explored with electrophysiology and 24 mathematical modeling, the impact of jitter on three forms of STDP at corticostriatal 25 synapses: NMDAR-LTP, endocannabinoid-LTD and endocannabinoid-LTP. We found that 26 NMDAR-LTP was highly fragile to jitter, whereas endocannabinoid-plasticity appeared more 27 resistant. When the frequency or number of pairings was increased, NMDAR-LTP became 28 more robust and could be expressed despite strong jittering. Our results identify 29 endocannabinoid-plasticity as a robust form of STDP, whereas the sensitivity to jitter of 30 NMDAR-LTP varies with activity frequency. This provides new insights into the mechanisms 31 at play during the different phases of learning and memory and the emergence of Hebbian 32 plasticity in *in vivo*-like activity.

33 INTRODUCTION

34 Long-term changes in synaptic efficacy between neurons are thought to underlie learning and memory¹ and can be assessed with a synaptic Hebbian paradigm such as spike timing-35 dependent plasticity (STDP)²⁻⁸. In STDP, the occurrence of timing-dependent-long-term 36 37 potentiation (tLTP) or -depression (tLTD) is the result of three factors. The two first ones are 38 the two coincident activity patterns on either side of the synapse, which depends on 1) the relative timing between pre- and postsynaptic spikes $(\Delta t_{\text{STDP}})^{4,8}$, 2) the dendritic and axonal 39 propagation delays⁹, 3) the number of paired spikes $(N_{\text{pairings}})^{10-13}$, 4) the frequency of the 40 paired spikes $(F_{\text{pairings}})^{6,7,10,11,14}$ and 5) membrane depolarization^{14,15}; the third factor involves 41 in STDP expression mainly refers to neuromodulators^{16,17} and glial cells¹⁸. 42

43 STDP has been attracting a lot of interest in computational and experimental neurosciences 44 because it relies on spike correlation and has emerged as a candidate mechanism for experience-dependent changes in neural circuits, including map plasticity^{8,19–23}. STDP is 45 46 classically investigated using regular repetitions of the same spike timing and fixed intervals 47 between successive paired stimulations. A typical experimental protocol consists in pairing pre- and postsynaptic stimulations with a fixed Δt_{STDP} (ranging from -30 to +30ms for 48 plasticity induction); $\Delta t_{\text{STDP}} < 0$ when the postsynaptic stimulation occurs before the paired 49 50 presynaptic one (post-pre pairings), whereas $\Delta t_{\text{STDP}} > 0$ when the presynaptic stimulation 51 occurs before the postsynaptic one (pre-post pairings). These pairings are then repeated 52 between 15 and 200 times (between 0.1 and 5Hz) with the spike timing and the time interval 53 between successive pairings being kept constant.

Regular stimulation paradigms produce patterns of activity that are likely to differ from the variability expected in *in vivo*-like firing. *In vivo* paired natural visual stimulations with postsynaptic action potential lead to Hebbian STDP with specific characteristics when compared to *ex vivo* STDP (broader membrane potential changes and STDP window), which

can be explained by jittered presynaptic inputs occurring in $vivo^{22}$. A theoretical study has 58 59 started to explore naturalistic stimulations and showed that when neurons fire irregularly, the impact of spike timing in plasticity becomes weaker than the influence of the firing rate²⁴. 60 61 However, whether STDP emergence and maintenance is robust against biological variability remains to be investigated. To address this question, it is important to take into account 62 63 various forms of STDP, involving distinct intracellular signal transduction pathways, i.e. NMDAR-, mGluR- or endocannabinoid(eCB)-mediated STDP²⁵⁻²⁹ (for reviews see^{8,30,31}). It 64 is thus expected that those various STDP forms might exhibit different robustness to spike 65 66 train variability.

Here, we show with patch-clamp recordings and mathematical modeling that corticostriatal NMDAR- and eCB-STDP do not exhibit the same sensitivity to noisy spike timings: eCBplasticity (eCB-tLTD and eCB-tLTP) appeared robust to jittering whereas NMDAR-tLTP was fragile. However, increasing the number or frequency of pairings improved NMDAR-tLTP robustness. Our results further indicate that the robustness of NMDAR-tLTP to jitter is also strengthened by the irregularity of the spike-train stimulations.

73 **RESULTS**

Corticostriatal synapses exhibit a bidirectional STDP in which NMDAR-tLTP, eCB-74 $tLTD^{29,32-34}$ or eCB-tLTP^{12,13} are induced depending on the number of pairings (N_{pairings}) and 75 the spike timing (Δt_{STDP}). While STDP is classically investigated using fixed Δt_{STDP} , we 76 77 examine here the effect of noisy Δt_{STDP} , closer to *in vivo*-like firing. Here, we define the Inter-78 Pairing-Interval (IPI) as the time interval between two consecutive presynaptic stimulations, 79 whereas Δt_{STDP} is the time interval between the post- and the presynaptic stimulation times within a given paired stimulation (in a "standard" STDP protocol both IPIs and Δt_{STDP} are 80 81 constant).

82

83 NMDAR- and eCB-mediated STDP triggered with fixed Δt_{STDP}

84 We observed bidirectional asymmetric STDP in MSNs (Fig. 1a): 100 post-pre pairings 85 $(\Delta t_{\text{STDP}} < 0)$ induced tLTP whereas 100 pre-post pairings ($\Delta t_{\text{STDP}} > 0$) induced tLTD. An 86 example of the tLTP induced by 100 post-pre pairings with fixed Δt_{STDP} (Fig. 1b) is illustrated 87 in Figure 1c1. The input resistance Ri remained stable over this period. Although we fixed the 88 values of Δt_{STDP} , they slightly vary from one pairing to another (Fig.1b). This low jitter can be written formally as $\Delta t_{\text{STDP}} = m_{\Delta t} + \xi_{\Delta t}$, were $m_{\Delta t}$ is the mean spike timing and $\xi_{\Delta t}$ is a 89 90 random variable with mean 0 and standard deviation $\sigma_{\Delta t}$. Overall, 100 post-pre pairings 91 induced tLTP (mean EPSC amplitude recorded 60 min after protocol induction: 149±13% of baseline, p=0.0028, n=13) (Fig. 1c2). $\sigma_{\Delta t}$ was 1.4±0.2ms (Fig. 1c3). This tLTP was NMDAR-92 93 mediated since prevented by D-AP5 (50 μ M), a NMDAR antagonist, (88±12%, p=0.375, n=5; 94 p < 0.0001 when compared with tLTP in control) (Fig. 1c2). Conversely, pre-post pairings 95 induced tLTD, as shown in the example in Figure 1d1. Overall, 100 pre-post pairings induced 96 tLTD (76±7%, p=0.0070, n=10) (Fig. 1d2). $\sigma_{\Delta t}$ was 1.4±0.2ms (Fig. 1d3). This tLTD was

97 CB₁R-mediated since prevented by AM251 (3μ M), a CB₁R specific inhibitor ($99\pm2\%$, 98 *p*=0.816, n=5; *p*<0.0001 when compared with tLTD in control) (Fig. 1d2). We thus find an 99 anti-Hebbian polarity for corticostriatal STDP. We previously showed that GABA operates as 100 a Hebbian/anti-Hebbian switch at corticostriatal synapses^{34,35} and corticostriatal STDP 101 polarity depends on the presence^{32,33} or the absence^{12,29,36,37} of GABA_A receptor antagonists.

102 Besides this bidirectional STDP (NMDAR-tLTP and eCB-tLTD) induced for 100 pairings, 103 we reported that low numbers of pairings ($N_{\text{pairings}} = 5-15$) induce an eCB-tLTP, dependent on the activation of $CB_1R^{12,13}$. Figure 1e1 shows an example of tLTP induced by 10 post-pre 104 105 pairings. Overall, 10 post-pre STDP pairings induced tLTP (145±9%, p=0.0005, n=15), which 106 was prevented by AM251 (3μ M) (76±10%, p=0.080, n=4; p<0.0001 when compared with tLTP in control) (Fig. 1e2) as recently reported¹². $\sigma_{\Delta t}$ for 10 post-pre parings STDP was 107 108 1.8±0.3ms (n=15) (Fig. 1e3). Note that there was no difference between the $\sigma_{\Delta t}$ measured for 109 NMDAR-tLTP, eCB-tLTD and eCB-tLTP (one way ANOVA, p=0.4094).

110

111 A computer model emulates NMDAR- and eCB-mediated STDP

112 To help interpret our experimental results, we used a mathematical model of the signaling pathways implicated in corticostriatal STDP, including NMDAR- and CB1R-plasticity^{12,29,32,33} 113 114 (Fig. 1f1). The model is detailed in S1 Text, with parameter values listed in S1 and S2 Tables. 115 We include in the Supporting Information a thorough description of the mechanisms by which 116 eCB- and NDMAR-plasticity are expressed in the model when STDP protocol (N_{pairings}=100) 117 is applied (S3 Text). Figure 1f2 shows the value of the total synaptic weight (W_{total}) predicted 118 by the model with various N_{pairings} and Δt_{STDP} . In agreement with the experimental data, with 119 small $\sigma_{\Delta t}$ (3ms), the model features three main plasticity regions: tLTD was observed for short pre-post pairings ($0 < \Delta t_{\text{STDP}} < +30 \text{ms}$), whereas short post-pre pairings ($-30 < \Delta t_{\text{STDP}} < 0 \text{ms}$) 120

induced a first tLTP region for low numbers of pairings ($3 < N_{pairings} < 25$) and a second tLTP region emerging for $N_{pairings} > 50$. Blocking the NMDAR-CaMKII pathway in the model (Fig. 1r3) suppresses the second tLTP region whereas blocking CB₁R activation (Fig. 1f4) prevents the expression of tLTD and of the first tLTP region. Therefore, the model emulates both CB₁R-tLTD and -tLTP, in agreement with the eCB-tLTD and eCB-tLTP illustrated in the experimental data of Figure 1d and 1e, respectively. The model also faithfully emulates the appearance of NMDAR-tLTP for $N_{pairings} > 50$.

128

129 Predicting the effects of jittered STDP with a mathematical model

130 The small jitter observed in the above experiments was the result of biological variability 131 inherent to the experimental setup. Our next goal was to extend this study to larger jitters $(3 < \sigma_{\Delta t} < 10 \text{ms})$ that we controlled experimentally. We first used constant intervals between 132 133 two consecutive pairings (Inter-Pairing Interval, IPI) and used the same definition as above for jittered Δt_{STDP} (*Methods*): $\Delta t_{\text{STDP}} = m_{\Delta t} + \xi_{\Delta t}$, where $\xi_{\Delta t}$ is a random variable with zero 134 135 mean and $m_{\Delta t}$ is the expected value of the spike timing Δt_{STDP} over the N_{pairings} . Next, we adjusted the probability distribution function and the variance of $\xi_{\Delta t}$ to perform a parametric 136 137 exploration of the impact of jitter.

In Figure 2, the jitter $\xi_{\Delta t}$ was sampled from a uniform distribution in $[-\sigma_{\Delta t}, +\sigma_{\Delta t}]$ (Fig. 2a1). STDP in the model was globally not robust to jitter, since the three forms of plasticity vanished when $\sigma_{\Delta t}$ was large enough (Fig. 2a2). However, the model delivered the prediction that the three forms of plasticity do not display the same sensitivity to jitter. In particular, NMDAR-tLTP is predicted to be very fragile when subjected to $\sigma_{\Delta t}$: its amplitude decreases as soon as $\sigma_{\Delta t}$ >2ms and completely vanishes for $\sigma_{\Delta t}$ >4ms (Fig. 2a2 and 2b2). In contrast, eCB-plasticity is much less sensitive: eCB-tLTP is still present for jitters as large as 7-8ms and eCB-tLTD is still expressed with $\sigma_{\Delta t} = 10$ ms (Fig. 2b3).

146 We next evaluated the validity of this prediction when the probability distribution function of 147 the jitter changes. Figure 2 illustrates the results we obtained with Gaussian (Fig. 2c) or 148 triangular (Fig. 2d) distributions of $\sigma_{\Delta t}$ (see *Methods*). In both cases, the profiles of the 149 robustness curves (Fig. 2c2 and 2d2) are almost identical to each other and similar to the 150 robustness curves obtained with a uniform distribution (Fig. 2a2). In Figure 2e, we combined a triangular distribution for $\sigma_{\Delta t}$ with random IPIs, i.e. in this case, stochasticity is applied not 151 152 only to the timing between the two stimulations of a given pairing, but also to the time 153 interval between two consecutive pairings: IPI distribution was Poisson with refractory period 154 τ_r and rate λ (see *Methods*). Despite this increased stochasticity, we found that eCB-tLTD 155 was again more robust, tolerating jitters up to 6-7ms (eCB-tLTP) or 10ms (eCB-tLTD) 156 whereas NMDAR-tLTP was fragile, vanishing for $\sigma_{\Delta t}$ >4ms (Fig. 2e2). Therefore, according 157 to our model, the observation that eCB-STDP is more robust to jitter than NMDAR-tLTP 158 could be a general property of the response of the signaling pathways to noisy paired 159 stimulations.

