

Environmental enrichment enhances precision and remodeling speed of synaptic nanoarchitecture revealed by crosstalk free two-color *in vivo* STED nanoscopy

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Abstract

Synaptic plasticity underlies long-lasting structural and functional changes to brain circuitry and its experience-dependent remodeling can be fundamentally enhanced by environmental enrichment. It is unknown, however, whether and how environmental enrichment alters the morphology and dynamics of individual synapses. Here, we present a virtually crosstalk free, two-color *in vivo* STED microscope to simultaneously superresolve the dynamics of the postsynaptic density protein PSD95 and spine geometry. With environmental enrichment, the size distribution of PSD95 and spine head sizes were narrower than in controls, indicating that synaptic strength is set more precisely with environmental enrichment. Spine head geometry and PSD95 assemblies were highly dynamic exhibiting multiplicative size changes. With environmental enrichment, the topography of PSD95 nanoorganization was more dynamic and the magnitude of reorganization increased systematically with time. Thus, two-color *in vivo* time-lapse imaging of synaptic nanoorganization uncovers a unique synaptic nanodynamics associated with the enhanced learning capabilities under environmental enrichment.

INTRODUCTION

Excitatory synapses in the brain are mainly formed on dendritic spines and are regarded as the central locus of memory formation. Cognitive, sensory and motor learning can induce specific changes to the cortical circuitry over the entire lifespan. Such experience-dependent plasticity is based on synaptic plasticity, plays a key role for normal brain function and is thought to be neuroprotective against neurodegenerative diseases^{1,2}. Memory relies on structures being persistent over long periods of time up to a life-time. There is, however, also evidence that synaptic structure can be highly volatile^{3,4}. Synaptic connections thus presumably undergo continuous spontaneous remodeling on top of which activity-dependent changes inscribe memory traces.

Neural activity per se is not required to form an intact synapse; for instance, despite abolishing presynaptic release⁵, or silencing a network, morphologically intact spines are formed⁶. At the beginning of the century, much evidence had shown a strong correlation between the morphological transformation of the spine, termed structural plasticity, and changes in synaptic transmission, termed functional plasticity⁷. However, recent evidence suggests that structural and functional plasticity are not fully interconnected⁸. For example, 'silent' synapses are largely silent at rest but the spine's shape is of mature morphological shape⁹. A main link between structure and function is F-actin, which drives spine morphological remodeling and anchors scaffolding proteins of the post-synaptic density (PSD) such as PSD95. PSD95 in turn anchors amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) receptors via transmembrane AMPAR-regulatory proteins (TARPs)^{10,11} which is the main contribution to synaptic strengthening after long-term potentiation (LTP). Both, anchoring of AMPA receptors and the actin remodeling, are highly dependent on Ca^{2+} influx after activation, but the intracellular signaling pathways are different as far as known⁸. In addition intrinsic turnover of the cytoskeleton and its different actin pools impose independent contributions to structural remodeling⁸. A plethora of studies show a strong temporal link between AMPA receptor mediated currents and spine volume; both increase rapidly after glutamate uncaging¹². However, there is recent evidence that not all synaptic proteins increase instantaneously but increase rather with a delay after stimulation. For instance, the number of the scaffolding proteins Homer, Shank and PSD95 at the synapse increases with a delay of ~1 hour after LTP^{13,14}. Thus, the correlation between spine and PSD size which is well documented by electron microscopy^{14–17} may be temporally much less correlated during LTP. These observations suggest that the PSD and spine head plasticity are regulated differentially in the short term and could be more independent than previously thought.

In animal models, experience dependent plasticity is approached by environmental enrichment (EE) which provides the animal with multi-sensory stimulation, cognitive activity, social interactions and physical exercise. Various alterations of brain structure and function by EE have been shown and include increased maturation of new neurons, enhanced expression of signaling molecules and enhanced synaptic plasticity². Structural changes include an increased dendritic branching and spine density in specific cortical regions^{18,19} and an increase in pre- and postsynaptic protein levels in major cortical brain regions²⁰. In mouse models of brain diseases strong effects of environmental enrichment are documented². However, none of these studies investigated the ultrastructure or dynamic of the PSD or spine morphology.

Here, we use nanoscale imaging to map temporal changes of the spine head and PSD *in vivo* and address mainly the following questions. Firstly, how does experience change the synaptic nanoorganization and plasticity? Therefore, we quantified structural correlates as well as the structural plasticity of dendritic spines and PSD95. Secondly, how are fluctuations of the PSD size correlated in time with spine head fluctuations *in vivo*? For that purpose, we developed a virtually crosstalk free two-color STED microscope to image the spine morphology and PSD95 *in vivo* at nanoscale resolution.

This is a significant technical advance over previous two-color live-cell STED microscopy approaches which were performed with a crosstalk of 92% and 27%²¹ and therefore required unmixing of the two channels. Hitherto, most *in vitro* and *in vivo* studies have been performed using two-photon microscopy to visualize spines and PSDs^{12–14,16,22–24}. However, due to its limited optical resolution of 300–500nm it was impossible to quantify precisely spine parameters such as head size and neck length or diameter. Moreover, PSD nanostructures such as perforations²⁵ are not detectable with two-photon microscopy. Several studies utilized superresolution microscopy to dissect the synaptic nanoorganization of PSD95 in neuronal cultures or fixed brain slices and showed an arrangement of PSD95 in nanoclusters of 50–250 nm in size^{26–29}. Recently, we studied the nanoplasticity of PSD95 with STED microscopy in a PSD95-EGFP transgenic mouse *in vivo*. We found that PSD95 assemblies were often perforated, formed ring-like or horse-shoe shapes or were clustered; all shapes changed within a few hours³⁰. Here, we utilized a transcriptionally regulated antibody-like protein to visualize endogenous PSD95 without introducing overexpression artefacts³¹. Together with a membrane label we recorded temporal morphological changes of the spine head and PSD95 nanoorganization of mice raised in an enriched environment (EE) versus raised in standard cages (Ctr). Effects of environmental enrichment on structure and function have been shown for multiple brain regions². We chose to investigate the synaptic plasticity in the visual cortex since several studies have shown enhanced plasticity of this area in rodents housed in EE cages^{32,33} and it is easily accessible with light microscopy.

Our data show changes in width but not mean value of spine head and PSD95 size distributions following EE. In EE housed mice, PSD95 underwent much stronger directional changes in size as well as in morphology while spine head dynamical changes were unaltered. The changes of PSD95 and spine head size were only mildly correlated and did not show directional change.

RESULTS

Virtually crosstalk free two-color STED microscope

To unambiguously dissect the dynamics of distinct synaptic components and examine their interactions crosstalk free nanoscale imaging is of critical importance. We thus established a virtually crosstalk free two-color STED microscope for imaging of EGFP and EYFP or Citrine, respectively. Previous attempts featuring two detection channels were suffering of high crosstalk²¹. In addition to two-color detection, we therefore implemented a two-color excitation to specifically excite the green or yellow fluorescent protein (Fig.1A). This was achieved by adding a pulsed 520nm green laser line to the 483nm excitation of our previously described *in vivo* STED microscope³⁴. 520nm is at the red edge of the EGFP excitation band and therefore predominantly excites EYFP (or Citrine) (Fig.1B). The STED beam passed a vortex phase plate to create a donut-shaped intensity profile in the focal plane of the objective as previously described³⁵. All three beams, two excitation beams and one STED beam, were spatially aligned by dichroic mirrors and the pulses were temporally aligned to overlap in the focal plane. The fluorescence emission was split by a dichroic mirror and detected in two spectrally separated detection channels. EGFP was preferentially detected by detection 1 (Det1, Fig.1B) at 498–510nm while EYFP/Citrine emission was detected mostly by detection 2 (Det2, Fig.1B) at 532–555nm. The 483 and 520nm excitation was switched on and off alternately while recording two consecutive line scans to temporally separate the EGFP and Citrine emission. To determine the crosstalk between both channels, we investigated one-color labelled cells expressing EGFP or EYFP. Live-cell imaging showed that we have designed a virtually crosstalk free microscope with only ~8% crosstalk of EYFP in the EGFP channel and ~5% of the EGFP signal in the EYFP channel (Fig.1C).

Two-color *in vivo* STED microscopy of endogenous PSD95 and spine morphology

Having established the virtually crosstalk free detection of two STED approved fluorescent proteins^{36,37}, we set out to simultaneously superresolve the nanoorganization of the post-synaptic density protein PSD95 and the associated spine morphology *in vivo*. To this end, we generated recombinant adeno-associated viral (AAV) particles expressing fusion proteins under control of the human Synapsin promoter (hSyn). To visualize PSD95, we expressed the transcriptionally regulated antibody-like protein PSD95-FingR that has been shown to label endogenous PSD95³¹ attached to the fluorescent protein Citrine. To label the membrane, we expressed myr-EGFP-LDLR(Ct), a combination of a myristoylation site (myr), EGFP and the C-terminal (Ct) cytoplasmic domain of low density lipoprotein receptor (LDLR), a potent marker for the dendritic membrane³⁸. In order to reduce the density of labelled neurons, the fluorescent labels were incorporated into AAVs in reverse orientation in the vector between double-floxed inverted open reading frames (DIO). Expression of the labels was enabled by Cre recombinase expressing AAVs that were co-injected at low concentration. Hence, in this system the density of labeled neurons is continuously adjustable by the dilution of the Cre expressing AAV. With this set of AAVs we were able to adapt the brightness of the PSD95 and membrane labelling independently of the cell density. Thus, we co-injected the three different AAVs into pyramidal cell layer 5 of the visual cortex. Three to six weeks after transduction, the mice were anaesthetized and a cranial window was inserted over the visual cortex as described before³⁹. STED microscopy visualized dendrites, spines, and attached PSD95 assemblies in layer 1 of the visual cortex at superresolution; the images were basically background-free and of high contrast due to the low number of fluorescently labelled neurons (Fig.1C). Due to the low crosstalk, a computational post-processing such as unmixing was not required. We observed mainly dendrites that expressed both, EGFP and Citrine, and only rarely found dendrites expressing either Citrine or EGFP alone. The inset in Fig.1D reveals a perforated nanoorganization of the endogenous PSD95 that would not be detectable with conventional *in vivo* microscopy (Fig.1D) due to the insufficient resolution of two-photon microscopy. In general, we obtained a resolution of at least ~70nm for Citrine and ~84nm for EGFP (Fig.S1A,B).

