### 1 Supporting information for

- 2 Quantum Dot conjugated nanobodies for multiplex imaging of protein dynamics at
- 3 synapses

### 4 Materials and methods

### 5 Quantum Dot functionalization with nanobody

6 Qdot 605 ITK<sup>™</sup> Amino (PEG) Quantum Dots were purchased from Life Technologies.

37 µl of QD605 was mixed with 13 µl 1M Borate buffer, pH 8, 0.5 mg 7 Bis[sulfosuccinimidyl] suberate (BS3) (Sigma Aldrich) and incubated for 30 min at RT with 8 9 mild vortex. This mixture then loaded onto a Nap-5 column (Life technologies) and QD-BS3 conjugate was eluted with 1×PBS pH 7.4. First 500 µl coloured solution was collected and 10 concentrated to 50 µl using a 50 kDa molecular weight cut-off filter (Life Technologies). 100 11 12 µl anti-GFP nanotrap (nanobody hence after) (ChromoTek GmbH) was added to the QD conjugate and incubated for 2h at RT with mild vortex. QD and nanobody mixture then 13 14 diluted to 400 µl and loaded on to a 50 kDa molecular weight cut-off filter and centrifuged at 15  $4000 \times g$  for 8 min. Flow through was discarded and supernatant was diluted to 400 µl and centrifuged again. This step was repeated 5-6 times and then the supernatant was 16 17 concentrated to 50 µl. Finally unreacted BS3 sites on OD-nanobody conjugates was quenched 18 by adding 2  $\mu$ l of 1M Tris buffer pH 7.4.

#### 19 Characterization of QD-nanobody conjugates in vitro

20 QD conjugates, free QD and QD-BS3 were analysed using 1% Agarose-TBE gel. Gel was ran for 2-3 h at 100 V and visualized under UV in a gel documentation system. These 21 conjugates were further analysed with 12% SDS page and stained with coomassie brilliant 22 blue to visualize protein conjugates and free nanobodies. GFP was purified from HeLa cells 23 by transfecting them with CAG GFP followed by affinity capture using GFP trap agarose 24 beads (ChromoTek GmbH). Affinity captured GFP was eluted from bead by addition of 50 µl 25 1M glycine pH 2.5 and then quenching with 1M Tris pH 7.4. 5 µl of GFP and BSA (1 mg/ml) 26 27 was blotted in a nitrocellulose membrane and allowed to dry at RT for 30 min. The membrane was blocked with PBST (PBS+0.05% Tween-20) containing 10% Horse serum 28 29 (Life Technologies) and 1% BSA. The blot was incubated with 1 in 10000 dilution of QD-30 nanobody conjugates overnight at 4°C. The membrane was washed 3 times in PBST and developed in a UV illuminated gel documentation system. Dot blot with QD-nanobody 31 32 conjugates and free QD was performed similarly with minor modifications. For labelling, 1 in 33 5000 dilution of QD-nanobody conjugates and 1 in 2000 dilution of free QD was used.

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### 35 Cell culture, transfection, labeling and staining

HeLa cells were cultured in DMEM medium (Gibco) supplemented with streptomycin (100  $\mu$ g/ml), penicillin (100 U/ml), and 10% fetal bovine serum on 13 mm glass coverslips in a 6 cm cell culture dishes. Cells were transfected with ~5  $\mu$ g of GFP-NrCAM or GFP-AnkG