160

161 Differential effect of jitter on NMDAR- and eCB-mediated plasticity

Based on the model predictions, we next investigated experimentally the sensitivity to jitter of NMDAR-tLTP induced by 100 post-pre pairings (Fig. 3a), using stochastic Δt_{STDP} , distributed according to Figure 2c1. As aforementioned, NMDAR-tLTP can be induced with a $\sigma_{\Delta t}$ <3ms (Fig. 1c3). Increasing $\sigma_{\Delta t}$ beyond 3ms was sufficient to obliterate tLTP for 100 post-pre pairings. As exemplified (Fig. 3b and c1), 100 post-pre pairings with $\sigma_{\Delta t}$ =7.9ms and centered on $m_{\Delta t}$ =-18ms failed to induce plasticity. Overall, we did not observe tLTP expression for 168 $\sigma_{\Delta t}$ >3ms (92±2%, *p*=0.560, n=7; *p*<0.0001 when compared with tLTP observed without 169 jitter) (Fig. 3c2 and c3) in agreement with the model prediction (Fig. 2c2).

We next investigated the robustness of the eCB-tLTD to the variability of Δt_{STDP} . We observed potent tLTD even with large values of $\sigma_{\Delta t}$. As shown in Figure 3d1, 100 pre-post pairings with $\sigma_{\Delta t}$ =7.0ms centered on $m_{\Delta t}$ =+18ms induced a large tLTD. Overall, we observed that tLTD was still induced for 100 pre-post pairings with 3< $\sigma_{\Delta t}$ <10ms (70±6%, *p*<0.0001, n=14; *p*=0.1661 when compared with tLTD observed without jitter) (Fig. 3d2 and d3). The mean value of tLTD, observed for 3< $\sigma_{\Delta t}$ <10ms, was not different from the one observed in control, i.e. for 0< $\sigma_{\Delta t}$ <3ms (*p*=0.400).

Finally, we investigated the robustness of the eCB-tLTP to the variability of Δt_{STDP} and observed potent tLTP even with large values of jitter, i.e. up to $\sigma_{\Delta t} \sim 8\text{ms}$. An example of tLTP induced by 10 post-pre pairings with $\sigma_{\Delta t}$ =6.4ms centered on $m_{\Delta t}$ =-21ms is illustrated in Figure 3e1. In summary, tLTP could be induced even for $\sigma_{\Delta t}$ =8ms (160±12%, p=0.0021, n=9; p=0.0732 when compared with tLTP observed without jitter) (Fig. 3e2 and e3); the mean value of tLTP was not different from the one observed in control (p=0.1660).

We ensured that the $m_{\Delta t}$ absolute values were not different in control and jittered conditions for 100 post-pre pairings (18±2ms, n=13, vs 21±3ms, n=7, p=0.4711), 100 pre-post pairings (21±2ms, n=10, vs 21±2ms, n=14, p=0.8984), 10 post-pre pairings (17±2ms, n=15, vs 20±2ms, n=9, p=0.2297) or between these different STDP protocols (one-way ANOVA: p=0.4851) (Supplementary Fig. 1).

In conclusion, in agreement with our model prediction, whereas NMDAR-mediated tLTP appears fragile against the variability of Δt_{STDP} , eCB-plasticity (eCB-tLTD as well as eCBtLTP) exhibits a large robustness for the temporal imprecision of Δt_{STDP} . 191

192 Model-based analysis of the effects of jittering

193 The above experimental results provide us with a validation of our mathematical model. We 194 next used this validated model as a tool to investigate the molecular mechanisms behind these 195 differences of robustness.

196 We have shown in a previous study that eCB-tLTP was expressed when the amount of eCBs 197 produced was large enough that the fraction of activated CB_1R , y_{CB1R} , overcomes a threshold $\theta_{\rm LTP}^{\rm start \ 13}$. In the absence of added noise, $\sigma_{\Delta t} = 0$, our experimental data shows that as few as 5 198 pairings at 1Hz and Δt_{STDP} =-15ms are enough to trigger eCB-tLTP¹². Accordingly, our model 199 produces large amounts of eCBs in the first 5-10 pre-post pairings so that y_{CB1R} reaches θ_{LTP}^{start} 200 after as few as 5 pairings (Fig. 4a1). When $\sigma_{\Delta t} = 5$ ms jitter was added to $m_{\Delta t} = -15$ ms, 201 202 some of the pairings failed to deliver y_{CB1R} transients of maximal amplitude (Fig. 4a2). But for $\sigma_{\Delta t}$ <7-8ms, a number of y_{CB1R} transients still had sufficient amplitude to overcome $\theta_{LTP}^{\text{start}}$, 203 204 which was enough to trigger eCB-tLTP (Fig. 4a2). When $\sigma_{\Delta t}$ was large though (e.g. 10ms), 205 the y_{CB1R} transients failed to reach the LTP zone and eCB-tLTP was not expressed (Fig. 4a3).

In experiments without added jitter, eCB-tLTD progressively accumulates when N_{pairings} increases and starts to be significant for $N_{\text{pairings}} > 25^{12}$. In the model with $m_{\Delta t} = +20$ ms and $\sigma_{\Delta t} = 0$, the y_{CB1R} transients remain in the LTD area for the most part of the STDP pairings (Fig. 4a3). To account for the progressive accumulation of eCB-tLTP with N_{pairings} , each transient in the model contributes a small decrease of the synaptic weight. Even with large amounts of jitter (see e.g. $\sigma_{\Delta t} = 5$ or 10ms in Figure 4a2-a3) y_{CB1R} transients remain mostly inside LTD area so eCB-tLTD remains expressed even with large jitter.

213

214 The robustness of NMDAR-tLTP is predicted to be frequency-dependent

Experimentally, NMDAR-tLTP ($m_{\Delta t} = -15$ ms and $\sigma_{\Delta t} = 0$) was observed at $F_{\text{pairings}} = 1$ Hz 215 when $N_{\text{pairings}} > 50$, beyond which its amplitude did not depend much on N_{pairings}^{13} . To account 216 for this feature, the steady-state concentration of activated CaMKII, that sets W_{post} in the 217 218 model, is bistable: 45 min after the stimulation, CaMKII was either almost fully inactivated ("no plasticity" state) or almost fully activated ("LTP" state)^{13,38}. When the frequency of post-219 pre pairings (-30 $\leq \Delta t_{\text{STDP}} \leq 0$ ms) was large enough (i.e. $F_{\text{pairings}} \geq 1$ Hz), the IPI was smaller than 220 221 the decay time of the CaMKII activation transient triggered by each post-pre pairings (Fig. 222 4b1). As a result, in the absence of added noise ($\sigma_{\Delta t} = 0$), the CaMKII activation transients 223 progressively built up on top of each other. The accumulated CaMKII activation overcomes the threshold between the "no plasticity" and the "LTP" states only for $N_{\text{pairings}} > 50$ (Fig. 224 $(4b1)^{13}$. With jitter (e.g. $\sigma_{\Delta t}$ =5ms in Fig. 4b2 or 10ms in Fig. 4b3), many of the IPIs were 225 226 either too long or too short to trigger maximal amplitude transients of activated CaMKII. As a 227 result, CaMKII activation never reached the threshold, thus explaining the fragility of 228 NMDAR-tLTP with respect to jittering (at 1Hz).

229 A major indication from the above analysis is the importance of the IPI frequency for the 230 robustness of NMDAR-tLTP to jitter. Figure 5 compares the robustness to jitter for constant IPI with F_{pairings} =1Hz (Fig. 5a) and 1.05Hz (Fig. 5b). The size of the green area that locates 231 232 eCB-tLTD on Figure 5a2-b2 was not much altered by the increase of the pairing frequency. 233 Likewise, the size of the red area that locates eCB-tLTP in Figure 5a1-b1 did not vary much 234 when the frequency was increased. However, the size of the red area locating NMDAR-tLTP increased drastically with frequency (Fig. 5a2-b2). Therefore, our model predicts that 235 236 NMDAR-tLTP should become more robust to jitter at larger frequencies, whereas eCB-tLTP 237 and eCB-tLTD are not significantly changed.

238 Several aspects of glutamate signaling at the corticostriatal synapse are known to display complex frequency-dependence³⁹. Our model is calibrated with experimental data at 1Hz and 239 240 features none of the above frequency dependencies. Therefore, we cannot expect a precise 241 quantitative match between experiments and model when frequency is varied below or above 242 1Hz. However, the model still yields correct predictions of the main qualitative trends 243 observed in the experiments, since the effects of a small change of F_{pairings} in the model (1.00 244 to 1.05Hz) were similar to the effects of larger changes (1 to 3Hz) in the experiments¹³. 245 Therefore, in the experiments, we expect to observe a change of robustness pattern similar to 246 Figure 5b, around F_{pairings} =3Hz rather than 1.05Hz.

247

248 Increasing F_{pairings} and N_{pairings} stabilized NMDAR-tLTP against jitter

249 We next tested experimentally whether an increase from 1 to 3Hz of STDP pairings would 250 protect the NMDAR-tLTP against jitter, as predicted by the model. As shown in Figure 6a1, 251 100 post-pre pairings at 3Hz with $\sigma_{\Delta t}$ =9ms and centered on $m_{\Delta t}$ =-22ms induced tLTP. Overall, we observed tLTP for 100 post-pre pairings applied at 3Hz with $4 < \sigma_{\Delta t} < 9$ ms 252 253 (181±30%, p=0.0274, n=9) (Fig. 6a2 and a3); the mean value of these tLTP observed at 3Hz with $4 < \sigma_{\Delta t} < 9$ ms were not significantly different from the one observed with stimulation-254 related $\sigma_{\Delta t}$ <3ms at 1Hz (p=0.520). When we plotted the magnitude of plasticity against $\sigma_{\Delta t}$ 255 for 100 post-pre pairings at 3Hz, we observed that tLTP was still induced even for $\sigma_{\Delta t}$ =9ms 256 257 (Fig. 6a3). We observed different synaptic efficacy changes following 100 post-pre pairings at 1Hz with and without jitter and at 3Hz with jitter (ANOVA, p<0.0001). The NMDAR-tLTP 258 259 thus acquires certain robustness to the variability of Δt_{STDP} with increasing frequency of 260 pairings.

261 We next investigated whether a higher N_{pairings} would secure the expression of NMDAR-tLTP 262 even with $\sigma_{\Delta t}$ >3ms (Fig. 6b). When we doubled N_{pairings} we did not observe expression of significant tLTP. Indeed, 200 post-pre pairings (-30 $< m_{\Delta t} < 0$ ms with $3 < \sigma_{\Delta t} < 10$ ms) did not 263 induce plasticity (112±9%, p=0.236, n=6; Fig. 6b3) but when we increased N_{pairings} up to 250-264 265 300, tLTP was reliably observed with $\sigma_{\Delta t}$ >3ms. Figure 6b1 shows an example of tLTP 266 induced by 250 pairings with $\sigma_{\Delta t}$ =5ms and centered on -20ms. Overall, 250-300 post-pre 267 STDP pairings (-30 $< m_{\Delta t} < 0$ ms with $3 < \sigma_{\Delta t} < 10$ ms) induced tLTP (170±10%, p=0.0012, n=6; Fig. 6b2 and b3); the mean values of these tLTP observed for $N_{\text{pairings}}=250-300$ at 1Hz with 268 $4 < \sigma_{\Delta t} < 9$ ms were not different compared to tLTP induced with 100 post-pre pairings at 1Hz 269 with $\sigma_{\Delta t} < 3$ ms (p=0.1952). We observed different synaptic efficacy changes following 100, 270 271 200 and 250-300 post-pre pairings (ANOVA, p<0.0001).

272

273 Transitions from spike-timing- to frequency-dependent plasticity

274 We next tested with our mathematical model whether the robustness of STDP to jitter of the spike timing depends on the regularity of the spike train. In Figure 2e, we show an example of 275 276 the combination of stochastic spike timings (triangular distribution) with stochastic IPIs 277 (Poisson distributed) with a constant and relatively large refractory period τ_r (see *Methods*-278 Mathematical model, Protocol 4). However, when τ_r decreases, short IPIs are more 279 frequently sampled and the train of presynaptic stimulation becomes more irregular. 280 Therefore, through variations of the refractory period τ_r , the presynaptic stimulation can be 281 progressively switched from highly irregular to regular spike trains.