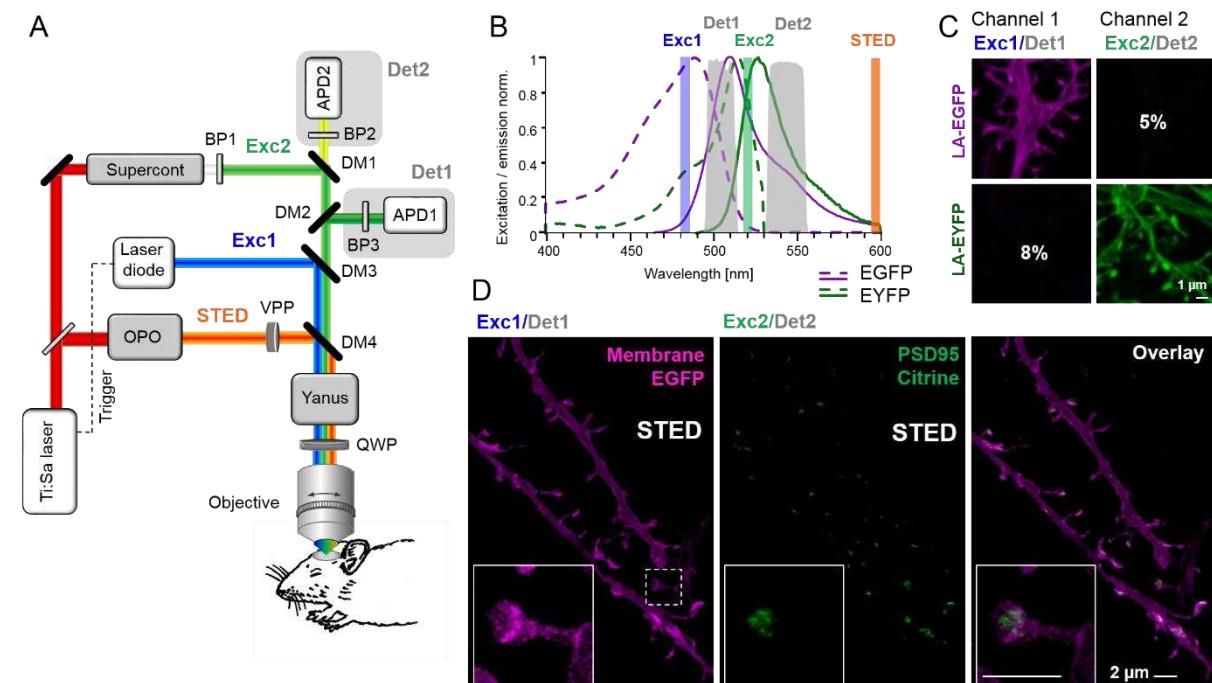


Figure 1: Virtually crosstalk free two-color *in vivo* STED microscopy. (A) Custom-designed *in vivo* STED microscope with pulsed 483 nm (Exc1) and 520 nm (Exc2) excitation and 595 nm STED laser. APD:

Avalanche photon detector, BP: bandpass filter, Det: detection, DM: dichroic mirror, OPO: optical parametric oscillator, QWP: quarter wave plate, VPP: vortex phase plate. Confer Material and Methods section for details. **(B)** Excitation (dashed line) and emission (solid line) spectrum of EGFP and EYFP and selective excitation and detection. **(C)** Cultured hippocampal neurons expressing the actin marker Lifeact (LA)-EGFP or LA-EYFP. Excitation of EGFP at 483 nm close to its excitation maximum (B) and detection at 498–510 nm (Det1) reduces the crosstalk to 5% in Channel 2 (C). EYFP or Citrine is excited at 520 nm close to its maximum (B) and detected at 532–555 nm (Det2) which results in a low crosstalk of 8% in Channel 1 (C). **(D)** *In vivo* STED microscopy image of apical dendrite in layer 1 of the visual cortex of an anaesthetized mouse. Labelling of membrane (myr-EGFP-LDLR(Ct)) and PSD95 (PSD95-FingR-Citrine) visualizes the spine morphology at superresolution and PSD95 nanoorganization (D, inset). Images are smoothed and represent maximum intensity projections (MIPs) but without unmixing due to the low crosstalk.

EE housed mice exhibit more sharply defined spine head and PSD95 sizes than Ctr

Having established the two-color *in vivo* STED microscope, we addressed the question of whether spine or post-synaptic density exhibit systematic differences between EE and Ctr mice. EE mice were housed in a large, two-floor cage equipped with three running wheels for physical exercise, a labyrinth for cognitive stimulation, a tube and a ladder to change between the two levels, and were kept in groups of up to 12 female mice to allow manifold social interactions (Fig.S2). Control mice were raised in pairs in standard cages without any equipment (Fig.S2). The mice were transduced with AAVs as described above and 3–6 weeks after transduction, a cranial window was implanted above the visual cortex. With the two-color STED microscope, we acquired z-stacks mainly of dendrites, which were in parallel to the focal plane. To analyze differences in morphology of the dendritic spines, we measured the length and width of the spine head and computed the spine head cross sectional area approximating the head by an ellipse (Fig.2A). Analogously, we measured the length and width of the PSD95 assembly and computed the elliptic area (Fig.2A, cf. section Methods for details). We only considered spines containing a PSD95 label and therefore spines, which most likely form functional synapses. We occasionally observed but did not analyze long, thin spines without a head, sometimes called filopodia. We determined a median spine head area for EE housed mice of 0.493 (interquartile range (IR): 0.328–0.813) μm^2 and 0.477 (IR: 0.297–0.854) μm^2 for Ctr mice. The area of PSD95 on the same set of spine heads was 0.153 (IR: 0.097–0.255) μm^2 in median for EE mice and 0.148 (IR: 0.089–0.280) μm^2 for Ctr mice which is in accordance with previous EM studies in the mouse visual cortex⁴⁰. For comparison, we also determined the brightness of PSD95 assemblies, an often used stand-in for spine size in two-photon microscopy, and found a moderate correlation of $R^2=0.76$ between the brightness and the size determined from our STED data (Fig.S3A). The spine head area correlated with the PSD95 area for EE and Ctr (Fig.2B), corroborating EM and two-photon microscopy studies^{14–16}. The histograms of spine head area (Fig.2C) and PSD95 area (Fig.2D) were positively skewed and the distributions were significantly different between EE and Ctr mice. To further dissect these differences, we next plotted the histograms of the logarithm of spine head areas (Fig.2E) and PSD95 areas (Fig.2F). All four histograms (Fig.2E,F) are symmetric and were well described by a Gaussian function. This in line with prior work showing that the spine and PSD95 fluorescence is log-normally distributed in general which is predicted by multiplicative processes of ongoing spine plasticity^{6,22}. The Gaussian functions fitting the spine head area (Fig.2E) and PSD95 area (Fig.2F) are almost centered and indeed the average area is not significantly different between EE and Ctr mice (Fig.2G,H). However, we observed a significant difference in variance (Fig.2I,J) visible in the different widths of the Gaussian distributions in Fig.2E,F. Spine density was not significantly different between EE and Ctr housed mice (Fig.S3B). These results

show that although the average areas of the spine head and PSD95 are similar for EE and Ctr mice, their variance in size is smaller for the enrichment mice than for the control mice, indicating that the synaptic strength are more sharply defined in EE mice.