using nucleofection (Amaxa, Lonza AG). After 24h post transfection, cells were labelled with 1 2 QD-nanobody conjugates (1 in 3000 dilution) for 5 min at RT in PBS containing 10% Horse 3 serum. Coverslips were washed 3 times by dipping in PBS and imaged in a widefield set up based on Olympus BX51WI microscope with a 60× Olympus objective coupled to an EM-4 CCD camera (Andor) and a perfusion chamber as described in.<sup>1</sup> Images were acquired at 8.4 5 Hz for 20-30s and analysed in ImageJ (NIH). 6 7 Rat hippocampal cultures were performed as described earlier and transfected on the day of dissection using nucleofection with 10 µg plasmid DNA either of GFP-NrCAM (A gift from 8 Dargent Lab) or with 5µg of α2-SEP (Gift from S. Moss) according to manufacturer's 9 protocol.<sup>1</sup> 2 days post transfection with GFP-NrCAM, neurons were labelled with 1 in 1000 10 dilution of QD-nanobody conjugates for 5-7 minutes and washed 2 times with PBS and 11 imaged as described above using 405 illuminations of QDs.<sup>1</sup> Growth cones were imaged after 12 identifying axon initial segment that generally have higher GFP expression. For depletion of 13 F-Actin network, neurons were treated with 39 µM Cytochalasin-D (Sigma) in their original 14 culture media for 30 min at 37°C and then labelled with QD-nanobody conjugates containing 15 39 µM Cytochalasin-D at RT for 5 min.<sup>2</sup> Cells were washed twice in PBS and imaged with 16 continuous perfusion of imaging media containing 1 μM Cytochalasin D. α2-SEP expressing 17 neurons were imaged 14 day post nucleofection after labelling with QDs (1 in 3000 dilution) 18 19 as described above.

### 20 Organotypic slice culture, transfection and labeling

Organotypic hippocampal slice cultures were made following the Stoppini interface method 21 (2), where by cultured slices are maintained on membrane insert.<sup>3</sup> Briefly, P7-10 Sprague-22 Dawley rats were sacrificed in accordance with schedule 1 Home Office protocol. The brain 23 24 was removed and placed in a petri dish containing ice-cold slicing media (EBSS + 25mM 25 HEPES). 300 µm sagittal slices were made on a vibratome (Leica) following which the hippocampus was dissected and placed on the membrane. Slices were cultured in an 26 27 incubator containing 5% CO<sub>2</sub> for 7 days at 37°C in a 6 well plate containing 1.6 ml of culture media (72% MEM (Gibco), 25% Horse Serum, 0.64% Glucose (Sigma), 100 U/ml Penicillin, 28 100 µg/ml Streptomycin, 6 U/ml Nystatin (Sigma)). Slices were fed with fresh culture media 29 30 every 3-5 days and transfected with a hand-held Helios gene gun (Bio-Rad) 7 days post 31 dissection. 50 µg DNA was coupled to gold microcarriers (1.6 µm diameter) by precipitation with calcium chloride according to the previous literature<sup>4,5</sup>. Finally, the gold coated 32 33 cartridges were inserted inside the gene gun and shot onto the slices with helium pulse. Slices 34 were further cultured for 7 days as described above and labelled with QDs. Slices with 35 membranes were removed from the wells and 1 in 1000 dilution of QD-nanobody conjugates 36 diluted in PBS+10% horse serum+1% BSA was added to the slices and incubated for 5-8 37 minutes unless specified, at 37°C. Slices were washed 2 times in PBS and imaged as 38 described earlier.

#### 39 Immunofluorescence and imaging

Slices were fixed with 4% paraformaldehyde containing 4% Sucrose for 10 minutes in PBS
and washed 5 times with PBS. Slices were transferred in a well of 24 well plate and

1 permeabilized with 0.2% TritronX-100 in PBS containing 10% horse serum and 1% BSA for 2 2 h at RT. Slices were then labelled with anti-GFP Rat IgG (Nacalai Tesque, 1 in 2000), anti-3 Gephyrin Mouse IgG (Syanptic System, 1 in 250) and anti-MAP2 guinea pig IgG (Synaptic Systems1 in 500) at 4°C overnight in a eppendorf thermomixer with gentle shaking. Slices 4 were washed 3 times with large volume of PBS with shaking and labelled with secondary 5 antibodies for 2 h at RT. 1 in 500 dilutions of Alexa-488 conjugated anti-Rat; Alexa-555 6 7 conjugated anti-Mouse and Alexa-647 conjugated anti-Guniea pig (Obtained from Molecular probes, Life Technologies) were used to visualize various molecular markers. Slices were 8 mounted in Vectashield H-1400 (Vector labs) and imaged in a zeiss LSM 700 confocal 9 microscope with a plan-apochromat  $63 \times$  oil-immersion lens with 1.4 numerical aperture. Z 10 stacks were acquired with 1 µm step size and then projected using maximum intensity 11 projection in ImageJ. 12