Figure 7 shows model predictions for the alteration of the robustness curves when τ_r varies. With high refractory period, *i.e.* when $\tau_r \approx F_{\text{pairings}}$, the IPIs are weakly noisy, so one recovers the robustness curves of Figure 2e, with eCB-STDP being more robust than

NMDAR-tLTP. However, when τ_r is 70% of F_{pairings} , the robustness of both eCB-tLTP and 285 NMDAR-tLTP increases and both exhibit similar robustness. With even smaller values of τ_r , 286 NMDAR-tLTP becomes more robust than eCB-tLTP. Moreover, eCB-tLTD ($F_{pairings}$ =1Hz, 287 $m_{\Delta t} = +20$ ms, $N_{\text{pairings}}=100$) progressively stops triggering LTD when τ_r decreases, even 288 inducing LTP for very low τ_r . Hence, when the presynaptic spike train is progressively 289 290 switched from regular to very irregular spike trains, two main changes occur: (i) the three 291 plasticity forms (NMDAR-tLTP, eCB-tLTD and eCB-tLTP) become increasingly robust to 292 jittering (at the limit of highly irregular spike trains, jittering hardly affects plasticity 293 amplitudes), and (ii) eCB-tLTD is progressively changed to LTP, so that highly irregular 294 paired stimulations with $F_{\text{pairings}}=1$ Hz only produce LTP regardless of the sign of the spike 295 timing.

296 **DISCUSSION**

297 Here, we have introduced jittered STDP protocols as noisy variants of STDP protocols where 298 the spike timing, *i.e.* the delay between a presynaptic and paired postsynaptic stimulation (or 299 vice-versa), is a random variable. Our results show that the magnitude of variability that 300 STDP can tolerate varies depending on the pairing frequency and on the STDP form, i.e. 301 NMDAR-tLTP, eCB-tLTP or eCB-tLTD. At 1Hz pairing frequency, eCB-STDP (eCB-tLTD 302 and eCB-tLTP) is much more robust than NMDAR-dependent tLTP. However, this 303 robustness depends on the number and frequency of pairings. Indeed, increasing the average 304 pairing frequency to 3Hz improved the robustness of NMDAR-tLTP. We observed a similar 305 improvement of the robustness of NMDAR-tLTP to jitter when the number of pairings 306 increased more than two folds.

307 These results have wide-ranging implications for STDP expression in *in vivo*-like firing. They 308 indicate that eCB-STDP could be more likely responsible for fast learning involving few trials 309 (eCB-tLTP) or for learning at lower activity frequency than NMDAR-STDP. NMDAR-STDP 310 could however take part in learning for larger frequencies or when the same spike timing 311 recurs a large number of times during sustained activity. It also implies that in synapses where NMDAR is the main coincidence detector for tLTD or tLTP^{3,5,6,40-42}, a high degree of 312 313 precision of the spike timing, and/or an increased number or frequency of pairings, are 314 required for the emergence of the NMDAR-mediated plasticity. This can be viewed as a 315 mechanism preventing the occurrence of spurious plasticity for noisy neural network activity. 316 At synapses in which NMDAR and endocannabinoids are both involved in STDP expression^{12,13,25–27,33,43} and in which eCB-LTP can occur, as it is the case in the striatum^{12,13}, 317 the hippocampus⁴⁴⁻⁴⁸ or the neocortex⁴⁹ the emergence of STDP would be possible even in 318 319 noisy conditions and could thus serve subsequently as a priming for the subsequent 320 expression of NMDAR-tLTP.

321 In the present study, we explored the robustness of STDP when noise affects not only the 322 spike timing but also the inter-pairing interval (IPI), i.e. the delay between two successive 323 pairings. In the absence of jitter of the spike timing, our model predicts that IPI irregularity 324 tends to consolidate tLTP at the expense of tLTD. This prediction that irregular IPIs 325 strengthen tLTP when the spike timing is constant confirms the result obtained independently and with different mathematical models^{7,19,22,24}. When IPIs are regular, adding jitter of the 326 327 spike timing in our model progressively suppresses STDP. However, with more irregular IPIs 328 our model predicts that the three STDP forms should exhibit larger robustness to noise. 329 Therefore our result suggests that jittering of the spike timing has larger impact on STDP 330 when the IPIs are regular, but that jittering has much less consequence with irregular IPIs. 331 Experimental testing of this model prediction would greatly improve our understanding of the induction and maintenance of STDP in in vivo-like firing. 332

Subthreshold postsynaptic depolarization is a key factor in the induction of plasticity⁵⁰. The 333 334 existence of plasticity induced by STDP-like protocols without postsynaptic spikes 335 (subthreshold-depolarization-dependent plasticity, SDDP) has been reported in the neocortex⁵¹, striatum⁵² and hippocampus⁵³. In the striatum, SDDP differs from STDP on two 336 aspects: (i) SDDP was induced in a larger temporal window (-100 $\leq \Delta t_{STDP} \leq +100$ ms) and (ii) 337 338 tLTP and tLTD were induced regardless of the spike timing. Therefore, the action potential 339 would not be necessary for plasticity induction, but determinant for the STDP polarity and the 340 width of the Δt_{STDP} . The effects of jitter remain to be determined on SDDP, *i.e.* subthreshold 341 events.

The polarity of plasticity depends greatly on the frequency of stimulus presentation. Indeed, prolonged firing activity at low and high frequency promote LTD and LTP, respectively, at the same synapses in hippocampus^{54–56}. However, alteration of a regular low-frequency stimulation (900 stimulations at 1Hz) with Poisson-distributed interstimulus intervals (as an

approximation of naturalistic patterns) prevents LTD expression in the neocortex⁵⁷. 346 347 Interestingly, this LTD induced with regular low-frequency stimulation has been reported to be NMDAR-mediated⁵⁸, and we show here that NMDAR-plasticity is fragile to jitter 348 especially for low firing rate (1Hz). It remains thus to investigate in a rate-based paradigm 349 whether Poisson stimulation would erase LTD induced by higher firing rate (3-5Hz⁵⁴). 350 351 Various forms of naturalistic patterns as an attempt to mimic in vivo activity have been 352 experimentally tested for plasticity expression: from single burst of dendritic spikes, which induced NMDAR-LTP in hippocampus⁵⁹ and NMDAR-LTD in neocortex⁶⁰, to noisy STDP-353 like protocols^{7,19,20,22} and burst-timing-dependent plasticity paradigms⁶¹. Based on data 354 355 harvested in the neocortex, a phenomenological computer modeling showed that for constant frequencies but random spiking, LTP wins over LTD with increasing frequency⁷. Activity 356 357 patterns recorded in vivo during specific behavioral tasks have been replayed ex vivo to test their ability to induce plasticity^{19,56,62}. When sequences of spike trains recorded *in vivo* from 358 359 visual cortex of the anaesthetized cat in response to natural scenes were replayed ex vivo in 360 layer 2/3 pyramidal cells of rat visual cortex, both LTP and LTD was observed 361 experimentally and modeling suggested the importance of suppressive interactions between spikes¹⁹. In the dentate gyrus, *in vivo* patterns recorded in granule cell during a task of delayed 362 nonmatch-to-sample were able in most of the cases to induce LTP at mossy fibers⁵⁶. 363 364 Similarly, when hippocampal place cells activity recorded in vivo were replayed ex vivo, NMDAR-LTP (also dependent on cholinergic tone) was observed⁶². Natural stimuli recruit 365 366 various neuronal circuits leading to presynaptic jitter, which can have important physiological 367 consequences. Indeed, in vivo visually driven presynaptic inputs induce presynaptic jitter leading to broader membrane potential events, and can consequently promote a spatial 368 369 reorganization of the responses of neurons to neighboring stimuli within a receptive field, as demonstrated in the rat visual $cortex^{22}$. 370

The debate about the nature of the neural coding, i.e. rate versus time $codes^{63-65}$, is prominent 371 372 when studying STDP. It remains to be tested with mathematical models and experiments 373 whether activity patterns of various vigilance states differently result in the emergence of 374 spike- or rate-coding plasticity. At most of the synapses both rate and time codes co-exist; for 375 example at CA1 synapses, stimuli induced LTD at 2-3Hz, no plasticity at 10Hz and LTP at 50 Hz demonstrating a rate code⁵⁴, and time codes are also supported as illustrated by the 376 expression of various forms of STDP at the same synapse¹¹. An answer to this debate could 377 378 be that both rate and time codes work together for plasticity expression, as recently shown in the somatosensory cortex⁶⁶, but their relative contribution *in vivo* would depend on the 379 380 ongoing activity (low or high), the cerebral structures, the noise level, the neuromodulatory 381 systems at play and the signaling pathways involved in plasticity expression (as shown in the 382 present study). Briefly, it appears that STDP can emerge form low frequency activity whereas rate codes would take over for higher frequency ranges^{7,19,20,22}. 383

384 A major conclusion from our present work is that the robustness of STDP to jitter depends on 385 the underlying signaling pathways. For instance, the robustness of NMDAR-tLTP in our 386 model is dependent on the amplitude of the activated CaMKII transients triggered by each 387 post-pre pairings or on alterations of the ratio between the decay time of these transients and 388 the IPIs. The robustness of STDP to jitter could be similarly controlled by quantitative 389 variations in the underlying pathways. Such variations are expected to occur between two 390 neuronal subtypes or brain regions but also as a result of the activation of a neuromodulatory 391 pathway. Therefore, our work suggests that the expression of STDP in *in vivo*-like firing 392 might appear or disappear as a result of modulations of its robustness to jitter, depending on 393 the properties of the incoming patterns (reflecting for example fast learning or heavy 394 training), but also on the brain region and, more transiently, on the activation of 395 neuromodulatory pathways.

396 **METHODS**

397 Animals and brain slice preparation

398 OFA (Oncins France Strain A) rats (Charles River, L'Arbresle, France) were used at postnatal 399 day 25-32 for brain slice electrophysiology. All experiments were performed in accordance 400 with the guidelines of the local animal welfare committee (Center for Interdisciplinary 401 Research in Biology Ethics Committee) and the EU (directive 2010/63/EU). The experimental 402 protocol (ref. 2017-02 N° 2016120110255176) was approved by the Ethics Committee in 403 Charge of Animal Experimentation (Paris Centre et Sud). Every precaution was taken to 404 minimize stress and the number of animals used in each series of experiments. Animals were 405 housed in standard 12 hours light/dark cycles and food and water were available ad libitum. 406 Horizontal brain slices containing the somatosensory cortical area and the corresponding corticostriatal projection field were prepared as previously described³⁶. Corticostriatal 407 408 connections (between somatosensory cortex layer 5 and the dorso-lateral striatum) are 409 preserved in the horizontal plane. Brain slices (330µm-thick) were prepared with a vibrating 410 blade microtome (VT1200S, Leica Microsystems, Nussloch, Germany). Brains were sliced in 411 an ice-cold cutting solution (125mM NaCl, 2.5mM KCl, 25mM glucose 25mM NaHCO₃, 412 1.25mM NaH₂PO₄, 2mM CaCl₂, 1mM MgCl₂, 1mM pyruvic acid) through which 95% O₂/5% CO₂ was bubbled. The slices were transferred to the same solution at 34°C for one hour and 413 414 then to room temperature.

415

416 Patch-clamp recordings

417 Patch-clamp recordings were performed as previously described^{12,18}. Briefly, for whole-cell 418 recordings in borosilicate glass pipettes of 5-7 M Ω resistance were filled with (in mM): 122 419 K-gluconate, 13 KCl, 10 HEPES, 10 phosphocreatine, 4 Mg-ATP, 0.3 Na-GTP, 0.3 EGTA

420 (adjusted to pH 7.35 with KOH). The composition of the extracellular solution was (mM): 421 125 NaCl, 2.5 KCl, 25 glucose, 25 NaHCO₃, 1.25 NaH₂PO₄, 2 CaCl₂, 1 MgCl₂, 10 µM 422 pyruvic acid bubbled with 95% O₂ and 5% CO₂. Signals were amplified using EPC10-2 423 amplifiers (HEKA Elektronik, Lambrecht, Germany). All recordings were performed at 34°C 424 (Bath-controller V, Luigs&Neumann, Ratingen, Germany) and slices were continuously 425 superfused with extracellular solution, at a rate of 2 ml/min. Slices were visualized under 426 microscope (BX51WI Olympus, Rungis, France), with a 4x/0.13 objective for the placement 427 of the stimulating electrode and a 40x/0.80 water-immersion objective for the localization of 428 cells for whole-cell recordings. Current- and voltage-clamp recordings were filtered at 5 kHz 429 and sampled at 10 kHz, with the Patchmaster v2x32 program (HEKA Elektronik).