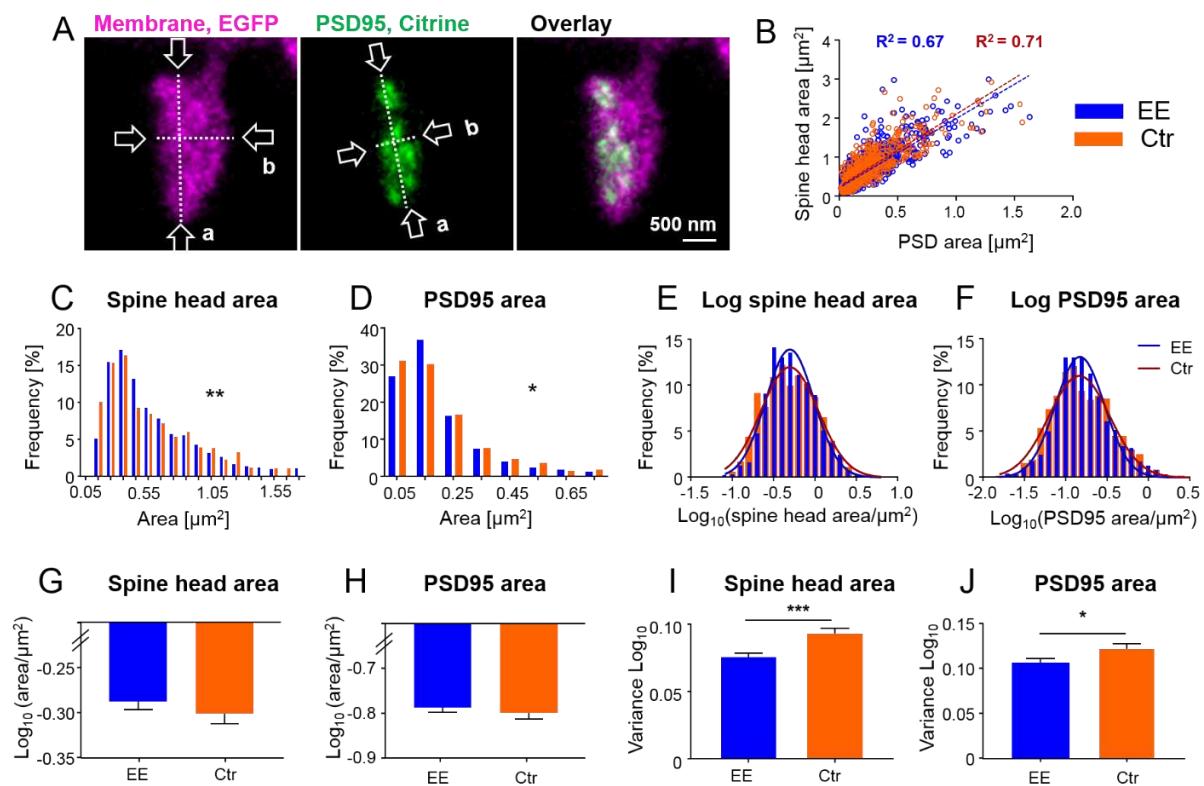


Figure 2: The size distributions of the spine head and PSD95 area are sharper for mice housed in environmentally enriched (EE) than standard (Ctr) cages. (A) STED image of dendritic spine head (left, magenta), associated PSD95 (middle, green) and overlay (right). For both, the length (a) and width (b) were measured to estimate the area. **(B)** Strong correlation of absolute spine head and PSD95 area in Ctr (orange) and EE (blue) housed mice. Linear regression lines are dashed and R^2 values are displayed (EE and Ctr, deviation from zero: $p < 0.0001$). **(C, D)** Frequency distributions of spine head area (C) and PSD95 area (D) are positively skewed and significantly different between EE and Ctr housed mice (Kolmogorov-Smirnov test, C: ** $p = 0.009$, D: * $p = 0.039$). Graphs display only sizes with a frequency $> 1\%$ and center of BIN. **(E, F)** Same data as shown in (C, D), plotted on a logarithmic scale. Solid lines represent Gaussian functions fitted to the respective histogram. **(G–J)** Mean (G: $\text{EE}_{\text{spine}}: -0.288$, $\text{Ctr}_{\text{spine}}: -0.301$, H: $\text{EE}_{\text{PSD}}: -0.787$, $\text{Ctr}_{\text{PSD}}: -0.799$) and variance (I, J) of logarithmic data shown in (E, F) + SEM (Unpaired *t*-test with Welch's correction: G: $p = 0.34$, H: $p = 0.47$, I: *** $p < 0.001$, J: * $p = 0.035$). Number of analyzed mice and spines: Ctr: 4x ♀mice, $n_{\text{spine}} = 737$; EE: 4x ♀mice, $n_{\text{spine}} = 1052$.

Correlated changes of PSD95 and spine head area over minutes to hours

A strong correlation of PSD and spine size has been reported previously and we confirmed it here for the PSD protein PSD95 (Fig.2B). Whether dynamic changes between these two features are also strongly linked, however, has remained unknown. We thus asked whether there were also strong correlations in temporal changes of these parameters. To answer this question, we performed time-lapse STED microscopy of spine morphology and PSD95 for EE and Ctr housed mice for up to 4 hours as described above. We recorded STED images at different fields of view (FOVs) for three time points

at varying time intervals Δt of 30min, 60min or 120min. Figures 3A and B show representative time-lapse STED images at a time interval of 30min. The average spine head area and PSD95 area did not change over the time period of 120min (Fig.3C). Thus, we did not observe directed changes as a result of light induced stimulation or phototoxicity and neither observed blebbing. We analyzed the spine head and PSD95 area for each time point as described above and computed their normalized size changes over time. Thus, we normalized the changes between two time points to the sum of the area at these time points. With this normalization positive changes (growth) as well as negative changes (shrinkage) are symmetric and have a boundary value of ± 1 . Scatter plots showed normalized changes of spine head area as a function of the changes of PSD95 area for the different time intervals (Fig.3D–F). In all three plots the data points are scattered over all 4 quadrants (for percentage changes refer to Fig. S4A–C). This means that we observed positively correlated changes of growing spine heads and growing PSD95 areas (or shrinking, respectively) (Fig.3G) as well as anti- or negatively correlated changes, i.e. shrinking PSD95 areas on a growing spine or vice versa (Fig.3H). A linear regression revealed a small positive correlation (R^2 between 0.02 and 0.11, Fig.3D–F), such that e.g. an increase of PSD95 area is more frequently accompanied by an increase in spine head area. To further quantify these changes, we asked how strong the changes were and asked whether they changed over time in magnitude. Growing spine heads grew in average by 23–29% (and shrinking spine heads shrunk by 17–24%) (Fig.S4D); growth and shrinkage increased slightly with increasing time intervals, but not significantly. PSD95 assemblies grew in average by 22–38% (Fig. S4E) and shrunk by 25–35% (Fig.S4E). About half of the spine heads and PSD95 assemblies grew while the other half shrunk (Fig.S4F,G). To sum up, spine heads and PSD95 assemblies were highly dynamic and grew in average by >22% and shrunk by >17% already over 30 min. These average changes, however, were not significantly different for EE and Ctr and did not increase significantly with increasing time interval.

We next assessed whether the spread of the temporal changes would be different, similar to what we have observed for the distribution of sizes. To this end, we examined the variance of our parameters and computed the total variance for spine head and PSD95 area changes for the different time intervals (Fig.S4H). A significant increase in variance was observed for EE housed mice after 30min while the increase after 120min was just not significant (Fig.S4H). We separately tested whether this difference was more prominent when computing the variance for correlated and anti-correlated changes separately. To this end, we performed a principal component analysis (PCA) for all three time intervals. Principal component 1 (PC1) is by definition the axis along which the data shows the maximum of variance and the eigenvalue of PC1 is the variance along this axis. The PC1 vector was for all three time intervals in the first (and third) quadrant (Fig.3D–F) and therefore attributing for positive correlation. The variance of PC1 was slightly higher for EE mice at all three time points with a significant increase for the 120min time interval (Fig.S4I). The second principal component (PC2) is by definition perpendicular to PC1 and therefore in the second (and fourth) quadrant (Fig.3D–F). Thus, PC2 quantifies negatively correlated changes in spine head and PSD95 area (Fig.S4J). Interestingly, the variance along PC2 was relatively high (~ 30 –40% of total, Fig.S4K), quantifying that a negative correlation of the size changes of PSD95 assemblies and spine head is not a rare phenomenon. In total, these results start to suggest a slightly higher temporal dynamics in EE mice due to the higher total variance for two time-points and higher PC1 value for EE mice. However, the average changes did not differ, neither over time nor between EE and Ctr housed mice.