#### 13 QD Image Analysis

#### 14 QD Detection and Tracking

Quantitative analysis of QD recordings can be separated into two discrete steps: detection and 15 tracking. Accurate QD detection was achieved using an image segmentation algorithm called 16 Basic Image Features (BIFs) which had previously been employed in a SPT study.<sup>1</sup> Briefly, a 17 set of 2-D Gaussian filters is applied to each frame so that each pixel is classified as one of 18 seven different basic image features (BIFs).<sup>6,7</sup> Pixels that possess high local rotational 19 symmetry, corresponding to local fluorescence maxima, are classified and grouped 20 21 accordingly. As QDs appear as local diffraction-limited bright blurs, these BIFs pixel clusters are considered QDs. Given a set of detected QDs for each frame, trajectories were formed 22 using tracking software written in Mathematica (Wolfram Research).<sup>1</sup> The algorithm works 23 by minimising the Euclidean Distance between QD tracks across consecutive frames. It can 24 25 account for QD blinking by allowing QD tracks to not link to any specific QD in one frame, 26 only to link to a re-emergent QD in a subsequent frame. The combinatorial complexity of the 27 algorithm is assisted by the user-specified maximum distance a QD is capable of moving 28 across successive frames. Furthermore, this step limits the search for linked QDs to a 29 bounded distance (no. frames since detection × max distance). This maximum interframe distance was generally set at 2 pixels (0.55µm) per timestep (120 ms). Trajectories were 30 31 required to consist of at least 50 points to be included in the analysis. Mean square displacement (MSD) versus time (t) was calculated for each QD track using the formula: 32

$$MSD(n*dt) = (N-n)^{-1} \sum_{i=1}^{N-n} ((x_{i+n}-x_i)^2 + (y_{i+n}-y_i^3)^2)$$

where xi & yi are the Cartesian coordinates of the QD at frame i, dt is the timestep between frames, n is number of timesteps and N is total number of frames in the movie. Note that MSD (0) = 0. Instantaneous diffusion coefficients (D) were then estimated by fitting a line through the origin to the first five points of the MSD curve and using the 2D diffusion law (MSD = 4 D×t). QDs exhibiting diffusion score lower than  $5 \times 10^{-3}$  were considered immobile and removed from the subsequent analysis.<sup>6-8</sup> Similarly, trajectories that produce D > 1 were removed from the analysis. While the QD diffusion score was estimated by linear fit of the

1 first five points of the MSD, in some cases the MSD was plotted as a function of time 2 (MSDt) for a population of QDs. In these instances, the MSD was calculated for each QD at 3 each timestep for at least 20 points. Then, the values at each timestep were subsequently averaged to produce a single curve. The standard error around each averaged timestep was 4 plotted. Some OD movies were acquired alongside a single image illustrating the position of 5 synaptic puncta. These images were taken immediately before the QD movie. Synapse 6 7 position was calculated from these images using the BIFs algorithm in a similar manner to the QD sub-pixel localisation. QD position was overlaid on synapse positions to determine 8 whether trajectories were synaptic or extra-synaptic. A trajectory was considered synaptic if 9 the distance between the two centroids (QD and synapse) was less than 2 pixels.<sup>1</sup> In these 10 instances, single trajectories can return both an inside and outside diffusion score. Identical 11 diffusion thresholds were applied to each set, as before  $(0.5 \times 10^{-3} < D < 1)$ . Representative 12 QD tracks were created in ImageJ using Mosaic particle tracker 2D/3D plugin.<sup>9,10</sup> 13

We examine drift by following methods, z project all the movies to assess if significant x-y drift is present in the movies. Any drift > 2.5  $\mu$ m (i.e ~10 pixel) are discarded. Any drift smaller than that are corrected as described elsewhere using StackReg macro (Rigid body or translation algorithm) embedded in imageJ.<sup>1</sup>



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Figure S1: SDS page analysis of QD-nanobody conjugates. 12% SDS page was performed to
confirm presence of Nanobody in QD-nanobody conjugates. M: PageRuler Plus Prestained
Protein Ladder (Thermo Scientific); QD: Free Quantum dot; QD-NT: QD-nanobody
conjugate; NT: Free nanobody. Gel was run at 100 V and stained with Coomassie Brilliant
Blue R250.