430

431 Spike timing-dependent plasticity protocols: regular and jittered patterns

432 Electrical stimulations were performed with a concentric bipolar electrode (Phymep, Paris, 433 France) placed in layer 5 of the somatosensory cortex. Electrical stimulation was monophasic, 434 at constant current (ISO-Flex stimulator, AMPI, Jerusalem, Israel). Cortical stimulations 435 evoked glutamatergic excitatory postsynaptic currents (EPSCs) (inhibited by CNQX 10µM 436 and D-AP5 50µM, n=6) and not significantly affected by GABAergic events; Indeed, 437 blocking GABA_ARs with picrotoxin (50µM) did not significantly affect EPSC amplitude at 438 corticostriatal synapses (123 \pm 31pA before and 118 \pm 26pA after picrotoxin, p=0.500, n=5). 439 MSNs were kept at their resting membrane potential (-74.6 ± 0.6 mV, n=90) and repetitive 440 control stimuli were applied at 0.1Hz. Currents were adjusted to evoke 50-200 pA EPSCs. 441 STDP protocols consisted of pairings of pre- and postsynaptic stimulations (at 1Hz) separated 442 by a specific time interval (Δt_{STDP}). The paired stimulations were applied at 1Hz throughout 443 the study except in Figure 6a1 and 6a2 in which 3Hz stimulations were tested. Presynaptic 444 stimulations corresponded to cortical stimulations and the postsynaptic stimulation of an

action potential evoked by a depolarizing current step (30ms duration) in MSNs. Δt_{STDP} <0ms 445 446 for post-before-pre pairings, and Δt_{STDP} >0ms for pre-before-post pairings. Neuronal 447 recordings on neurons were made over a period of 10 minutes at baseline, and for at least 60 448 minutes after the STDP protocols. EPSC baseline stability was assessed by comparing the 449 average of the EPSC magnitude of the two first and the two last minutes of baseline; if the variation of amplitude exceeded 10%, the experiment was discarded⁴. Recordings were 450 451 stopped if the injected current was larger than 50 or 100-150 pA during the baseline and after 452 pairings, respectively. We individually measured and averaged the amplitude of 60 successive 453 EPSCs from both baseline and 45-55 minutes after STDP protocol, in which the latter was 454 normalized by the former to calculate long-term synaptic efficacy changes. Neuron recordings 455 were made in voltage-clamp mode during baseline and for the 60 minutes of recording after 456 the STDP protocol, and in current-clamp mode during STDP protocol. Experiments were 457 excluded if the input resistance (Ri), measured every 10 sec all along the experiment, varied 458 by more than 20% during the very same time period where the EPSC amplitude were 459 measured for plasticity assessment: 10 minutes of baseline and the 10 last minutes of the 460 recording (generally 45-55 minutes after pairings). After recording of 10 min control baseline, 461 drugs were applied in the bath. A new baseline with drugs was recorded after a time lapse of 462 10 min (to allow the drug to be fully perfused) for 10 min before the STDP protocol. Drugs 463 were present until the end of the recording. All chemicals were purchased from Tocris 464 (Ellisville, MO, USA), except for picrotoxin (Sigma). DL-2-amino-5-phosphono-pentanoic 465 acid (D-AP5; 50µM) and 6-cvano-7-nitroquinoxaline-2,3-dione (CNOX; 10µM) were 466 dissolved directly in the extracellular solution and bath applied. N-(piperidin-1-yl)-5-(4-467 iodophenyl)-1-(2,4-dichlorophenyl)-4-methyl-1H-pyrazole-3-carboxamide (AM251; 3µM) 468 and picrotoxin (50µM) were dissolved in ethanol and added to the external solution, such that 469 the final concentration of ethanol was 0.01-0.1%.

470 For the jittered Δt_{STDP} patterns, we used the following algorithm (see Protocol 2, below). We note t_{pre_i} and t_{post_i} the times of the presynaptic and postsynaptic stimulations in the paired 471 472 stimulation number *i*, respectively. To set them, we first fixed the times of each presynaptic stimulation using the pairing frequency, i.e. $t_{\text{pre}_i} = \frac{i}{F_{\text{pairings}}}$. The postsynaptic times were 473 then chosen randomly according to $t_{\text{post}_i} = t_{\text{pre}_i} + m_{\Delta t} + \xi_{\Delta t,i}$ where $m_{\Delta t}$ is the average 474 spike timing and $\xi_{\Delta t}$ is a random variable with mean 0 and variance $\sigma_{\Delta t}^2$. We defined the 475 476 Inter-Pairing-Interval (IPI) as the time interval between the presynaptic stimulation times of two consecutive paired stimulations $(IPI_i = t_{pre_{i+1}} - t_{pre_i})$, whereas the spike timing Δt is 477 478 the time interval between the postsynaptic and the presynaptic stimulation times within a given paired stimulation ($\Delta t_{\text{STDP}_i} = t_{\text{post}_i} - t_{\text{pre}_i}$). In a standard STDP protocol both the 479 spike timing and the IPI are constant. The above algorithm yields constant IPIs and stochastic 480 481 spike timings.

482

483 Electrophysiological data analysis

484 Off-line analysis was performed with Fitmaster (Heka Elektronik) and Igor-Pro 6.0.3 485 (Wavemetrics). Statistical analysis was performed with Prism 5.02 software. In all cases "n" 486 refers to an experiment on a single cell from a single slice. All results are expressed as mean \pm 487 SEM in the text and as mean \pm SD in the figures. Statistical significance was assessed in 488 unpaired *t* tests, one way ANOVA, or in one-sample *t* tests, as appropriate, using the indicated 489 significance threshold (*p*).

490

492 **Jittered STDP protocols.** In the model, we define the spike timing as $\Delta t_{\text{STDP}_i} = t_{\text{post}_i} - t_{\text{pre}_i} + \delta$ where δ accounts for the time elapsed between the onset of the postsynaptic step 494 current and the action potential it triggers (~3ms in MSNs). In agreement with the experiment 495 protocol above, we introduced stochasticity of the spike timing by adding a random jitter $\xi_{\Delta t}$ 496 to the spike timing: $\Delta t_{\text{STDP}} = m_{\Delta t} + \xi_{\Delta t}$, were $m_{\Delta t}$ is the average spike timing and $\xi_{\Delta t}$ is a 497 random variable whose distribution is given by the STDP protocol. Here, we explored five 498 STDP protocols:

499 1) *Protocol 0* consisted of pairings with no jittering, i.e. $IPI_i = \frac{1}{F_{pairings}}$ and $\xi_{\Delta t} = 0$.

500 2) *Protocol 1* consisted of deterministic IPIs and uniformly distributed spike timings (Fig. 501 2a1), i.e. $IPI_i = \frac{1}{F_{pairings}}$, $\forall i$, and the probability distribution function of the jitter is given by

502
$$P(\xi_{\Delta t}) = \begin{cases} \frac{1}{2\sigma_{\Delta t}} & \text{if } |\xi_{\Delta t}| < \sigma_{\Delta t}, \\ 0 & \text{otherwise} \end{cases}$$
(1)

503 In eq. (1), $\sigma_{\Delta t}$ defines the maximal jitter amplitude *i.e.* the level of noise in the protocol. To 504 simulate *Protocol 1*, we first set each presynaptic stimulation time using $t_{\text{pre}_i} = \frac{i}{F_{\text{pairings}}}$. 505 Postsynaptic times were then fixed by $t_{\text{post}_i} = t_{\text{pre}_i} + m_{\Delta t} + \xi_{\Delta t,i} - \delta$.

506 3) *Protocol 2* (Fig. 2c1) shared the same definition as Protocol 1, except that the jitter 507 followed a normal distribution with zero mean and variance $\sigma_{\Delta t}^2$: $P(\xi_{\Delta t}) = \mathcal{N}(0, \sigma_{\Delta t}^2)$.

508 4) For *Protocol 3* (Fig. 2d1), we used a triangular distribution of the jitter by adding 509 uniformly distributed jitter to both presynaptic and postsynaptic times. We set t_{pre_i} =

510
$$\frac{i}{F_{\text{pairings}}} + \zeta_{\text{pre},i}$$
 and $t_{\text{post}_i} = \frac{i}{F_{\text{pairings}}} + m_{\Delta t} + \zeta_{\text{post},i} - \delta$, where ζ_{pre} and ζ_{post} are i.i.d.

random variables with the uniform distribution of eq. (1). The resulting spike timing is $\Delta t_{\text{STDP}} = m_{\Delta t} + \xi_{\Delta t}$ where $\xi_{\Delta t} = \zeta_{\text{post}} - \zeta_{\text{pre}}$ has a triangular distribution. Note that in this case, one still has $\text{IPI}_i \approx \frac{1}{F_{\text{pairings}}} \forall i$ as long as $2|m_{\Delta t}|F_{\text{pairings}} \ll 1$, which can safely be assumed here since we used $|m_{\Delta t}| < 50$ ms and $F_{\text{pairings}} < 2$ Hz.

515 5) Protocol 4 includes both a stochastic spike timing and stochastic IPIs (Fig. 2e1). We first sampled N_{pairings} IPIs according to an exponential distribution with rate λ and refractory 516 period τ_r : $P(IPI) = \Theta(IPI - \tau_r)\lambda \exp(-\lambda(IPI - \tau_r))$, where $\Theta(x)$ is the Heaviside function 517 $\Theta(x) = 1$ if $x \ge 0$; 0 otherwise. The spike trains defined with protocol 4 are thus Poisson 518 519 process. We used these IPIs to fix the stimulation times, adding a triangularly distributed jitter as in Protocol 3 above: $t_{\text{pre}_i} = \sum_{j < i} \text{IPI}_j + \zeta_{\text{pre},i}$ and $t_{\text{post}_i} = \sum_{j < i} \text{IPI}_j + m_{\Delta t} + \zeta_{\text{post},i} - \delta$. 520 To keep the average stimulation frequency at $1/F_{\text{pairings}}$, we constrained the value of the rate 521 λ as $\lambda = (1/F_{\text{pairings}} - \tau_r)^{-1}$. 522

523

524 **Stimulations**. A detailed account of our mathematical model can be found in S1 Text. We 525 modeled glutamate concentration in the synaptic cleft, G(t), as a train of exponentially-526 decaying impulses triggered by presynaptic stimuli at time t_{pre_i} . We modeled the electrical 527 response to these stimulations in a postsynaptic element considered as a single isopotential 528 compartment with AMPAR, NMDAR, VSCC and TRPV1 conductances:

529
$$C_{\rm m} dV/dt = -g_L(V - V_{\rm L}) - I_{\rm AMPAR}(V, G(t)) - I_{\rm NMDAR}(V, G(t)) - I_{\rm VSCC}(V) - I_{\rm TRPV1}(V, AEA) - I_{\rm action}(t)$$
(2)

530 where *V* is membrane potential; g_L and V_L are leak conductance and reversal potential 531 respectively; I_{AMPAR} , I_{NMDAR} , I_{VSCC} and I_{TRPV1} are currents through AMPAR, NMDAR, L-532 type VSCC (v1.3) and TRPV1 respectively. I_{action} is the action current resulting from the 533 postsynaptic stimulation (backpropagating action potential on top of a 30ms depolarization).

534 Details about the analytical expressions of these currents are given in S1 Text.

535 **Biochemical signaling**. We modeled the kinetics of the biochemical pathways activated by 536 the above electrical stimulations using the model of ref[13]. We give below a quick overview 537 of this model and refer to S1 Text for the details. Free cytosolic calcium is one of the main 538 signaling actors in the model. To model its dynamics, we assumed calcium can be transferred from/to two main sources: (i) extracellular calcium, via the plasma membrane channels of eq. 539 540 (3) above and (ii) the endoplasmic reticulum (ER), via the IP3-dependent calcium-induced 541 calcium release system. These calcium transients activated a network of biochemical pathways that collectively set the synaptic weight. Hence, in this model, the synaptic weight 542 is entirely fixed by the underlying biochemical signaling network¹³. More precisely, we 543 544 assumed that the total synaptic weight is the product of a pre- and a postsynaptic contribution $W_{\text{total}} = W_{\text{pre}}W_{\text{post}}$. Postsynaptic plasticity was based on the activation by calcium of 545 calmodulin and CaMKII^{67,68}. We assumed that postsynaptic plasticity is directly proportional 546 547 to the calcium-dependent activation of CaMKII:

548
$$W_{\text{post}} = 1 + 3.5 \frac{CaMKII^*}{CaMKII_{\text{max}}^*}$$
 (5)

where *CaMKII** and *CaMKII*^{*}_{max} are the current concentration of activated (phosphorylated)
CaMKII and its maximum value.

551 Our model also accounts for the biochemical pathways leading to the production of the 552 endocannabinoids 2-arachidonoylglycerol (2-AG) and AEA, and their subsequent activation 553 of CB₁R (see S1 Text for further details). In our model, CB₁R activation (y_{CB1R}) controls the 554 presynaptic weight W_{pre} according to the following rule:

555
$$\Omega(y_{CB1R}) = \begin{cases} 1 - A_{LTD} & \text{if } \Theta_{LTD}^{\text{start}} < y_{CB1R} < \Theta_{LTD}^{\text{stop}} \\ 1 + A_{LTP} & \text{if } \Theta_{LTP}^{\text{start}} < y_{CB1R} \\ 1 & \text{otherwise} \end{cases}$$
(7)

556 and
$$dW_{\rm pre}/dt = \left(\Omega(y_{\rm CB1R}) - W_{\rm pre}\right)/\tau_{W_{\rm pre}}$$
 (8)

557 Details about model implementation and numerics are given in the supplementary558 information.

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564

565 **Author contribution**:

L.V. and H.B. designed the study; Y.C. and A.M. performed electrophysiological experiments
and analysis; H.B. carried out the conception and the design of the mathematical model; H.B.
and I.P. performed the acquisition and analysis of data from the mathematical model; H.B.
and L.V. wrote the manuscript and all authors have edited and corrected the manuscript; L.V.
supervised the whole study.