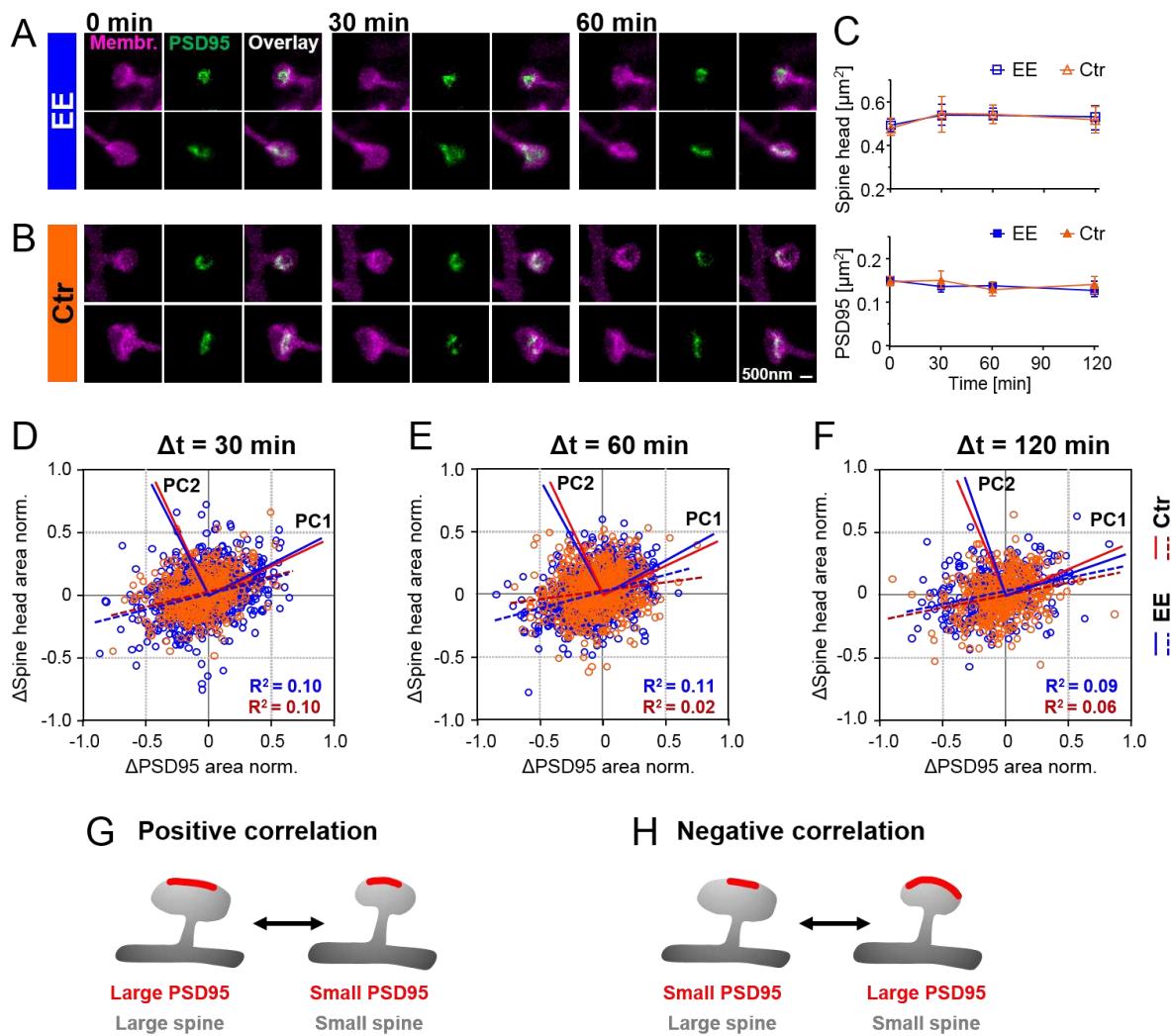


Figure 3: Temporal changes of spine head and PSD95 area are positively and negatively correlated.

(A, B) Representative sections of spine heads and corresponding PSD95 of time-lapse *in vivo* two-color STED microscopy images. EE (A) and Ctr (B) housed mice at time points of 0 min, 30 min and 60 min. Images are smoothed and shown in MIP. Scale bar 500 nm. (C) Average of spine head areas and PSD95 areas of Ctr (triangle, orange) and EE (square, blue) housed mice over time. Plots show median \pm 95% confidence interval (CI). (D–F) Normalized changes in spine head and PSD95 area after 30 min, 60 min and 120 min time intervals. Linear regression lines are dotted (deviation from zero: $p < 0.0001$). Solid lines represent the principal components 1 and 2 (PC1, PC2) of the principal component analysis (PCA) for Ctr (red) and EE (blue), respectively. (G, H) Illustration of temporal changes between spine head size and PSD95 area. (G) Growth and shrinkage of spine head size and the area of PSD95 assemblies goes hand-in-hand, i.e. the changes are (positively) correlated. (H) An anti-correlation or negative correlation is observed for a growing PSD95 assembly on a shrinking spine and vice versa. (C–F) Number of mice and spines: $\Delta t = 30$ min: 4x ♀mice, $n_{\text{spine}} = 453$ for Ctr and 4x ♀mice, $n_{\text{spine}} = 958$ for EE; $\Delta t = 60$ min: 4x ♀mice, $n_{\text{spine}} = 646$ for Ctr and 4x ♀mice, $n_{\text{spine}} = 1027$ for EE; $\Delta t = 120$ min: 3x ♀mice, $n_{\text{spine}} = 386$ for Ctr and 4x ♀mice, $n_{\text{spine}} = 358$ for EE.

Multiplicative downscaling of PSD95 area increases over time in EE but not Ctr housed mice

Since our parameters were log-normally distributed, indicating a multiplicative dynamics, we asked whether the temporal changes were also regulated by multiplicative processes^{6,22}. Thus, we plotted

the synaptic size after different time points as a function of the original size (Fig.4A–F). A linear regression analysis revealed a slope <1 and positive y-intercept for spine head and PSD95 area, for both EE and Ctr housed mice, indicating that large spines tend to shrink and small spines tend to grow which is often denoted as multiplicative dynamics. A recent study modelling spine dynamics as a Kesten process termed the slope as shown in Fig.4 as time-dependent multiplicative downscaling factor^{6,41}. Using this metric of ongoing synaptic dynamics we observed significant increase in downscaling, i.e. a decreasing slope over time for the PSD95 area of EE housed mice (Fig.4E,F) but lower downscaling over time in Ctr house mice (Fig.4B). Interestingly, this effect was specific to PSD95 as for the spine head size the slope of the linear regression did not change over time and therefore no time dependent downscaling was observed (Fig.4A,D,F). The R^2 value of the linear regression was relatively constant over all time intervals, for spine head and PSD95 area of EE and Ctr (Fig.4C). Plotting the normalized changes as a function of the normalized spine head or PSD95 area, respectively, also showed a significant change in slope over time for the PSD95 area of EE housed mice (Fig.S5A–D). In summary, we found a significant increase in downscaling over time for PSD95 size in EE mice compared to Ctr. This pronounced dynamics more strongly restores fluctuations away from the average parameters in EE in line with a more concentrated distribution of PSD95 and spine head sizes that are also associated with EE. Comparing a silenced network with an active network *in vitro* Hazan *et al.* found a significant decrease in downscaling of synaptic sizes over time after silencing and a decrease of the R^2 value⁶. By analogy, the more pronounced downscaling we find might be explained by a higher activity level in EE mice and potentially causing the narrower size distribution of EE housed mice (Fig.2F)⁶. However, we did not observe such a time-dependent downscaling for spine head size and did not observe a change in its R^2 value such as described *in vitro*⁶.

Faster loss of size correlation in EE housed mice

Next, we asked how spine size change correlated over time. Do growing spines continue to grow and do shrinking spines continue to shrink? Thus, we computed the auto-correlation of the spine head and PSD95 areas over all time points (Fig.4G,H). Both the spine head areas and the PSD95 areas showed positive correlation with a large offset for both EE and Ctr housed mice. This indicates that large spine heads tend to stay rather large and smaller ones small over the observation time of up to 120min. However, at all time intervals the correlation was significantly smaller for EE housed mice. In other words, the spine size of EE housed mice is less similar after this time. This suggests that the spine population and attached PSD95 are able to change and adapt faster in EE housed mice. To analyze the coupling between spine head and PSD95 area changes, we computed the cross-correlation between these measures. Fig.4I shows that the cross-correlation was also lower in EE housed mice ($\Delta t=0$) and we did not observe directional changes of these parameters. In particular, we did not observe that PSD95 increase would systematically precede a subsequent spine head increase or vice versa as both would result in an asymmetric cross-correlation.

