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1 TITLE:

Single Synapse Indicators of Glutamate Release and Uptake in Acute Brain Slices from Normal
 and Huntington Mice

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Glutamate release, glutamate clearance, glutamate spread, presynaptic terminal, astrocyte, glutamate uptake, decay kinetics, spill-over, optogenetics, iGlu_u, sCMOS

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SUMMARY:

We present a protocol to evaluate the balance between glutamate release and clearance at single corticostriatal glutamatergic synapses in acute slices from adult mice. This protocol uses the fluorescent sensor $iGlu_u$ for glutamate detection, a sCMOS camera for signal acquisition and a device for focal laser illumination.

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ABSTRACT:

Synapses are highly compartmentalized functional units that operate independently on each other. In Huntington's disease (HD) and other neurodegenerative disorders, this independence might be compromised due to insufficient glutamate clearance and the resulting spill-in and spill-out effects. Altered astrocytic coverage of the presynaptic terminals and/or dendritic spines as well as a reduced size of glutamate transporter clusters at glutamate release sites have been implicated in the pathogenesis of diseases resulting in symptoms of dys-/hyperkinesia. However, the mechanisms leading to the dysfunction of glutamatergic synapses in HD are not well understood. Improving and applying synapse imaging we have obtained data shedding new light on the mechanisms impeding the initiation of movements. Here, we describe the principle elements of a relatively inexpensive approach to achieve single synapse resolution by using the new genetically encoded ultrafast glutamate sensor iGluu, wide-field optics, a scientific CMOS (sCMOS) camera, a 473 nm laser and a laser positioning system to evaluate the state of corticostriatal synapses in acute slices from age appropriate healthy or diseased mice. Glutamate transients were constructed from single or multiple pixels to obtain estimates of i) glutamate release based on the maximal elevation of the glutamate concentration [Glu] next to the active zone and ii) glutamate uptake as reflected in the time constant of decay (TauD) of the perisynaptic [Glu]. Differences in the resting bouton size and contrasting patterns of short-term plasticity served as criteria for the identification of corticostriatal terminals as belonging to the intratelencephalic (IT) or the pyramidal tract (PT) pathway. Using these methods, we discovered that in symptomatic HD mice ~40% of PT-type corticostriatal synapses exhibited insufficient glutamate clearance, suggesting that these synapses might be at risk to excitotoxic damage. The results underline the usefulness of TauD as a biomarker of dysfunctional synapses in Huntington mice with a hypokinetic phenotype.

INTRODUCTION:

The relative impact of each synaptic terminal belonging to a "unitary connection" (i.e., the connection between 2 nerve cells) is typically assessed by its influence on the initial segment of the postsynaptic neuron^{1,2}. Somatic and/or dendritic recordings from postsynaptic neurons represent the most common and, until now, also the most productive means to clarify information processing under a top-down or vertical perspective³⁻⁵. However, the presence of astrocytes with their discrete and (in rodents) non-overlapping territories may contribute a horizontal perspective that is based on local mechanisms of signal exchange, integration and synchronization at synaptic sites⁶⁻¹⁰.

Because it is known that astroglia play, in general, a major role in the pathogenesis of neurodegenerative disease^{11,12} and, in particular, a role in the maintenance and plasticity of glutamatergic synapses¹³⁻¹⁶, it is conceivable that alterations in synaptic performance evolve in accordance with the state of astrocytes in the shared target area of afferent fibers with diverse origin. To further explore the target-/astroglia-derived local regulatory mechanisms in health and disease, it is necessary to evaluate individual synapses. The present approach was worked out to estimate the range of functional glutamate release and clearance indicators and to define criteria that may be used to identify dysfunctional (or recovered) synapses in brain areas most closely related to movement initiation (i.e., first of all in the motor cortex and dorsal striatum).

The striatum lacks intrinsic glutamatergic neurons. Therefore, it is relatively easy to identify glutamatergic afferents of extrastriatal origin. The latter mostly originate in the medial thalamus and in the cerebral cortex (see¹⁷⁻²⁰ for more). Corticostriatal synapses are formed by the axons of pyramidal neurons localized in cortical layers 2/3 and 5. The respective axons form bilateral intra-telencephalic (IT) connections or ipsilateral connections via a fiber system that more caudally constitutes the pyramidal tract (PT). It has further been suggested that IT- and PT-type terminals differ in their release characteristics and size^{21,22}. In view of these data, one could also expect some differences in the handling of glutamate.

The striatum is the most affected brain area in Huntington's disease (HD)⁵. Human HD is a severe genetically inherited neurodegenerative disorder. The Q175 mouse model offers an opportunity to investigate the cellular basis of the hypokinetic-rigid form of HD, a state that has much in common with parkinsonism. Starting at an age of about 1 year, homozygote Q175 mice (HOM) exhibit signs of hypokinesia, as revealed by measuring the time spent without movement in an open field²³. The present experiments with heterozygote Q175 mice (HET) confirmed the previous motor deficits observed in HOM and, in addition, showed that the

observed motor deficits were accompanied by a reduced level of the astrocytic excitatory amino acid transporter 2 protein (EAAT2) in the immediate vicinity of corticostriatal synaptic terminals²⁴. It has therefore been hypothesized that a deficit in astrocytic glutamate uptake could lead to dysfunction or even loss of respective synapses^{25,26}.