571

572 **Competing financial interest:**

573 The authors declare no competing interests.

574

576 Figure 1: NMDAR- and eCB-mediated STDP induced with fixed Δt_{STDP}

577 (a) STDP protocol: a single postsynaptic spike was paired with a single cortical stimulation, 578 10 or 100 times at 1Hz. (b) Example of 5 successive pairings (#4-8, from c1) exhibiting 579 relatively fixed Δt_{STDP} (#4-8: -8.0, -7.8, -7.4, -8.8 and -8.0ms). Plotting the successive Δt_{STDP} 580 illustrates their low variance $\sigma_{\Delta t}$ (2.6ms). (c) Corticostriatal NMDAR-mediated tLTP induced 581 by 100 post-pre pairings with fixed Δt_{STDP} . (c1) Example of tLTP induced by 100 post-pre pairings ($m_{\Delta t}$ =-9ms, $\sigma_{\Delta t}$ =2.6ms). The mean baseline EPSC amplitude, 95±3pA, increased by 582 583 77% to 168±6pA 45 minutes after pairings. Upper panel, time course of Ri (baseline: 584 $172\pm 2M\Omega$ and 35-45 min after pairings: $170\pm 3M\Omega$; change of 1%) and holding current (I_{hold}). 585 Inset: distribution of the 100 Δt_{STDP} . (c2) Averaged time-courses of tLTP induced by 100 586 post-pre pairings; this tLTP was NMDAR-mediated because prevented by D-AP5 (50µM). (c3) Relationship between the STDP magnitude and the STDP jitter for each of the recorded 587 588 neurons. (d) Corticostriatal eCB-mediated t-LTD induced by 100 pre-post pairings with fixed 589 $\Delta t_{\text{STDP.}}$ (d1) Example of tLTD induced by 100 pre-post pairings ($m_{\Delta t}$ =+21ms, $\sigma_{\Delta t}$ =2.1ms); 590 the mean baseline EPSC amplitude, 206±4pA, decreased by 25%, to 154±8pA, 45 minutes 591 after pairings. Upper panel, time course of Ri (baseline: $195\pm1M\Omega$ and 45-55 min after 592 pairings: 196±2MΩ; change <1%) and I_{hold}. Inset: distribution of the 100 Δt_{STDP} . (d2) 593 Averaged time-courses of tLTD induced by 100 pre-post pairings; this tLTD was CB₁R-594 mediated because prevented by AM251 (3µM). (d3) Relationship between the STDP 595 magnitude and the jitter. (e) Corticostriatal eCB-mediated tLTP induced by 10 post-pre 596 pairings with fixed Δt_{STDP} . (e1) Example of tLTP induced by 10 post-pre pairings ($m_{\Delta t}$ =-11ms, $\sigma_{\Delta t}$ =3.1ms); the mean baseline EPSC amplitude, 147±3pA, increased by 43% to 597 598 209±9pA 45 minutes after pairings. Upper panel, time course of Ri (baseline: 227±2MQ and 599 45-55 min after pairings: $239\pm 2M\Omega$; change of 5%) and I_{hold}. Inset: distribution of the 10

600 $\Delta t_{\text{STDP.}}$ (e2) Averaged time-courses of tLTP induced by 10 post-pre pairings; this tLTP was 601 CB₁R-mediated because prevented by AM251. (e3) Relationship between the STDP 602 magnitude and the jitter. (c1-d1-e1) Insets: average of 12 successive EPSC amplitude at 603 baseline (black traces) and at 40-50 min after pairings (grey traces). (f) Predictions of the 604 mathematical model in the absence of added jitter. The model expresses the biochemical 605 pathways schematized in (f1). (f2) Changes of W_{total} with various Δt_{STDP} and N_{pairings} . (f3) 606 Removing NMDAR/CaMKII signaling suppresses the tLTP observed for $\Delta t_{\text{STDP}} > 0$ and 607 $N_{\text{pairings}} > 50.$ (f4) Removing CB₁R suppresses the tLTP observed for $\Delta t_{\text{STDP}} > 0$ and $N_{\text{pairings}} < 20$ 608 and the tLTD observed for $\Delta t_{\text{STDP}} < 0$.

609

610 Figure 2: The mathematical model predicts that NMDAR-tLTP is less robust to jitter

611

than eCB-tLTP and eCB-tLTD.

612 (a) Using a jittered STDP protocol combining uniformly-distributed jitter of the spike timing 613 with deterministic IPIs at 1Hz (a1), the amplitudes of NMDAR-tLTP, eCB-tLTD and eCB-614 tLTP decrease with increasing jitter amplitudes $\sigma_{\Delta t}$ (a2). eCB-tLTP (red full line) was triggered with 10 pairings of average spike timing $m_{\Delta t} = -15$ ms and eCB-tLTD (blue full 615 line) with 100 pairings of $m_{\Delta t} = +20$ ms. NMDAR-tLTP (red dashed line) corresponded to 616 100 pairings of $m_{\Delta t} = -15$ ms. eCB-tLTD and eCB-tLTP are weakly affected by the jitter. 617 618 NMDAR-tLTP is less robust. Full and dashed lines show averages and light swaths show ±1sem. (b) Evolution of NMDAR-tLTP, eCB-tLTD and eCB-tLTP domains for $\sigma_{\Delta t} = 0$ (b1), 619 5 (b2) and 10ms (b3) for the STDP protocol of (a). Note the persistence of the eCB-tLTD 620 621 domain (blue box) while eCB-tLTP (red box) and NMDAR-tLTP domain (red dashed box) 622 disappeared for 5 and 10ms, respectively. (c-e) Similar results are obtained with other jitter distributions, including Gaussian-distributed jitter (c) and triangularly-distributed jitter (d). 623 624 We also considered a combination of triangularly-distributed jitter with random IPIs distributed according to an exponential distribution with refractory period τ_r =0.95 s (see *Methods*) in (e). The plots of the amplitude of NMDAR-tLTP, eCB-tLTD and eCB-tLTP for increasing jitter amplitude $\sigma_{\Delta t}$ (c2, d2, e2) show similar profiles and are similar to panel (a2), indicating that the model predictions are not qualitatively modified by the probability distribution function of the jitter.

630

631 Figure 3: Differential effect of jitter on NMDAR- and eCB-mediated plasticity.

632 (a) STDP protocol; Δt_{STDP} indicates the time between pre- and postsynaptic stimulations. 633 $\Delta t_{\text{STDP}} < 0$ and $\Delta t_{\text{STDP}} > 0$ refer to post-pre and pre-post pairings, respectively. (b) Example of 5 634 successive pairings (#5-9, from c1) exhibiting jittered values of Δt_{STDP} (#5-9: -10.3, -15.2, -635 30.5, -20.7 and -17.3ms); these pairings are indicated with the grey frame within the 100 636 pairings displaying jittered Δt_{STDP} . The distribution of the successive Δt_{STDP} illustrate their 637 standard deviation $\sigma_{\Delta t}$ =7.9ms. (c) NMDAR-tLTP is prevented with $\sigma_{\Delta t}$ >3ms. (c1) Example of the absence of plasticity after 100 post-pre pairings with $m_{\Delta t}$ =-18ms and $\sigma_{\Delta t}$ =7.9ms; the 638 639 mean baseline EPSC amplitude, 194±3pA, did not show significant change 45 minutes after 640 pairing, 186±4pA. Upper panel, time course of Ri (baseline: 198±2MΩ and 45-55 min after pairings: 194±2MΩ; change of 2%) and holding current (I_{hold}). Inset: distribution of the 100 641 642 $\Delta t_{\text{STDP.}}$ (c2) Averaged time-courses of the absence of plasticity after 100 post-pre pairings 643 with $3 < \sigma_{\Delta t} < 10$ ms. (c3) Relationship between the STDP magnitude and the jitter. (d) eCB-644 mediated t-LTD is not affected by $3 < \sigma_{\Delta t} < 10$ ms. (d1) Example of tLTD induced by 100 prepost pairings with $m_{\Delta t}$ =+18ms and $\sigma_{\Delta t}$ =7.1ms; the mean baseline EPSC amplitude, 645 646 160±3pA, decreased by 50%, to 81±5pA, 45 minutes after pairings. Upper panel, time course 647 of Ri (baseline: $112\pm1M\Omega$ and 35-45 min after pairings: $118\pm1M\Omega$; change of 5%) and I_{hold}. 648 Inset: distribution of the 100 Δt_{STDP} . (d2) Averaged time-courses of tLTD induced by 100 pre-649 post pairings with $3 < \sigma_{\Delta t} < 10$ ms. (d3) Relationship between the STDP magnitude and the jitter. (e) eCB-mediated tLTP is not affected by $3 < \sigma_{\Delta t} < 10$ ms. (e1) Example of tLTP induced by 10 post-pre pairings with $m_{\Delta t}$ =-21ms and $\sigma_{\Delta t}$ =6.4ms; the mean baseline EPSC amplitude, 113±2 pA, increased by 115% to 243±6pA 45 minutes after pairings. Upper panel, time course of Ri (baseline: 185±1MΩ and 45-55 min after pairings: 197±2MΩ; change of 6%) and I_{hold}. Inset: distribution of the 10 Δt_{STDP} . (e2) Averaged time-courses of tLTP induced by 10 post-pre pairings with $3 < \sigma_{\Delta t} < 10$ ms. (e3) Relationship between the STDP magnitude and the jitter.

Insets: average of 12 successive EPSC amplitude at baseline (black trace) and at 40-50 minafter STDP pairings (grey trace). Error bars represent the SEM.

659

Figure 4: Model-based analysis of the effects of jittering.

661 Model-predicted time-courses of (a) CB₁R activation y_{CB1R} and (b) the concentration of active 662 CaMKII during a protocol with jitter amplitude $\sigma_{\Delta t} = 0$ (a1-b1), 5 (a2-b2) or 10 (a3-b3) ms. 663 The STDP protocol was the same as Figure 2a1 *i.e.* uniformly-distributed jitter of the spike 664 timing and deterministic IPIs at 1Hz. eCB-tLTP (a, red full line) and NMDAR-tLTP (b, red dashed line) were triggered with average spike timing $m_{\Delta t} = -15$ ms and eCB-tLTD (**a**, blue 665 full line) with $m_{\Delta t} = +20$ ms. In (a) the shaded boxes locate the areas where y_{CB1R} triggers 666 eCB-tLTP or eCB-tLTD. Those areas are defined by the thresholds $\theta_{LTP}^{\text{start}}$ and $(\theta_{LTD}^{\text{start}}, \theta_{LTD}^{\text{stop}})$, 667 respectively (see *Methods*). For $m_{\Delta t} = -15$ ms (red), increasing jitter amplitudes 668 669 progressively hinders the build up of the y_{CB1R} transients. y_{CB1R} still reaches the LTP area for 5-25 pairings with $\sigma_{\Delta t} = 0$ (a1) and 5ms (a2) thus triggering eCB-tLTP but fails to do so for 670 671 $\sigma_{\Delta t} = 10 \text{ms}$ (a3). Conversely, eCB-tLTD is robust to jitter because y_{CB1R} remains in the LTD region for most of the pairings with $m_{\Delta t} = +20$ ms (blue), for all tested jitter amplitudes. In 672 (b), with $m_{\Delta t} = -15$ ms (red) the switch from the low activation to the high activation state 673

of CaMKII is obtained in the absence of jittering $\sigma_{\Delta t} = 0 \text{ms} (\mathbf{b1})$. The progressive build up of activated CaMKII is efficiently suppressed as soon as $\sigma_{\Delta t} = 5 \text{ms} (\mathbf{b2})$ thus effectively suppressing NMDAR-tLTP (**b3**).

677

Figure 5: Model-based analysis of the change of robustness with frequency.

679 (a) Summary of the robustness of the model plasticity at 1Hz. The jittered STDP protocol 680 combined uniformly-distributed jitter of the spike timing and constant IPIs at 1Hz. Two-681 dimensional maps show alterations of the plasticity obtained with 10 (a1) or 100 (a2) 682 pairings, when the average spike timing $m_{\Delta t}$ and the jitter amplitude $\sigma_{\Delta t}$ are changed. The 683 locations of the plasticity blotches are indicated with horizontal lines: full red for eCB-tLTP, 684 full blue for eCB-tLTD and dashed red for NMDAR-tLTP. The length of each of these 685 blotches along the y-axis reflects their robustness to jitter. (a3) A cross-section along the 686 dashed and full horizontal lines of (a1-a2) illustrates the robustness of NMDAR-tLTP, eCB-687 tLTD and eCB-tLTP. (b) Robustness of the model with larger frequency (1.05Hz) and same 688 jittered STDP protocol as in (a). Note the distortion between the pairing frequency in the 689 model and in the experiments. The behavior of the model for $F_{\text{pairings}}=1.05$ Hz corresponds to experimental results with $F_{\text{pairings}} \approx 3 \text{Hz}^{13}$. The two-dimensional maps for 10 (**b1**) or 100 (**b2**) 690 691 pairings as well as the cross-sections (b3) predict that for frequencies larger than 1Hz, the 692 robustness changes, with NMDAR-tLTP becoming more robust.