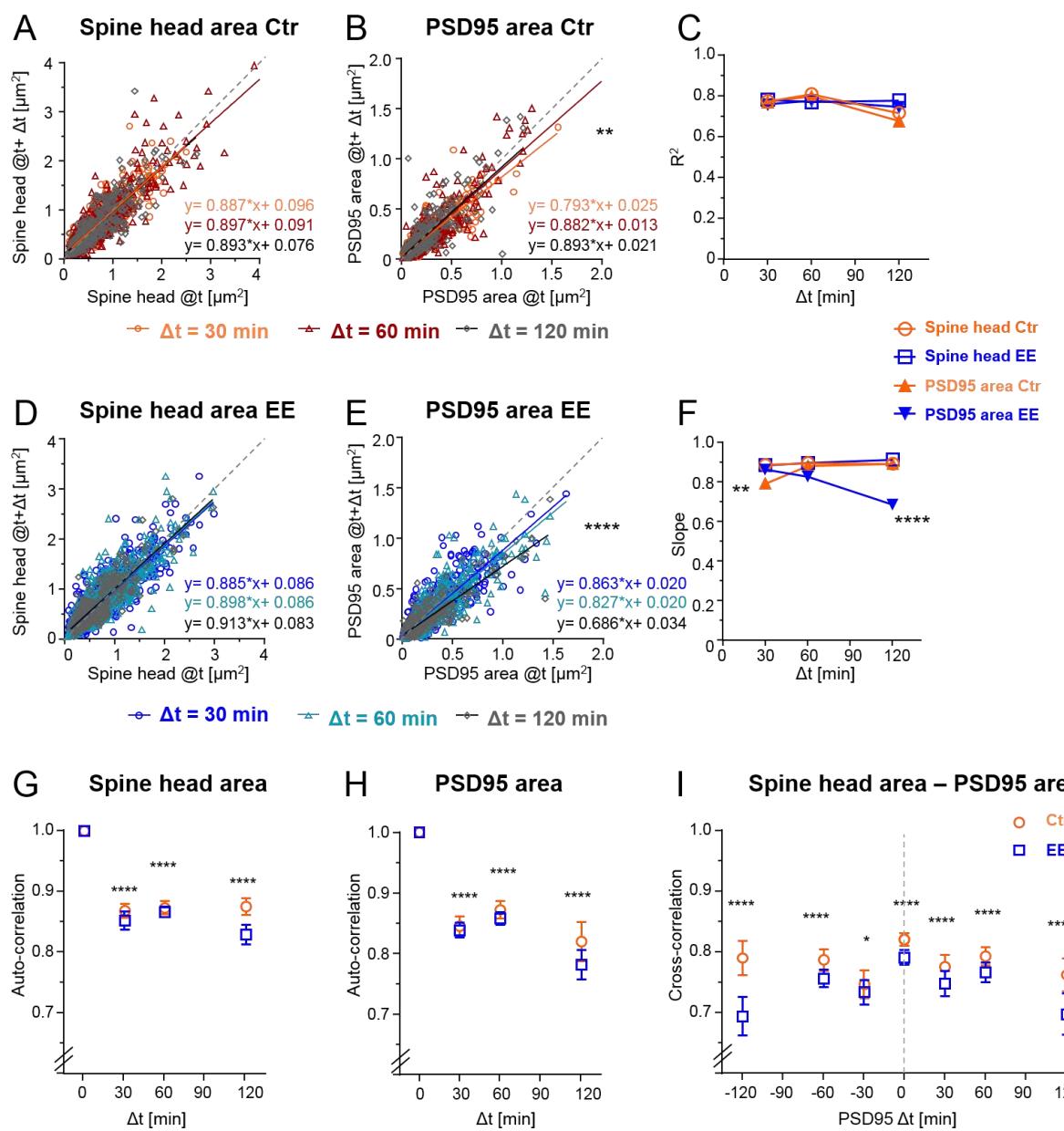


Figure 4: EE housed mice show an increase in multiplicative downscaling of PSD95 area and a lower correlation in spine head and PSD95 area over time and between these parameters. (A–F) Spine head and PSD95 area after different time intervals as function of area at t. Spine head area at time point t and after time intervals Δt of 30 min, 60 min and 120 min for Ctr (A) and EE housed mice (D). Analogous plots of PSD95 area for Ctr (B) and EE housed mice (E). Solid lines shows linear regression fits of the displayed equation; the identity line is dashed (A, B, D, E). R^2 value (C) and slope (F) of linear regressions (A, B, D, E) for different time intervals Δt . (Analysis of Covariance, Slopes equal? A: $p=0.95$, B: $**p = 0.005$, D: $p = 0.64$, E: $****p < 0.0001$). (G–I) Auto-correlation of spine head (G) and PSD95 area (H) and cross-correlation (I) of these parameters for different time intervals Δt (Unpaired t-test with Welch's correction, $*p < 0.05$, $****p < 0.0001$). Error bars are bootstrap SD. (A–I) Same data as Fig. 3.

PSD95 nanoorganization changes faster in EE mice

Previous works with EM have shown that PSDs on large spines, often called mushroom spines, are frequently perforated²⁵. In a previous publication³⁰, we investigated PSD95 morphology in homozygous

knock-in mice and showed that PSD95 assemblies on large spines were also often perforated and appeared ring-like or clustered. This nanoorganization was highly dynamic and changed in shape within a few hours. Now, we asked whether these changes would be different for mice housed in EE. Therefore, we selected perforated PSD95 assemblies and investigated their morphological changes analogously to the study by Wegner *et al.*³⁰. Fig.5A shows examples of two-color STED images of perforated PSD95 assemblies and their associated spine revealing temporal changes in the PSD95 nanoorganization. Morphological changes were categorized as no changes, subtle changes, or strong changes. The classification of morphological changes was performed blinded by three independent persons and the changes were analyzed with respect to their morphology at baseline t=0 (Fig.5B,C). Ctr mice showed a clear trend: the longer the measurement interval, the greater the morphological changes (Fig.5C). Within the time interval of 30–120min, the percentage of PSD95 nanoorganization showing no change decreased from ~40% to ~10%, while the percentage for strong changes nearly tripled for Ctr. For EE housed mice, we observed similar changes within all investigated time intervals; ~20% of the PSD95 assemblies did not change at all and ~80% underwent a change at all time points (Fig.5B). Although the percentage changes with respect to t=0 were similar for different time intervals, it should be noted that this does not exclude dynamic changes between the time points. Since we observe an equilibrium of changes in EE mice but not in Ctr mice, it argues for a time scale of less than 30min, i.e. our shortest observation interval, to perform a change in EE mice but approximately 30min in Ctr mice. These results show that not only the PSD95 total size but also the PSD95 nanoorganization is more dynamic in EE mice than in mice raised under control conditions within the investigated time course. Now, we asked whether this dynamic would be reflected as well in a different nanoorganization. Therefore, the nanoorganization was analyzed at t=0 and categorized into perforated (a continuous shape with a sub-structure such as ring- or horseshoe-like) or clustered. Indeed, fewer nanoorganizations of EE housed mice were perforated and a larger fraction was showing three or more clusters compared to Ctr mice (Fig.5D). This suggest that the PSD95 nanoorganization might play a substantial role in synaptic remodelling and is specifically shaped by experience or activity.

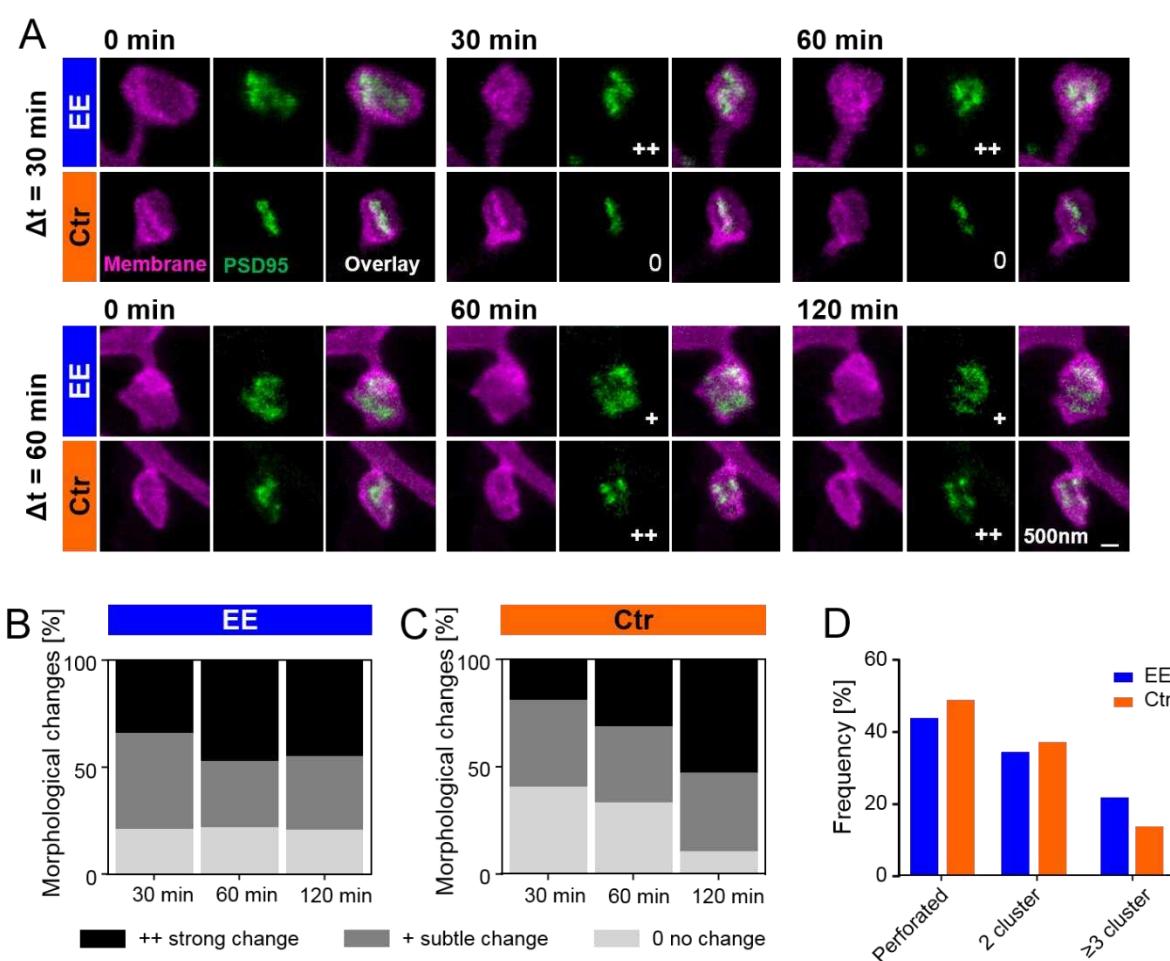


Figure 5: PSD95 nanoorganization is different between EE and Ctr and changes faster for EE housed mice. (A) Sections of two-color STED images (smoothed, MIP) of EE and Ctr housed mice at the indicated time points showing the spine membrane (magenta) and PSD95 (green). Morphological change is indicated with: 0 = no change; + = subtle change; ++ = strong change. **(B, C)** Stacked histogram of the relative frequency of morphological changes in PSD95 nanoorganization in EE (B) and Ctr (C) housed mice; all changes refer to $t = 0$ min. **(D)** Morphometry of PSD95 nanoorganization of EE and Ctr housed mice at $t=0$. Number of analyzed PSD95 assemblies in EE: 30 min: n = 38, 60 min: n = 68, 120 min: n = 29; Ctr: 30 min: n = 27, 60 min: n = 48, 120 min: n = 38.

DISCUSSION

Interactions with the environment play a key role in restructuring and refining the neuronal circuitry in the brain. Previous work has shown that an enriched environment results in structural as well as functional changes of synapses^{2,18,20,42} and stimulates synaptic plasticity in the mouse visual cortex³³. However, most studies were performed *post mortem* or *ex vivo* in slices. For the first time we here used *in vivo* STED microscopy, a superresolution light microscopy technique with nanoscale resolution, to perform an analysis of synaptic morphology as well as its nanoplasticity in the intact brain of EE housed and control mice. We designed a two-color, virtually crosstalk free *in vivo* STED microscope with a resolution of 70–80nm for simultaneous imaging of spine morphology and PSD95 in the visual cortex of anesthetized mice. We found a significantly smaller variance in size of the spine head and of

the post-synaptic density in EE mice while their average sizes were similar to Ctr mice. Both parameters were highly volatile and we observed a growth of >23% and shrinkage of >17% for spine heads and PSD95 assemblies within 30min. About 2/3 of these changes were positively correlated and ~1/3 showed a negative correlation; the amplitude of changes did not differ clearly between EE and Ctr. All parameters exhibited multiplicative downscaling, which is an indication for multiplicative changes. PSD95 was more dynamic in EE mice as denoted by an increase in multiplicative downscaling over time, faster decrease in size correlation over time and faster changes of the nanoorganization.