- 1 SI Figure 2



Figure S2: Specificity of QD-nanobody conjugates towards GFP. HeLa cells were transfected with either GFP-NrCAM or GFP-AnkyrinG as described in the methods section. 24h post transfections, cells were labelled with a 1 in 1000 dilution of QD-nanobody conjugates for 5 min and then washed 3 times in PBS and imaged with a Zeiss inverted microscope with 63× Plan-apochromat objective having photometrics evolve CCD camera under constant perfusion. Images were acquired at a rate of 1fps and projected using maximum intensity projection method in ImageJ. It can be seen that when GFP is present at the cell membrane (GFP-NrCAM), QD-nanobody conjugates recognizes it, while in ankyrinG expressing cells where GFP is expressed in the cytoplasm, no labelling is observed by the probe. Scale bar: 5 µm 



Figure S3: QD-nanobody conjugates trajectories overlaid on rat hippocampal neurons from post-natal day 0 rat cultures expressing GABA<sub>A</sub>-SEP ( $\alpha$ 2-SEP). Transfection of  $\alpha$ 2-SEP was performed on the day of dissection using nucleofection as described in the methods section. After 12-14 days of transfection, neurons were labelled with QD-nanobody conjugates for 5 min at RT and imaged in an Olympus microscope with 60× objective coupled to an Andor CCD camera under continuous perfusion. Imaging was performed at 8.4 Hz and representative tracks were created and overlaid using a Mosaic plugin available in Image J. Scale bar: 10 µm.

#### 1 SI Figure 4



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Figure S4: QD-nanobody conjugate trajectories in rat brain slices after longer exposure with 17 QDs. Organotypic slices were transfected with α2-SEP using biolistic method and imaged 7 18 days post transfection. Slices were labelled with QD-nanobody conjugate for 30 min prior to 19 imaging at 37°C, washed 3 times and placed in a perfusion chamber attached to a wide field 20 microscope. Transfected neurons were identified using GFP filters and then QD images were 21 22 acquired at 8.4 Hz. Tracks were created in imageJ, overlaid with the GFP image and stitched using MosaicJ plugin available in ImageJ.<sup>10</sup> Note: Due to longer labelling protocol, mobile 23 QDs not only labelled complete neuron but also increases nonspecificity which is reported in 24 earlier literature.<sup>4</sup> Scale bar: 30 µm. 25



Figure S5: Dynamics of nonspecific QDs diffusing through extracellular space compared to
 specific QD-nanobody conjugate diffusing in dendrites. After transfection and labelling with
 QDs, nonspecific QDs (QD trajectories not present on transfected neuron) and specific QDs
 (QD trajectories present on dendrites) were identified and analysed as described earlier.

## 1 SI Video 1

2 GFP-NrCAM dynamics in untreated cells labelled with QD-nanobody conjugate.

# 3 SI Video 2

- 4 GFP-NrCAM dynamics in cells treated with Cytochalasin D followed by labelling with QD-
- 5 nanobody conjugate.

# 6 SI Video 3

7 GABA<sub>A</sub> receptor dynamics in *ex vivo* slices labelled with QD-nanobody conjugate.

# 8 SI Video 4

9 A representative track of a GABA<sub>A</sub> receptor in *ex vivo* slices labelled with QD-nanobody
10 conjugate showing confinement inside synapse.

# 11 SI Video 5

- Simultaneous imaging of GABAA receptors (green) and GPI anchored proteins (red) at
- 14 dendritic shaft and synapses (violet).

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