693

694 Figure 6: Increasing frequency and N_{pairings} stabilized NMDAR-tLTP against jitter.

(a) 100 post-pre pairings at 3Hz induced a tLTP, which is not affected by $3 < \sigma_{\Delta t} < 10$ ms. (a1) Example of tLTP induced by 100 post-pre pairings at 3Hz with $m_{\Delta t}$ =-22ms and $\sigma_{\Delta t}$ =9ms; the mean baseline EPSC amplitude, 194±4pA, increased by 212%, to 606±7pA, 45 min after pairings. Upper panel, time course of Ri (baseline: 123±1MΩ and 45-55 min after pairings:

699 138±1MΩ; change of 12%) and I_{hold} for this cell. Inset: distribution of Δt_{STDP} for the 100 700 pairings. (a2) Averaged time-courses of tLTP induced by 100 post-pre pairings at 3Hz with 701 $3 < \sigma_{\Delta t} < 10$ ms. (a3) Relationship between the STDP magnitude and the jitter for each of the recorded neurons after 100 pairings at 1Hz (n=7; blue circles) or at 3Hz (n=10; red circles). 702 703 (b) 250-300 post-pre pairings induced tLTP, which is not affected by $3 < \sigma_{\Delta t} < 10$ ms. (b1) Example of tLTP induced by 250 post-pre pairings with $m_{\Delta t}$ =-20ms and $\sigma_{\Delta t}$ =5ms; the mean 704 705 baseline EPSC amplitude, 126±3pA, increased by 57% to 198±4pA 45 minutes after pairings. 706 Upper panel, time course of Ri (baseline: $65\pm1M\Omega$ and 45-55 min after pairings: $65\pm2M\Omega$; 707 change <1%) and I_{hold}. Inset: distribution of Δt_{STDP} for the 250 pairings. (b2) Averaged time-708 courses of tLTP induced by 250-300 post-pre pairings with $3 < \sigma_{\Delta t} < 10$ ms. (b3) Relationship 709 between the STDP magnitude and the jitter for each of the recorded neurons after 100 (n=7; 710 blue circles), 200 (n=6; black circles) or 250-300 (n=7; red circles) post-pre pairings.

Insets: average of 12 successive EPSC amplitude at baseline (black trace) and at 40-50 min
after STDP pairings (grey trace). Error bars represent the SEM.

713

Figure 7: The robustness of plasticity depends on the regularity of the presynaptic spike train.

716 The figure shows the robustness of the model plasticity (eCB-tLTP with red, eCB-tLTD with 717 blue and NMDAR-tLTP with dashed red) with the jittered STDP protocol of figure 2e1, *i.e.* 718 exponentially-distributed IPIs with refractory period τ_r and triangularly-distributed jitter $\xi_{\Delta t}$. From right to left, the refractory period τ_r is gradually decreased. When $\tau_r \rightarrow 1/F_{\text{pairings}}$ 719 where $F_{\text{pairings}} = 1$ Hz is the average pairings frequency, one gets regular presynaptic spike 720 trains, similar to classical STDP protocols. In this case (see e.g. $\tau_r = 0.95/F_{\text{pairings}}$), the 721 722 robustness pattern is similar to the previous observations at 1Hz: eCB-STDP is robust, 723 whereas NMDAR-tLTP is fragile. However, when $\tau_r \rightarrow 0$, the presynaptic spike train

- becomes highly irregular. In this case, the robustness patterns changes totally, with NMDAR-
- 725 LTP becoming the most robust, and eCB-tLTD turning into a LTP. Hence, irregular
- 726 presynaptic spike trains are predicted to produce robust LTP at the expense of LTD.

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b3











 $\sigma_{\Delta t} = 5 \text{ ms}$

 \diamond









 $\sigma_{\Delta t}$ = 10 ms

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Robustness of STDP to spike-timing jitter

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SUPPORTING INFORMATION

S1 Text - Description of the mathematical model

We describe here the mathematical model used to predict the outcome of STDP stimulations with spike timing jitter. We give here the information necessary to implement the model. Further detail and validation can be found in Cui et al (2016).

Stimulations. A STDP protocol consists in a series of N_{pairings} post- and presynaptic pairs of stimulations. We note t_{pre_i} the time at which the *i*th presynaptic stimulation is delivered and t_{post_i} the onset time of the *i*th postsynaptic step current stimulation, where $i = 1, ..., N_{\text{pairings}}$. We modeled glutamate concentration in the synaptic cleft, G(t), as a train of exponentially-decaying impulses triggered by presynaptic stimuli at time t_{pre_i} :

$$G(t) = G_{\max} \sum_{i} \exp\left(-\frac{t - t_{\text{pre}_i}}{\tau_{\text{G}}}\right) \Theta\left(t - t_{\text{pre}_i}\right)$$
(SI1)

where G_{max} is the peak glutamate concentrations and τ_G is the glutamate clearance rate. On the postsynaptic side, action currents resulting from postsynaptic stimulations were modeled according to

$$I_{\text{action}}(t) = -\text{DC}_{\max}\sum_{i} R(t, t_{\text{post}_{i}}, \text{DC}_{\text{dur}}) - \text{AP}_{\max}\sum_{i} \Theta(t - \delta - t_{\text{post}_{i}}) \exp\left(\frac{-t + \delta + t_{\text{post}_{i}}}{\tau_{\text{bAP}}}\right) (SI2)$$

where the rectangle function $R(t, a, l) = \Theta(t - a) - \Theta(t - a - l)$, DC_{max} and DC_{dur} are the amplitude and duration of step-current injected by the patch pipette in the postsynaptic soma, AP_{max} is the amplitude of the action current producing the bAP; and τ_{bAP} is the time scale for bAP decay.

We modeled the electrical response to these stimulations in a postsynaptic element considered as a single isopotential compartment with AMPAR, NMDAR, VSCC and TRPV1 conductances

$$C_{\rm m} \frac{dV}{dt} = -g_L (V - V_{\rm L}) - I_{\rm AMPAR} (V, G(t)) - I_{\rm NMDAR} (V, G(t)) -I_{\rm VSCC} (V) - I_{\rm TRPV1} (V, AEA) - I_{\rm action} (t)$$
(SI3)

where V is membrane potential; g_L and V_L are leak conductance and reversal potential respectively; I_{AMPAR} , I_{NMDAR} , I_{VSCC} and I_{TRPV1} are currents through AMPAR, NMDAR, Ltype VSCC (v1.3) and TRPV1, respectively, and AEA stands for anandamide. NMDAR and AMPAR were modeled with two-state kinetic models and 1.0mM Mg²⁺ (Destexhe *et al.*, 1995) whereas the model and parameters for the Ca_v1.3 VSCC current was taken from Wolf *et al.* (2005). The TRPV1 current, including its dependence on AEA, was modeled as:

$$I_{TRPV1}(V, AEA) = g_{TRPV1} \cdot V \cdot P_{TRPV1}^{open}(V, AEA)$$
(SI4)

where g_{TPRV1} is the maximal conductance of TRPV1 and the mathematical expression for the P_{TRPV1}^{open} was taken from Matta and Ahern (2007).

Biochemical signaling. Free cytosolic calcium is one of the main signaling actors in the model. To model its dynamics, we assumed it can be transferred from/to two main sources: (i) extracellular calcium, via the plasma membrane channels of eq. (SI3) above and (ii) the endoplasmic reticulum (ER), via the IP3-dependent Calcium-Induced Calcium Release (CICR) system. Hence, the concentration of free cytosolic calcium C was computed according to

$$T_{\rm C}(C)\frac{dC}{dt} = J_{IP_3R} - J_{\rm SERCA} + J_{\rm leak} + J_{\rm NMDAR} + J_{\rm VSCC} + J_{\rm TRPV1} - \frac{C-C_{\rm b}}{\tau_{C_{\rm b}}}$$
(SI5)

where the fluxes J_{IP3R} , J_{SERCA} , J_{leak} from and to the ER in the CICR system were taken from the model of De Pittà *et al.* (2009) - see also Cui et al. (2016) and J_{NMDAR} , J_{VSCC} and J_{TRPV1} are the calcium fluxes from the plasma membrane channels (eq. SI3), computed as $J_x = \xi_x \cdot I_x$ were the ξ_x are constants, $x \in \{NMDAR, VSCC, TRPV1\}$. C_b is the basal cytosolic calcium level resulting from equilibration with calcium diffusion out of the cell and τ_{C_b} is the corresponding time scale. The CICR model also includes the dynamics of the fraction of inactive IP3 (inositol triphosphate) receptor channels (IP3R), h

$$\frac{dh}{dt} = a_2 d_2 \frac{IP3 + d_1}{IP3 + d_3} (1 - h) - a_2 Ch$$
(SI6)

Likewise, the dynamics of C_{ER} , the calcium concentration in the endoplasmic reticulum (ER) was given by

$$T_{C_{ER}}\left(C_{ER}\right)\frac{dC_{ER}}{dt} = -\rho_{ER}\left(J_{IP_{3}R} - J_{SERCA} + J_{leak}\right)$$
(SI7)

where ρ_{ER} is the ER to cytoplasm volume ratio. In eq.(SI5) and (SI7), T_x , x = C or C_{ER} , is a time scaling factor accounting for the presence of endogenous calcium buffers and expressed according to

$$T_{x}(x) = 1 + \frac{B_{T}}{K_{dB}(1 + x / K_{dB})^{2}}$$
(SI8)

Our model also accounts for the biochemical pathways leading to the production of the endocannabinoids 2-Arachidonoylglycerol (2-AG) and AEA, and their subsequent activation of cannabinoid receptors type-1, CB₁R. Principally, the model expresses the kinetics of the following enzymatic reactions: (*i*) synthesis of DAG (diacylglycerol) and IP3 (inositol triphosphate) as a result of glutamate (*G*) binding of metabotropic glutamate receptors (mGluR) and subsequent activation of PLC β as well as Ca-dependent synthesis by PLC δ , (*ii*) transformation of DAG into 2-AG by DAG Lipase α and (*iii*) synthesis of AEA by calcium-dependent fatty acid amide hydrolase (FAAH). The dynamics of this system was modeled with the following equations:

$$\frac{dIP_3}{dt} = v_\beta \frac{G}{G + K_R + K_P C/(C + K_\pi)} + \frac{v_\delta}{1 + IP_3 / \kappa_d} \frac{C^2}{K_\delta^2 + C^2} - v_{3K} CaMKII * \frac{IP_3^{n_3}}{K_3^{n_3} + IP_3^{n_3}} - r_{5P} IP_3$$
(SI9)

$$\frac{dDAG}{dt} = v_{\beta} \frac{G}{G + K_R + K_P C/(C + K_\pi)} + \frac{v_{\delta}}{1 + IP_3 / \kappa_d} \frac{C^2}{K_{\delta}^2 + C^2} - \frac{r_{DGL} \cdot DAGL \cdot \varphi_{DAGL} \cdot DAG}{DAG + K_{DAGL}} - r_{DAGK} \cdot DAG$$
(SI10)

$$\frac{d2AG}{dt} = \frac{r_{DGL} \cdot DAGL \cdot \varphi_{DAGL} \cdot DAG}{DAG + K_{DAGL}} - r_{MAGL} 2AG$$
(SI11)

$$\frac{d\varphi_{DAGL}}{dt} = r_k C^{n_c} \left(1 - \varphi_{DAGL}\right) - r_p \varphi_{DAGL}$$
(SI12)

$$\frac{dAEA}{dt} = v_{AT}C - r_{FAAH} \frac{AEA}{K_{FAAH} + AEA}$$
(SI13)

where *CaMKII** is the amount of activated (phosphorylated) CaMKII (see below); φ_{DAGL} is the fraction of active DAG Lipase α and DAGL its total (activated+not activated) concentration. Eq. (SI9-SI11) account for IP3 degradation by IP3 3-kinase (3K) and inositol polyphosphate 5-phosphatase (5P), while DAG and 2AG are degraded by DAG kinase (DAGK) and monoacylglycerol lipase (MAGL), respectively.

2-AG and AEA are retrograde signaling molecules that are produced in the postsynaptic neuron but diffuse to the presynaptic cell, where they activate CB₁R. We modeled CB₁R activation by 2-AG and AEA using a simple three-state kinetic model: open (x_{CB1R}), desensitized (d_{CB1R}) and inactivated (i_{CB1R}):

$$\frac{dx_{CBIR}}{dt} = \alpha_{CBIR} \cdot eCB \cdot i_{CBIR} - (\beta_{CBIR} + \gamma_{CBIR}) x_{CBIR}$$
(SI14)

$$\frac{dd_{CBIR}}{dt} = -\varepsilon_{CBIR}d_{CBIR} + \gamma_{CBIR}x_{CBIR}$$
(SI15)

where eCB = 2-AG + 0.10 AEA and mass conservation implies $x_{CB1R} + d_{CB1R} + i_{CB1R} = 1$. The open fraction x_{CB1R} was then used to compute CB₁R activation as

$$y_{\rm CB1R} = k_{\rm CB1R} x_{\rm CB1R} + C_1 \tag{SI16}$$

where C_1 is a constant that accounts for the modulation of presynaptic plasticity by other pathways and k_{CB1R} quantifies the strength of CB₁R activation on presynaptic plasticity.