Two-color STED nanoscopy of synaptic nanoorganization

Previous studies mainly used two-photon microscopy to study synapse and spine plasticity *in vivo* and *in vitro*^{13,14,16,43}. Due to its limited resolution of 300–500nm, spine brightness is often taken as a measure of size^{12,16,23}. This is not very accurate since the brightness can be influenced by many factors such as expression level of the fluorescence label, the excitation laser intensity, group velocity dispersion or scattering in the tissue. Therefore, spine brightness is typically normalized to the brightness of the dendrite, making comparative studies and determination of absolute sizes difficult. Two-photon microscopy can be combined with STED to achieve nanoscale resolution^{44,45}. However, a drawback of this approach is the relatively high crosstalk when using two colors. The excitation and emission of the fluorescent labels must be spectrally close to each other in order to use the same two-photon excitation and stimulated emission laser for both fluorophores⁴⁵. Thus for EGFP and EYFP that are typically used, the reported crosstalk is as high as 92% and 27%, respectively, requiring linear unmixing²¹. Unmixing works well for structures which are relatively bright and spatially separated such as for volume labeled spines and astrocytes⁴⁵ but can produce artefacts in dark and overlapping structures. Here, we performed STED microscopy of an endogenously tagged synaptic protein, which has a considerably lower brightness than a volume label and spatially overlaps with the second label, the spine membrane, and therefore unmixing would be difficult. We achieved this by designing a two-color STED microscope with two different excitation lasers (one-photon excitation) and two spectrally separated detection channels. This new approach gave us an almost negligible crosstalk of 5% for EGFP and 8% for Citrine and making unmixing obsolete.

Temporal changes of PSD95 assemblies and spine heads are largely uncorrelated

It has been shown that PSD95 overexpression promotes the maturation of glutamatergic synapses, increases spine size, and enhances AMPA receptor currents^{46,47}. Therefore, we used the transcriptionally regulated antibody-like PSD95-FingR, which efficiently labels endogenous PSD95^{31,48} and excludes overexpression artefacts. While we observed a substantial correlation between the absolute spine head and PSD95 assembly size ($R^2\sim 0.7$, Fig.2B), corroborating previous EM studies^{15,17}, we found only weak correlations between their temporal changes ($R^2\sim 0.1$, Fig.3D–F). Most of the changes were positively correlated while about ~1/3 was attributed to negatively correlated changes. LTP experiments have shown a rapid increase in spine head size and AMPA receptor current but a delayed increase of PSD95 and other scaffolding proteins of ~60min^{13,14}. Although these experiments were performed in cell culture and with overexpressions, they indicate that spine and PSD changes might not be temporally concerted and therefore might be regulated differently^{10,11}. These results are thus congruent with the largely uncorrelated changes we observed at baseline *in vivo* at similar time scales of 30–120min. However, we also found no directional changes that would demonstrate the PSD95 size to systematically follow an increase in spine head size, as the cross-correlation is symmetric (Fig.4I). Nevertheless, why is the time-line of AMPA receptor incorporation and PSD95 increase so different although AMPA receptors are linked to PSD95 by TARP_s? At the moment different hypotheses are discussed. There is evidence that LTP promotes the exocytosis of AMPA receptor containing vesicles and thus increases the number of receptors at the synapse¹⁰. On the other hand, a slot model

proposes that the actin polymerization during LTP increases the number of slots which can harbor AMPA receptors¹⁰. Both hypotheses are in line with our observations. However, we found that spine heads and PSD95 area changed by about the same percentage of >23% already after 30min. Meyer *et al.* reported a three times larger increase in spine volume than PSD95 volume after LTP and only ~19% of change in PSD95 volume after one hour¹⁴. While the overexpression of PSD95 might have influenced the results by Meyer *et al.* we observe a PSD95 baseline volatility which is in the range of LTP driven changes.

Synapses of EE housed mice are sharper defined in size and PSD95 assemblies change more rapidly

We next compared adult mice housed in standard cages after the critical period with age matched EE housed mice and analyzed synaptic morphology and plasticity. All histograms of spine head and PSD95 area were positively skewed and log-normally distributed. While the sizes were similar between EE and Ctr housed mice, their variances were different. Both distributions, spine head and PSD95 area, were significantly narrower or sharper in EE housed mice. At the moment different models of how neuronal networks adapt to changes in activity to undergo homeostatic plasticity are considered⁴⁹. In the model of synaptic scaling, an increase in activity leads to a downscaling of excitatory synapses. The downscaling is multiplicative and therefore changes affect the average as well as broadening of the size distribution as shown, for example, in a silenced network⁶. *In vivo* and with EE, however, we observed no change in average size and therefore not pure synaptic scaling. This is in line with recent *in vivo* observations that in the adult cortex homeostatic plasticity is rather input-specific and not multiplicative for the whole ensemble of synapses⁴⁹.

Changes of spine heads and PSD95 assemblies correlated with their absolute size for EE and Ctr house mice. Small spines tended to grow and large spines tended to shrink. Such morphological changes can be modeled by a Kesten process including additive and multiplicative changes. This model entails a time-dependent multiplicative downscaling factor⁶ corresponding to the slope in Fig.4A–D. Silencing a neuronal network not only resulted in an average size increase and broader distribution but also in a weaker multiplicative downscaling⁶. Our observations are in line with these results in such that the increase in activity by enrichment strengthened the multiplicative downscaling over time for PSD95. This in turn explains the narrower size distribution. However, we did not observe these changes between EE and Ctr for the spine head. We note in passing that it should not be surprising that our *in vivo* experiments do not completely match expectations from artificially silenced neuronal culture experiment. As discussed above, there is evidence that homeostatic plasticity is not global but input-specific⁴⁹.

Enhanced nanoplasticity by experience

With STED nanoscopy we often observed a perforated nanoorganization of PSD95 undetectable by two-photon imaging. In EE housed mice, the nanoorganization of PSD95 was different and more dynamic. Accumulating evidence indicate that pre- and postsynaptic structures are clustered. The functional consequence of this clustering, however, are not fully understood. An emerging view is that these clusters are trans-synaptically aligned in ‘nanocolumns’ which are organized by activity⁵⁰. Interestingly, this implies that a reorganization of receptors might alter synaptic strength – independently of changes in the amount of receptors⁵⁰. Our finding of an increased structural plasticity of the PSD95 nanoorganization but not increased average size in EE mice might therefore reflect an increase of synaptic plasticity^{42,51}.

Outlook

Our results demonstrate that two-color STED microscopy opens novel avenues for the *in vivo* investigation of the synaptic nanoorganization and its dynamics in the living brain. Our two-color STED

approach can be extended in the future to study the dynamics of further synaptic supramolecular complexes and uncover the plasticity of nanocolumns after experience, as well as over longer time-intervals in chronic experiments and different brain regions. The crosstalk free two-color imaging introduced here will be critical to obtain unequivocal observations on the dynamics of synaptic molecular architecture in the living brain.

MATERIALS AND METHODS

DNA constructs

The transcriptionally regulated antibody-like protein to label PSD95 was obtained via plasmid pAAV-ZFN-hSyn-DIO-PSD95-FingR-Citrine-CCR5TC, with a double-floxed inverted open reading frame (DIO). It was cloned sequentially by creating the following plasmids. 1) pAAV-hSyn-DIO-EYFP: PCR based insertion of the human Synapsin promoter (hSyn) instead of the human elongation factor-1 alpha (Ef1a) promoter into the vector backbone of pAAV-Ef1a-DIO-EYFP, kindly provided by the laboratory of Dr. K. Deisseroth of Stanford University, CA. 2) pAAV-ZFN-hSyn-DIO-EYFP: Insertion of the zinc finger binding motif (ZFN)³¹ 5' to the hSyn promoter by hybridization of two 5' phosphorylated primers. 3) pAAV-ZFN-hSyn-DIO-PSD95-FingR-Citrine-CCR5TC: PCR based insertion of the antibody like PSD95-FingR (FingR: Fibronectin intrabodies generated with mRNA display)³¹, the transcriptionally regulatory elements CCR5ZFL-KRAB(A) (CCR5ZFL: left zinc finger from an engineered zinc-finger nuclease against the gene CCR5; KRAB(A): transcriptional repressor domain from rat Kid-1³¹), and Citrine to obtain the final plasmid. The sequences of PSD95-FingR and CCR5ZFL-KRAB(A) were amplified from pCAG_PSD95.FingR-eGFP-CCR5TC (gift from Dr. Don Arnold (Addgene plasmid # 46295)). All primers and restriction endonucleases used for cloning are listed in Table 1.

The plasmid pAAV-hSyn-DIO-myr-EGFP-LDLR(Ct) for dendritic membrane labelling was cloned as follows: first, we PCR amplified a myristoylation (myr) site-attached EGFP including the myristoylation sequence ATGGGCTGTGTGCAATGTAAGGATAAGAAGCAACAAACTGACG in the forward primer. Second, we split the C-terminal (Ct) cytoplasmic domains of low density lipoprotein receptor (LDLR; gb: AF425607, amino acid residues 813–862)³⁸, finally designated as LDLR(Ct), into two parts (part 1 and part 2), designed 5' phosphorylated forward and reverse primers for each part and hybridized each pair (Table 1). In the third and final step, the endonuclease-digested myr-EGFP PCR together with the LDLR(Ct) part1 and LDLR(Ct) part 2 were ligated into plasmid pAAV-hSyn-DIO-EYFP digested with Ascl and Nhel to finally obtain pAAV-hSyn-DIO-myrEGFP-LDLR(Ct). Generation of the Cre recombinase expression plasmid pAAV-hSyn-Cre was performed as previously described⁵². To determine the crosstalk of our two-color *in vivo* STED microscope, we used two plasmids, both encoding a fusion protein consisting of the small peptide Lifeact (LA), that directly binds to F-actin, and a fluorescent protein: pAAV-hSyn-LA-EYFP³⁴ and pAAV-hSyn-LA-EGFP. pAAV-hSyn-LA-EGFP was generated by the exchange of the fluorescent protein EYFP with EGFP. All generated or used AAVs contain as the 3' untranslated regions the woodchuck hepatitis virus posttranslational regulatory element (WPRE) and the human growth hormone polyadenylation site.

Table 1: Overview of the primers and endonucleases used to clone new plasmids. a: produced by PCR, b: generated by hybridization, P: phosphorylated; underlined nucleotides: restriction sites or part of them

Name of the plasmid to be generated	Primers	Restriction sites	DNA-insert
pAAV-hSyn-DIO-EYFP	5'- ccgtacgcgtgtcttagactgcagaggcccctg -3'	MluI	hSyn ^a
	5'- gcatgcacggatccggtaccctctgactgcgc -3'	SalI	
pAAV-ZFN-hSyn-DIO-EYFP	P-5'- <u>ccgtgtcatccatcggt</u> -3'	MluI + XbaI	ZFN ^b
	P-5'- <u>ctagacacgatgaggatgaca</u> -3'	MluI + XbaI	
pAAV-ZFN-hSyn-DIO-PSD95-FingR-Citrine-CCR5TC	5'- gacaccatggatctactggagccgtacc -3'	NcoI	PSD95-FingR ^a
	5'- tccgt <u>ctagcatgc</u> taaggcatagaaggaa -3'	NheI	
	5'- atcgg <u>cgcccta</u> agecatagaaggaa -3'	Ascl	CCR5ZFL-KRAB(A) ^a
	5'- gtc <u>ggccggagcgggtgc</u> aggggc -3'	KasI	
	5'- agegg <u>ccagtc</u> cccttagactgtacagctgc -3'	KasI	Citrine ^a
	5'- gatccatggtagcaaggcgaggagctgtcac -3'	NcoI	
pAAV-hSyn-LA-EGFP	5'- gat <u>ccatggtagcaaggcgaggagctgtcac</u> -3'	NcoI	EGFP ^a
	5'- gegcc <u>gaattcttacttgtacagctcgccatgc</u> -3'	EcoRI	
pAAV-hSyn-DIO-myrEGFP-LDLR(Ct)	5'- agttat <u>ctagcatggctgtgtcaatgt</u> aaaggataag aagcaacaaaactgc <u>atgtggtagcaaggcgaggag</u> -3'	NheI	Myristol (myr)-EGFP ^a
	5'- cgc <u>accggtttgc</u> atcg <u>ctcgccatg</u> -3'	AgeI	
	P-5'- <u>ccggcggaa</u> ctggcg <u>ctgaagaat</u> atcaacagc atcaatt <u>tcgataacccgtgtaccagaagaccacagaggat</u> -3'	AgeI	LDLR(Ct)-part1 ^b
	P-5'-cag <u>tcatctgtggcttctgt</u> acacgggttatcgaaa ttgat <u>gtgttatcttcaggcgccatcg</u> -3'	AgeI	LDLR(Ct)-part1 ^b
	P-5'-gag <u>ctcacattgcaggccaa</u> gacgggtacacccatcc aagt <u>ccgcagatggtcagccgtggacgtggctgagg</u> -3'	Ascl	LDLR(Ct)-part2 ^b
	P-5'- <u>cgcgcctcaggccacatgtccgtggacgtaccatgtcc</u> gacttggat <u>agggtgtacccgttgggacactcaatgt</u> -3'	Ascl	LDLR(Ct)-part2 ^b

Virus production

Recombinant rAAV particles with mixed serotype 1 and 2 of pAAV plasmids encoding the proteins of interest were produced in HEK293-FT cells. The entire procedure was previously described in detail⁵² and is applied here with the following modifications: After DNaseI treatment (30 min at 37 °C), the suspension was centrifuged at 1200 g for 10 min. The supernatant was filtered through an 0.45 µm sterile filter (Merck/Millipore, Darmstadt, Germany) and first applied to an Amicon Ultra-15, MWCO 100 kDa, centrifugal filter unit (Merck/Millipore, Darmstadt, Germany), followed by a Vivaspin® 500, MWCO 100 kDa, centrifugal concentrator (Sartorius, Göttingen, Germany) to wash the virus in Opti-MEM™ Medium (ThermoFisher Scientific, Darmstadt, Germany) and concentrate to a final volume of 150 µl.

Animals

All animal experiments were performed with C57BL/6J female mice reared at the animal facility of the Max Planck Institute of Experimental Medicine in Göttingen and housed with a 12-h light/dark cycle, with food and water available *ad libitum*. Experiments were performed according to the guidelines of the national law regarding animal protection procedures and were approved by the responsible

authorities, the Niedersächsisches Landesamt für Verbraucherschutz (LAVES, identification number 33.9-42502-04-14/1463). All efforts were made to minimize animal suffering and the number of animals used.

Housing conditions

Environmentally enriched (EE) animals were born and raised in the commercially available Marlau™ cage (Marlau™, Viewpoint, Lyon, France)⁵³, 580 x 400 x 320 mm in size including 2 floors, providing an extensive exploration area (Fig. S2). Three pregnant female mice were placed in an EE cage about 1 week before delivery. Pups were weaned and split by sex at post-natal day 30. The ground floor of the EE cage consisted of two separate compartments: the smaller part contained food and the larger part contained water, running wheels and a red house. To get food, mice had to use a stairway to reach the second floor where a maze was placed. After passing the maze, mice slided through a tunnel back to the ground floor, directly into the smaller compartment and had free access to food. By passing a one-way door they could enter the larger part of the ground floor to get water. To increase novelty and maintaining cognitive stimulation, the maze was changed 3 times per week with a total of 12 different configurations. Control mice (Ctr) were born and raised in standard cages of 365 x 207 x 140 mm in size. They were kept in 2–3 animals per cage, which were solely equipped with nesting material.

AAV transduction

Adult (>12 weeks) female C57BL/6J mice were stereotactically transduced with a mixture of rAAV with mixed serotype 1 and 2. rAAV1/2-ZFN-hSyn-DIO-PSD95-FingR-Citrine-CCR5TC, rAAV1/2-hSyn-Cre and rAAV1/2-hSyn-DIO-myEGFP-LDLR(Ct) were stereotactically injected at the same time into the visual cortex of the left hemisphere at a depth of ~ 500 µm below the pia transducing mainly layer 5 pyramidal neurons by using the following coordinates: 3.4 mm posterior to the bregma, 2.2 mm lateral to the midline, at an angle of 70° to the vertical, and with a feed forward of ~750 µm. Anesthesia was initiated by intraperitoneal injection of 0.05 mg/kg fentanyl, 5 mg/kg midazolam, and 0.5 mg/kg medetomidin (MMF). The mouse was head fixed with a stereotaxic frame and placed on a heating pad throughout the whole procedure to maintain a constant body temperature. The depth of the anesthesia was controlled by monitoring the pulse rate and O₂ saturation with a pulse oximeter at the thigh of the mouse and body temperature with a rectal temperature probe. A gas mixture with a high O₂ (47.5 vol%) and CO₂ (2.5 vol%) content was applied to the mouse's nose, significantly increasing the oxygenation. The eyes of the mouse were covered with eye ointment and the head was disinfected with 70% ethanol. The skin was cut by a ~ 0.5 cm long incision and a drop of local anesthetic (0.2 mg mepivacaine) was applied. Then, a 0.5 mm hole was drilled into the skull. 150 nl of rAAV containing solution was pressure injected through a glass micropipette attached to a microinjector (Picospritzer® III, Parker Hannifin Corp, Cleveland, Ohio) with an injection rate of ~50 nl per minute. After injection, the pipette was kept at the target location for an additional 2 minutes to allow the virus to disperse away. After retracting the micropipette, the incision was closed with a suture. Anesthesia was then antagonized by intraperitoneal administration of 0.1 mg/kg buprenorphine and 2.5 mg/kg atipamezole. The mouse was kept in a separate cage until full recovery and then put back into its original cage, and group-housed in the animal facility until the final experiment.

Craniotomy for *in vivo* STED microscopy

Three to six weeks after viral transduction, a craniotomy was performed as described previously³⁹. In brief, the mouse was anaesthetized with MMF, placed on a heating pad and vital functions and depth of anesthesia were controlled throughout the final experiment as described above. The head was mounted in a stereotaxic frame. The scalp was removed and a flat head bar was glued to the right hemisphere to leave enough space for the cranial window above the visual cortex, rostral to the former

viral injection site. After drilling a circular groove (\sim 2–3 mm) into the skull, the bony plate was carefully removed without causing a trauma. The remaining dura and arachnoid mater were carefully removed with a fine forceps. A small tube was positioned at the edge of the opening to drain excess cerebrospinal fluid if necessary. The craniotomy was sealed with a 6 mm diameter cover glass glued to the skull. The mouse was mounted on a tiltable plate which can be aligned perpendicular to the optical axis of the microscope in a quick and easy routine before being placed under the microscope ³⁹.