In our model, CB₁R activation (y_{CB1R}) controls the presynaptic weight W_{pre} according to the following rule: W_{pre} decreases for intermediate values of y_{CB1R} , i.e. when y_{CB1R} is comprised between two tLTD thresholds ($\theta_{LTD}^{start} < y_{CB1R} < \theta_{LTD}^{stop}$) whereas W_{pre} increases when y_{CB1R} is larger than a tLTP threshold ($y_{CB1R} > \theta_{LTP}^{start}$). Following Cui et al. (2016), we implemented this rule as

$$\Omega(y_{CB1R}) = \begin{cases}
1 - A_{LTD} & \text{if } \Theta_{LTD}^{\text{start}} < y_{CB1R} < \Theta_{LTD}^{\text{stop}} \\
1 + A_{LTP} & \text{if } \Theta_{LTP}^{\text{start}} < y_{CB1R} \\
1 & \text{otherwise}
\end{cases}$$
(SI17)

and

$$\frac{\mathrm{d}W_{\mathrm{pre}}}{\mathrm{d}t} = \frac{\Omega(y_{\mathrm{CB1R}}) - W_{\mathrm{pre}}}{\tau_{W_{\mathrm{pre}}}(y_{\mathrm{CB1R}} + C_2)} \tag{SI18}$$

Here Ω determines the change of presynaptic plasticity, with a time scale $\tau_{W_{\text{pre}}}$ set to yield rapid changes of W_{pre} for large y_{CB1R} values and very slow changes at very low y_{CB1R} :

$$\tau_{W_{\text{pre}}}(\mathbf{x}) = \frac{P_1}{P_2^{P_3} + x^{P_3}} + P_4 \tag{SI19}$$

To account for experimental observation that the presynaptic weight ranges from about 50 to 300%, W_{pre} was clipped to 3.0 (hard bound).

Postsynaptic plasticity in the model was based on the activation by calcium of calmodulin and CaMKII and the regulation of this system by PKA, calcineurin and protein phosphatase 1 (PP1). The model of Cui et al (2016) for this subsection is based on the model proposed in Graupner & Brunel (2007). In this model the concentration of the calcium/calmodulin complex with four calcium ions bound (CaM) is considered at equilibrium and computed as:

$$CaM = \frac{CaMT}{1 + \sum_{i=1}^{4} (C^{i} \cdot \prod_{j=5-i}^{4} K_{j})}$$
(SI20)

where *CaMT* is the total calmodulin concentration and K_i stands for the equilibrium constant of the binding of the *i*th calcium ion to calmodulin. Each CaMKII enzyme consists of two subunits, each of which contains 6 phosphorylation sites, thus defining 14 phosphorylation states per subunit. Denoting y_i , *i*=0...13 the concentrations of subunit in phosphorylation state *i*, the model of Graupner & Burnel (2007) expresses their dynamics as:

$$\begin{aligned} \frac{dy_1}{dt} &= 6k_6\gamma^2 y_0 - (4k_6\gamma^2 + k_7\gamma + k_{10})y_1 + 2k_{10}\sigma_2^4 \\ \frac{dy_2}{dt} &= (k_7\gamma + k_6\gamma^2)y_1 - (3k_6\gamma^2 + k_7\gamma + 2k_{10})y_2 + k_{10}(y_5 + \sigma_5^7) \\ \frac{dy_3}{dt} &= 2k_6\gamma^2 y_1 - 2(k_7\gamma + k_6\gamma^2 + k_{10})y_3 + k_{10}(\sigma_5^7 + 3y_8) \\ \frac{dy_4}{dt} &= k_6\gamma^2 y_1 - 2(k_7\gamma + k_6\gamma^2 + k_{10})y_4 + k_{10}(y_6 + y_7) \\ \frac{dy_5}{dt} &= k_7\gamma(\sigma_2^3 - y_5) + k_6\gamma^2(y_2 - 2y_5) + k_{10}(2y_9 + y_{10} - 3y_5) \\ \frac{dy_6}{dt} &= k_6\gamma^2(\sigma_2^3 - y_6) + k_7\gamma(2y_4 - 2y_6) + k_{10}(-3y_6 + \sigma_9^{11} + y_{11}) \\ \frac{dy_7}{dt} &= k_6\gamma^2(y_2 + 2y_4 - y_7) + k_7\gamma(y_3 - 2y_7) + k_{10}(-3y_7 + \sigma_9^{11} + y_{11}) \\ \frac{dy_8}{dt} &= k_6\gamma^2 y_3 - 3k_7\gamma y_8 + k_{10}(y_{10} - 3y_8) \\ \frac{dy_9}{dt} &= k_6\gamma^2 y_5 + k_6\gamma^2 y_6 + k_7\gamma(y_7 + 3y_8 - 2y_{10}) + k_{10}(2y_{12} - 4y_{10}) \\ \frac{dy_{11}}{dt} &= k_6\gamma^2 y_9 + k_7\gamma(2\sigma_9^{11} - y_9 - y_{12}) + k_{10}(6y_{13} - 5y_{12}) \\ \frac{dy_{13}}{dt} &= k_7\gamma y_{12} - 6k_{10}y_{13} \end{aligned}$$

where $\sigma_i^j = \sum_{k=i}^j y_k$ and mass conservation implies $y_0 = 2CaMKII_0 - \sigma_1^{13}$, where $CaMKII_0$ is the total CaMKII concentration. In eq. (SI21) above the probabilities that CaM binds to phosphorylated (γ) and dephosphorylated (γ^*) subunit of CaMKII are computed as $\gamma = CaM/(CaM + K_5)$ and $\gamma^* = CaM/(CaM + K_9)$. Moreover, the rate of subunit dephosphorylation (k_{10}) is given by $k_{10} = k_{12}PP1/(K_M + CaMKII^*)$ where K_M is a constant, *PP1* is the concentration of PP1. *CaMKII*^{*} is the total concentration of phosphorylated subunits of CaMKII computed accross all possible states of phosphorylation $CaMKII^* = \sum_{i=0}^{13} m_i y_i$ where m_i is the number of the phosphorylated subunits of CaMKII in state *i* and y_i .

Finally, the model of Graupner & Brunel (2007) assumes that, in addition to calcineurin (CaN), PKA activity depends on CaM according to a Hill-function activation:

$$\nu_x(CaM) = k_x^0 + \frac{k_x}{1 + (K_x/CaM)^{n_x}}$$
(SI22)

where x = CaN or *PKA*, The interaction between PP1 and I1 is then described as:

$$\frac{dI_1}{dt} = \frac{dPP_1}{dt} - \nu_{CaN}I1 + \nu_{PKA}I1_0$$
(SI23)
$$\frac{dPP_1}{dt} = -k_{13}I1 \cdot PP1 + k_{-13}(PP1_0 - PP1)$$

where $PP1_0$ and $I1_0$ are total PP1 and I1 concentrations, respectively.

In agreement with Cui et al. (2016) and Graupner & Brunel (2007), we assumed that postsynaptic plasticity is directly proportional to the calcium-dependent activation of CaMKII and set:

$$W_{\text{post}} = 1 + 3.5 \frac{CaMKII^*}{CaMKII^*_{\text{max}}}$$
(SI24)

where *CaMKII*^{*}_{max} is the maximal concentration of activate (phosphorylated).

Finally, the total synaptic weight was taken as the product of the presynaptic and the postsynaptic contributions above:

$$W_{\text{total}} = W_{\text{pre}} W_{\text{post}} \tag{SI25}$$

Model implementation and numerics. Our mathematical model comprises 36 ordinary differential equations and close to 150 parameters, among which more than one half is constrained by experimental data. Initial conditions were set to the steady-state of each variable in the absence of stimulation. Numerical solution was obtained with the LSODA solver from the ODEPACK fortran77 library with absolute and relative tolerances both equal to 10^{-7} . Initial conditions were set to the steady-state of each variable in the absence of stimulation proceeded until the synaptic weights reach stable values (typically observed around t \approx 5min after the stimulation protocol). We used the final values of the pre- and postsynaptic weights to compute the total synaptic weight change due to the stimulations. Importantly, with the exception of the stimulation protocols, *we have used the exact same equations and parameter values as in* Cui et al. (2016). The list of parameters and their estimated values is given in S2 Table. The current study employs stochastic simulations since the stimulation protocol is stochastic (while the rest of the model is deterministic). Therefore, the model was calibrated using experimental data from deterministic stimulation

protocols (Cui et al., 2016) and we tested here whether it can make successful predictions when we applied stochastic stimulation protocols, for which the model was not calibrated. The results are thus averages over N_{trials} realizations. N_{trials} was varied from 30 to 500 depending on the smoothness of the averaged curves and values of SEM, but $N_{\text{trials}}=50$ in most of the simulations. The computer code of the model is available online from https://github.com/iprokin/Cx-Str-STDP and

https://senselab.med.yale.edu/modeldb/ShowModel.cshtml?model=187605.

S2 Table – Parameter values

A. Intracellular dynamics

Description	Name	Values	Units	Reference
I _{NMDAR} to Ca flow conversion	ξ_{NMDAR}	98	µM/pCol	set to match Ca2+ amplitudes in [1]
I _{VSCC} to Ca flow conversion	ξ_{VSCC}	140	µM/pCol	set to match Ca2+ amplitudes in [2]
I _{TRPV1} to Ca flow conversion	ξ_{TRPV1}	290	µM/pCol	set to match Ca2+ amplitudes in [1]
Total endogenous Ca buffer	B_T	4.5	μΜ	Estimated from our experimental data
Basal cytoplasmic [Ca]	C_{b}	0.1	μΜ	[1,3]
Time scale for Ca exit	τ _{Cb}	0.007	S	Idem
Endogenous Ca buffer affinity	K_{dB}	0.5	μΜ	[3,4]
ER-to-cytosol volume ratio	$ ho_{\scriptscriptstyle E\!R}$	0.3	-	Adapted from [4]
IP3R binding rate (inactivation)	a_2	0.5	$\mu M/s$	Idem
Maximal SERCA pump rate	v_{ER}	8	$\mu M/s$	Idem
IP3R affinity for IP3	d_3	0.9434	μΜ	[5]
Maximal IP3R rate	r_{c}	4	1/s	Adapted from [5]
Basal [Ca] in the ER	$Ca_{ER,b}$	65	μΜ	[6]
Ca leak from the ER	r_l	0.1	1/s	[5]
IP3R affinity for Ca	d_5	0.12	μΜ	Adapted from [5]
IP3R dissociation constant	d_2	3.049	μΜ	Idem
SERCA pump affinity for Ca	K_{ER}	0.05	μΜ	[5]
IP3R affinity for IP3	$d_{_1}$	0.13	μΜ	Idem
PLC ₀ product inhibition	κ_d	1.5	μΜ	Idem
PLCo Ca-activation	K_{δ}	0.1	μΜ	Idem
5P-IP maximal rate	<i>r</i> _{5<i>P</i>}	0.2	1/s	Adapted from [5]
PI3K maximal rate	v_{3K}	0.001	$\mu M/s$	Idem
PI3K Ca-activation constant	K_D	1.5	μΜ	Idem
PLC6 maximal rate	v_{δ}	0.02	$\mu M/s$	[5]
PI3K affinity for IP3	<i>K</i> ₃	1	μΜ	Idem
Glutamate affinity to mGluR	K_R	1.3	μΜ	Idem
regulation by PLCβ termination	K_P	10	μΜ	Idem
PLCβ maximal rate	v_{β}	0.8	$\mu M/s$	Adapted from [5]
PKC Ca-activation constant	K_{π}	0.6	μΜ	[5]
Total CaMKIIa concentration	CaMKII ₀	16.6	μΜ	[7]
Total Calmodulin concentration	CaMT	0.07085	μΜ	Adapted from [7]
PKA Hill number	n _{PKA}	3	-	Idem

Referenced articles: [1] Sabatini, B.L., Oertner, T.O. and Svoboda, K. (2002) *Neuron* 33(3):439-452 [2] Carter, A.G. and Sabatini, B.L. (2004) *Neuron* 44(3):483-493 [3] Jackson, M.B. and Redman, S.J. (2003) *J Neurosci* 23:1612–1621 [4] Nägerl, U.V. *et al.* (2000) *Biophys J* 79: 3009–3018 [5] De Pittà M. *et al.* (2009) *J Biol Phys* 35:383-411 [6] Solovyova, N. *et al.* (2002) *EMBO J* 21:622-630 [7] Graupner, M. and Brunel, N. (2007) *PLoS Comput Biol* 3:e221.