Culture of primary hippocampal neurons, transduction and imaging

Primary cultures of rat hippocampal neurons were prepared from P0-P1 Wistar rats (RjHan:WI; The Jackson Laboratory, Bar Harbor, ME) of both sexes according to ⁵⁴. Neurons were cultured at 37 °C in a humidified atmosphere with 5% CO₂ and transduced between 8 and 10 days *in vitro* (DIV) with the respective rAAVs. After an incubation time of 7 days, live cell STED imaging was performed at room temperature.

***In vivo* two-color STED microscope**

A previously described custom-designed STED microscope ^{30,34} was modified to accommodate for virtually crosstalk free two-color imaging as follows (Fig. 1). Excitation light (Exc1) provided by a pulsed laser diode emitting blue light at 483 nm (PiLas, Advanced Laser Diode Systems, Berlin, Germany) was complemented by a second excitation beam (Exc2). The beam of a Ti:Sapphire laser (MaiTai; Spectra-Physics, Santa Clara, CA) was split into two, pumping an optical parametric oscillator (OPO; APE, Berlin, Germany) emitting 80 MHz pulses at 595 nm for stimulated emission depletion (STED) and a supercontinuum device (FemtoWHITE800, NKT photonics, Birkerød, Denmark) generating white light. The white light was spectrally filtered for green light with a bandpass filter (BP1, BrightLine HC 520/5, Semrock, IDEX Health & Science, Rochester, NY) for selective excitation of EYFP or Citrine at 520 nm (Exc2, Fig. 1A,B). Exc1, Exc2 and the STED beam were co-aligned with dichroic mirrors. After passing a Yanus scan head (Till Photonics-FEI, Gräfelfing, Germany) consisting of two galvanometric mirrors for x-/y-scanning and relay optics, the three beams were focused by a 1.3 numerical aperture objective lens (PL APO, 63x, glycerol; Leica, Wetzlar, Germany). Additionally, the STED beam was passing a vortex phase plate (VPP; RPC Photonics, Rochester, NY) to create a doughnut shaped focal intensity pattern featuring zero intensity in the center. Temporal overlap of all three pulsed laser beams was achieved electronically by synchronizing the blue laser diode, Exc1, to the MaiTai and optically by an optical delay line for Exc2. Z-scanning was performed by moving the objective with a piezo (MIPOS 100PL; piezosystem jena GmbH, Jena, Germany). The back-projected fluorescence light was split at 515 nm with a dichroic mirror (DM2, ZT502rdc-UF3; Chroma Technology Corporation, Bellow Falls, VT) into two beams. The shorter wavelength was reflected, filtered by a bandpass filter (BP3, BrightLine HC 504/12; Semrock) and focused onto a multimode fibre for confocal detection connected to an avalanche photodiode (APD, Excelitas, Waltham, MA). The transmitted, longer wavelength fluorescence light was also filtered with a bandpass filter (BP2, H544/23, AHF analysentechnik, Tübingen, Germany) and detected with an APD, respectively.

***In vivo* two-color STED imaging of anesthetized mice**

The *in vivo* STED microscopy was performed in layer 1 of the visual cortex at a depth of 5–20 μ m below the pia. Spherical aberrations due to the tissue penetration were corrected on the first order by adapting the correction collar of the glycerol immersion objective for the best image quality at each field of view. Both excitation colors were alternated line-by-line; i.e. a line was recorded by excitation with blue light (Exc1) and then the same line was recorded again with green excitation light (Exc2). Potential drift between images and movement of the spine or PSD95 was therefore negligible. For imaging we picked FOVs with dendrites parallel to the focal plane. After recording a two-color STED

image stack, the position of the motorized micrometer stage (MS-2000, Applied scientific instruments, Eugene, OR) was noted and an overview image taken. This process was repeated several times for different positions. After 30 min the micrometer stage was moved back to the first position. The position was confirmed by recording coarse overview images of the dendrite and thereby adjusting the z-position to the right depth. STED images were recorded as z-stacks of ~20–40 μm in x and y with 500 nm axial steps at different positions of the cranial window. All positions were repeatedly imaged 3–4 times at intervals of 30 min to 2 hours. Therefore, some dendritic regions were investigated at the time points 0 min, 30 min and 60 min, while other dendritic region within the cranial window were examined at the time points 0 min, 60 min and 120 min or even at 0 min, 120 min and 240 min. The benefit of a membrane label over the often used volume label is that the brightness of the spines and dendrites is similar whereas with a volume label spine heads and dendrites often outshine the small spine neck; thus, either the much darker neck is not visible between the bright head and dendrite or the detector is saturated at the bright heads and dendrites.

All images were recorded with a pixel dwell time of 5 μs , a pixel size of 30 \times 30 nm in x and y, and z-stacks of 500 nm step size. Blue and green excitation power was 4.5 μW , respectively, in the back aperture of the objective. The average STED power in the back aperture was 37–45 mW for static images and 15 mW for time-lapse imaging.

Data processing and analysis

Confocal and STED images were acquired by the software Imspector (Abberior Instruments, Göttingen, Germany). Analysis of size, shape, and brightness was performed manually in Fiji⁵⁵. For spine head and PSD95 analysis, the first and last plane were omitted to ensure that the spine (or PSD95, respectively) was located completely in the focal plane. We only analyzed spines that also bore a PSD and we only analyzed spines that extended from the dendrite parallel to the focal plane. PSDs directly on the dendrite most likely representing shaft synapses and spines pointing upwards or downwards that could not be clearly resolved were not analyzed. First, images of each channel were processed as follows (Fiji commands shown in capital letters): 1. Smoothing: PROCESS > SMOOTH once. 2. Brightness adjustment: IMAGE > ADJUST > BRIGHTNESS/CONTRAST set minimum to 1 instead of zero. 3. Overlay both one-color images: IMAGE > COLOR > MERGE CHANNELS. 4. Open ROI manager: ANALYSE > TOOLS > ROI MANAGER. To determine the length (a) and width (b) of PSD95 and the spine head, a line from the top to the bottom, or from left to right as shown in Fig. 2A was drawn (from tip to tip of the corresponding arrowheads) and saved to the ROI manager. For PSD95 assemblies showing a perforation such as shown in Fig. 2A, we took the maximal length and width to cover the whole assembly. From these values, the area (A) was estimated by computing the area of an ellipse by $A=\pi*a/2*b/2$. For the analysis of brightness, PSD95 images were processed as described above. Using the ELLIPTICAL SELECTION tool, each PSD95 assembly was encircled and the brightness was displayed by the variable "RawIntDen", which is the sum of the intensity values of all pixels in the selected area. Spine density was obtained by dividing the total number of spines minus one by the length of the parent dendrite between the first and last spine. The length was measured with the FREEHAND LINE tool, in Z PROJECTION (maximum intensity) images. Branched spines were counted once.

Absolute changes of spine head area or PSD95 area between two time points, t and $t+1$, were calculated by $\Delta A_{\text{abs}}=A(t+1)-A(t)$. 'A' denotes the spine head area or the PSD area, respectively. Normalized relative changes of spine head area or PSD area were computed by $\Delta A_{\text{norm}}=(A(t+1)-A(t))/(A(t+1)+A(t))$ and percentage changes by $\Delta A\%=(A(t+1)-A(t))/A(t)*100$.

Data were further analyzed with MATLAB[®] (MathWorks[®], Natick, MA). The principal component analysis (PCA) was performed with the built in MATLAB function 'princomp'.

The correlation function was computed for different time intervals Δt by

$$\text{Corr}(\Delta t) = \langle \delta A_1(t+\Delta t) * \delta A_2(t) \rangle / \sqrt{\langle \delta A_1(t+\Delta t)^2 \rangle * \langle \delta A_2(t)^2 \rangle},$$

with δ denoting the fluctuation of the mean: $\delta A(t) = A(t) - \langle A(t) \rangle$. The bracket $\langle \dots \rangle$ denotes the average over time and over the empirical distribution of spines or PSDs, respectively. The auto-correlation was computed with $A_1 = A_2$ either for spine head area or PSD95 area. The cross-correlation was computed with A_1 denoting the spine head area and A_2 the PSD95 area.

Nanoorganization and morphological changes of PSD95

All non-macular PSD95 nanoorganizations were selected. Their shape was categorized by 3 blinded scientists into perforated or clustered shapes. Perforated PSDs were of continuous shape but complex such as a ring or horseshoe-like or more twisted. Clustered nanoorganizations were characterized by two or more separated assemblies of PSD95 per spine and the number of clusters was counted. Temporal changes in PSD95 morphology were analyzed similarly. Assemblies which did not alter their overall morphology or number of clusters were marked as 'no change'. Assemblies showing minor modifications, such as a small movement of a sub-cluster, were classified as 'subtle change'. Changes of the overall morphology such as a smooth continuous PSD falling apart into different clusters were characterized as 'strong change'. All changes were referred to time point $t = 0$ min.

Statistical analysis

Statistical analysis was performed in GraphPad Prism (version 7.04, Graphpad Software, San Diego, CA). Positively skewed data sets were compared by the Kolmogorov Smirnov test. Log-transformed data and normalized relative changes were normally distributed and compared by an unpaired t -test with Welch's correction. The number of analysed mice and spines, p-value, and the specific statistical test performed for each experiment are included in the appropriate figure legend. All tests were applied two-sided where applicable. Probabilities are symbolized by asterisks as follows: * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$, **** $p < 0.0001$

Materials availability

This study did not generate new unique reagents.

Data and materials availability

Image data files are provided by reasonable request through the corresponding author.

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Author contributions

Conceptualization (WW, KIW); STED microscope design and construction (KIW); Cloning and virus production (WW, CG); Surgery (HS, WW); Imaging (WW, KIW); Analysis (WW, KIW, FW); Writing (WW, KIW).

Declaration of interest

The authors declare no competing interests.

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