Description	Name	Values	Units	Reference
TRPV1R max. conductance	g_{TRPV1}	0.0003	nS	Estimated from our experimental data
Permeability of L-type VSCC	p_{VSCC}	0.00000102	$\mu M/s$	Value of [1], scaled to ~5,000 spines, radius 1 μm
AMPAR maximal conductance	$g_{\rm AMPAR}$	5.1	nS	Estimated from our experimental data
AMPAR closing rate constant	$eta_{\scriptscriptstyle AMPAR}$	190	1/s	Adapted from [2]
AMPAR opening rate constant	α_{AMPAR}	1.02	1/(µM.s)	Idem
NMDAR maximal conductance	$g_{\rm NMDAR}$	1.53	nS	Estimated from our experimental data
Magnesium concentration	Mg	1	mМ	Directly measured in our experiments
NMDAR opening rate constant	$\alpha_{\scriptscriptstyle NMDAR}$	0.072	1/(µM.s)	Set to emulate the experimental kinetics of NMDAR- transported Ca in [3]
NMDAR closing rate constant	$eta_{\scriptscriptstyle NMDAR}$	100	1/s	Idem
Resting membrane potential	V_L	-70	mV	Directly measured in our experiments
Leak conductance	g_L	10	nS	Idem
Membrane capacitance	C_m	0.1	nF	Idem
Extracellular Ca2+ concentration	Ca _{out}	5000	μΜ	[1]
Duration of depolarization step	DC_{dur}	0.03	S	From the experimental stimulation protocol
Amplitude of depolarization step	$DC_{\rm max}$	495	pA	Idem
Amplitude of action current	$AP_{\rm max}$	7020	pA	[4]
Time constant of action current	$ au_{\scriptscriptstyle bAP}$	0.001	S	Idem
Glutamate peak concentration in the cleft	G_{\max}	2000	μΜ	Estimated from our experimental data
Glutamate decay time constant in the cleft	$ au_{_G}$	0.005	S	Estimated from our experimental data
Delay to bAP outset	δ	0.015	S	From the experimental stimulation protocol
Stimulation frequency	F _{pairings}	1	Hz	From the experimental stimulation protocol

B. Electrophysiology

Referenced articles: [1] Wolf, J.A. *et al.* (2005) *J Neurosci* 25(40):9080-9095 [2] Destexhe, A., Mainen, Z., and Sejnowski, T. (1994) *J Comput Neurosci* 1:195-230 [3] Sabatini, B.L., Oertner, T.O. and Svoboda, K. (2002) *Neuron* 33(3):439-452 [4] Fino, E., *et al.* (2010) *J Physiol* 588:3045-3062.

Description	Name	Values	Units	Reference
Scaling factor for endocannabinoid contribution to plasticity	k _{CB1R}	3000	$1/\mu M$	Estimated from our experimental data
presynaptic plasticity time scale	P_1	1e-9	S	Set to yield rapid / slow changes of Wpre for high / low 2-AG values, respectively
presynaptic plasticity time scale	P_2	1e-5	-	Idem
presynaptic plasticity time scale	P_3	7	-	Idem
presynaptic plasticity time scale	P_4	2	S	Idem
The lower limit of LTD induction	$ heta_{\scriptscriptstyle LTD}^{\scriptscriptstyle start}$	0.027	-	Estimated from our experimental data
The upper limit of LTD induction	$ heta_{\scriptscriptstyle LTD}^{\scriptscriptstyle stop}$	0.047	-	Idem
The constant determining the rate of LTD induction	A_{LTD}	0.65	-	Idem
The lower limit of LTP induction	$ heta_{LTP}^{start}$	0.087	-	Idem
The constant determining the rate of LTP induction	A_{LTP}	10.8	-	Idem
CB1R opening rate constant	$lpha_{_{CB1R}}$	0.2402	1/(µM.s)	Idem
CB1R closing rate constant	$\beta_{_{CB1R}}$	11.072	1/s	Idem
CB1R desensitization rate constant	γ_{CB1R}	416.38	1/s	Idem
CB1R closing rate constant	\mathcal{E}_{CB1R}	0.047796	1/s	Idem
DAGL α affinity for DAG	K _{DAGL}	30	μΜ	[1]
Maximal rate of MAG lipase	r _{MAGL}	0.5	$\mu M/s$	Set for rapid turnover dynamics.
Maximal DAGL α rate	r _{DGL}	20000	$\mu M/s$	Idem
DAG kinase rate	r _{DAGK}	2	s ⁻¹	Idem
Hill number for DAGL activation by Ca	n _c	6	-	Estimated from our experimental data
DAGL dephosphorylation rate	r_{P}	380	s ⁻¹	Idem
DAGL phosphorylation rate	r_{K}	50	$\mu M^{-1}.s^{-1}$	Idem
Total DAGL concentration	DAGL	1	μΜ	Idem
N-acetyltransferase activity	v_{AT}	0.2	1/s	Idem
FAAH Michaelis-Menten constant	K _{FAAH}	1	μΜ	[1]
FAAH enzyme activity	r _{FAAH}	4	μM/s	Estimated from our experimental data

C. Endocannabinoid dynamics

Referenced articles: [1] Okamoto, Y. et al. (2004) J Biol Chem 279:5298-5305.

S3 Text – Mechanisms of STDP expression in the model

We give here an overview of the mechanisms that give rise to the expression of eCB-tLTP, eCB-tLTD and NMDAR-tLTP in our mathematical model. All the examples given here concern regular (deterministic) STDP protocols with constant IPI and spike timings.

Our model combines the two signaling pathways involved in cortico-striatal STDP: a first signaling pathway leading from NMDAR to calmodulin and CaMKII with a second that couples mGluR and cytosolic calcium to eCB production and the resulting activation of CB₁R (see Fig. 1f1). In the model, we assume that the total synaptic weight (W_{Total}) is the product of presynaptic (W_{Pre}) and postsynaptic (W_{Post}) weights (see Supporting Information S1).

On the postsynaptic side, W_{Post} is proportional to the amount of CaMKII activated by the NMDAR pathway. CaMKII in our model forms a bistable system that settles at long times either on a "DOWN" state where CaMKII is mostly inactive (no plasticity) or on an "UP" state characterized by high levels of activated CaMKII (Fig. 2b1) (NMDAR-tLTP). In the example of the figure, the time needed for CaMKII dephosphorylation is larger than the interval between two successive pairings, so activated CaMKII accumulates with the number of pairings during the protocol. For a transition from the UP to the DOWN state to occur, the amount of activated CaMKII needs to overcome the separatrix between the two states. The separatrix is reached in the example of Supplementary Figure S2 (in Supporting Information S4 below) for circa 50-60 post-pre pairings (with Δt_{STDP} =-15 ms) (Fig. S2a1). Therefore, in the conditions of Figure S2 (short post-pre pairings), W_{Post} converges to the UP state (potentiation) only when the STDP protocol comprises > 50 post-pre pairings. With pre-post pairings, the calcium released after each pairing activates less calmodulin than with post-pre protocols. As a result, activated CaMKII never reaches the threshold for the UP state (Fig. S2a2). To summarize, the model successfully reproduces the experimental observations that (i) NMDAR-dependent tLTP emerges after 50-75 short post-pre pairings and (ii) no NMDAR-dependent LTP is obtained with pre-post pairings (Cui et al., 2016).

When a stimulation protocol composed of 100 pairings is applied, the evolution of the amplitude of the triggered calcium peaks is biphasic (Fig. S2c): it increases up to 10-20 pairings and decreases afterwards to settle to constant amplitude after circa 50 pairings. For the first 10-20 stimulations, the iterated activation of mGluRs leads to the accumulation of IP3, which contributes an extra influx of calcium from the endoplasmic reticulum through calcium-induced calcium release. This extra-boost of cytoplasmic calcium however vanishes when the number of pairings increases further because the concentration of calcium in the ER

decreases. This decay ER calcium eventually compensates the effect of IP3 accumulation on the calcium-induced calcium release, thus stabilizing the amplitude of the calcium peaks for N_{pairings} >50. Moreover, since the width of the postsynaptic calcium peak in the model is larger with post-pre than pre-post pairings (Fig S2b), the fraction of calcium-activated DAGLa is much larger for post-pre pairings. As a result, the amplitude of eCB peaks and, ultimately, the amplitude of the fraction of activated CB₁R (y_{CB1R}), show a more pronounced biphasic profile (Fig S2d). The biphasic trend is further amplified at the level of CB₁R activation because of CB₁R receptor desensitization that amplifies the decay above 20 pairings (Fig S2d). The amplitude of the y_{CB1R} peaks increases sharply for the first 10-20 pairings, and progressively decays afterwards to converge to constant amplitude. As show in the figure, y_{CB1R} reaches large values only for short post-pre pairings (Δt_{STDP} around -15 ms) while even short pre-post pairings ($0 < \Delta t_{STDP} < 10$ ms) do not give rise to such large amplitude peaks.

The dynamics of the level of y_{CB1R} is responsible for the emergence of eCB-dependent plasticity in the model. Indeed, in the model, W_{Pre} depends on y_{CB1R} with the piecewise constant function shown in Figure S2e and its 3 thresholds Θ_{LTD}^{start} , Θ_{LTD}^{stop} and Θ_{LTP}^{start} : if y_{CB1R} reaches moderate amounts, i.e. lies within $[\Theta_{\text{LTD}}^{\text{start}}, \Theta_{\text{LTD}}^{\text{start}}], W_{Pre}$ decays (thus emulating eCB-tLTD); whereas W_{Pre} increases (eCB-tLTP) if $y_{CB1R} > \Theta_{LTP}^{start}$ (see the dashed lines in Fig. S2d and summary in Fig. S2e). With short pre-post pairings ($10 < \Delta t_{STDP} < 40$ ms), y_{CB1R} enters the LTD range during most of protocol so each pairing contributes a increment of W_{Pre} reduction. Since W_{Post} is not changed with positive, pre-post pairings (Fig. S2a2), pre-post pairings globally lead to a progressive reduction of W_{Total} , i.e. the expression of eCB-tLTP. The situation is different for post-pre pairings. The amplitude of y_{CB1R} peaks overcomes Θ_{LTP}^{start} for ≈ 5 to 30 post-pre-pairings with short Δt_{STDP} (around -15 ms), yielding a strong increase of W_{Pre} . Given that more that 50 post-pre pairings are needed to alter W_{Post} (Fig. S2a1), the expansion of W_{Pre} leads to eCB-tLTP. However, when $N_{pairings} > 30$ post-pre pairings, y_{CB1R} amplitude gets back below Θ_{LTP}^{start} so that the initial increase of W_{Pre} does not continue beyond 30 post-pre pairings. This property of the model dynamics emulated the experimental observation that eCB-tLTP disappears when the number of pairings becomes larger than 25-30.

Therefore, the mechanisms that lead to the expression of eCB-dependent plasticity with regular deterministic stimulation protocols in our model can be summarized as follow. eCB-tLTD requires moderate levels of CB₁R activation that are reached with pre-post but not pre-post pairings. The expression of eCB-tLTP however demands larger amounts of CB₁R

activation, which are reached only during the first 5-30 post-pre pairings, where every component of the system contributes maximally to CB_1R activation. When the number of post-pre pairings becomes larger than 30, calcium efflux from the internal calcium stores decreases and CB_1R desensitization increases: CB_1R activation becomes insufficient to support such high values of CB_1R activation, so eCB-tLTP vanishes.

S4 Figure – Supplementary Figures





Supplementary Figure 1: $m_{\Delta t}$ values were not significantly different among STDP paradigms. $m_{\Delta t}$ were not significantly different in control and jittered conditions for 100 post-pre, 100 pre-post and 10 post-pre pairings (empty circles: individual neurons; black circle: average). Error bars represent the SEM. ns: not significant.

Figure S2



Supplementary Figure 2: Mechanisms of STDP expression in the model with deterministic stimulation protocols.

(a) Changes in the amount of active CaMKII starting from the down (non-activated) state. The number of pairings, N_{pairings} , is indicated for spike-timing $\Delta t_{STDP} = -15$ (a1) or + 15 (a2) ms. (b) Intracellular calcium changes for the first pairing in post-pre (b1) or pre-post (b2) pairing protocols. (c) Predicted dynamics of the cytoplasmic calcium for 100 post-pre (c1) or 100 pre-post pairings (c2) and corresponding activation of CB1R, y_{CB1R} (d). The spike-timing Δt_{STDP} for each curve is indicated in the legend. (e) The presynaptic weight W_{Pre} changes as a function of y_{CB1R} : it decreases (LTD) when y_{CB1R} reaches intermediate values and it increases (LTP) if y_{CB1R} overcomes the LTP threshold. The corresponding thresholds are reported in D as dashed lines. For all the simulations shown, the stimulation protocols were deterministic (regular) with a frequency of 1Hz.

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