Here, we describe a new approach that allows one to evaluate single synapse glutamate clearance relative to the amount of the released neurotransmitter. The new glutamate sensor iGlu_u was expressed in corticostriatal pyramidal neurons. It was developed by Katalin Török²⁷ and represents a modification of the previously introduced high-affinity but slow glutamate sensor iGluSnFR²⁸. Both sensors are derivatives of the enhanced green fluorescent protein (EGFP). For spectral and kinetic characteristics, see Helassa et al.²⁷. Briefly, iGlu_u is a low-affinity sensor with rapid de-activation kinetics and therefore particularly well suited to study glutamate clearance at glutamate-releasing synaptic terminals. The dissociation time constant of iGlu_u was determined in a stopped-flow device, which rendered a Tau_{off} value of 2.1 ms at 20 °C, but 0.68 ms when extrapolated to a temperature of 34 °C²⁷. Single Schaffer collateral terminals probed at 34 °C with spiral laser scanning in the CA1 region of organotypic hippocampal cultures under a 2-photon microscope exhibited a mean time constant of decay of 2.7 ms.

PROTOCOL:

All work has been carried out in accordance with the EU Directive 2010/63/EU for animal experiments and was registered at the Berlin Office of Health Protection and Technical Safety (G0233/14 and G0218/17).

NOTE: Recordings from Q175 wild-type (WT) and heterozygotes (HETs) can be performed at any age and sex. Here we studied males and females at an age of 51 to 76 weeks.

1. Injection of the glutamate sensor iGlu_u for expression in corticostriatal axons

1.1. Use the pipette puller (one step mode) to prepare borosilicate glass pipettes for the injection of the virus. After pulling, break the pipette manually to obtain a tip diameter of 30–50 µm. Autoclave the pipette and surgical instruments, including the drill for opening the skull.

1.2. Store the virus AAV9-CaMKIIa.iGlu_u.WPRE-hGH (7.5 x 10^{13} gc/mL) at -80 °C in 10 μ L aliquots. If injections are performed shortly after virus production (within 6 months), keep at 5 °C. Before surgery, take out the vial and maintain it at room temperature.

1.3. Fill the glass pipette and remove any bubbles.

1.4. Anesthetize the animal with an intraperitoneal injection of a solution containing 87.5 mg/kg ketamine and 12.5 mg/kg xylazine. Subcutaneously inject 0.25% bupivacain (8 mg/kg) for additional pain relief. Check the depth of anesthesia by monitoring the muscle tone and observing the absence of pain-induced reflexes.

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1.5. Shave the skin on the head and sterilize it with 70% alcohol. Fit the mouse into the stereotaxic frame.

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1.6. Use a scalpel to remove the skin and a high speed (38,000 rpm) -drill to make a 1.2 mm hole in the bone above the motor cortex.

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1.7. Mount the syringe with the attached injection pipette in the holder of a precision manipulator. Insert the vertically oriented pipette into the cortex at 4 different sites. The injection coordinates are, with respect to bregma (in mm): anterior 1.5, lateral 1.56, 1.8, 2.04, 2.28. The depth with respect to *dura mater* is (in mm): 1.5–1.7.

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1.8. Using the injection system, inject 0.3 μ L per site of the undiluted virus solution with a velocity of 0.05 μ L/min. After each injection, leave the pipette in place for 1 min before withdrawing it slowly (1 mm/min).

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1.9. Finish by closing the surgical wound with a nylon suture.

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1.10. Leave the mouse for 0.5 to 1 h on a heating pad in a clean cage before returning it to its original cage.

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1.11. Maintain the mouse on a 12 h day-night cycle for 6 to 8 weeks before the preparation of acute brain slices.

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160 161 NOTE: To avoid immune reactions resulting in cell damage and synapse loss, injection of multiple viral constructs must be performed simultaneously or within 2–3 hours after the primary injection. The coordinates for $iGlu_u$ injection were selected according to the Paxinos and Franklin²⁹ brain atlas. They correspond to the M1 motor cortex. Immunostaining of injected brains visualized numerous but mostly well isolated axons and axon varicosities in the ipsi- and contralateral striatum and in the contralateral M1 and S1 cortices.

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2. Search for glutamatergic terminals expressing the glutamate sensor iGlu_u

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2.1. Calibration mode

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- 2.1.1. Prepare the sCMOS camera and the camera control software with the following settings.
- On the **Readout** page set **Pixel readout rate**: 560 MHz (= fastest readout), **Sensitivity/Dynamic**
- 170 Range: bit, Spurious Noise Filter: Yes, Overlap Readout: Yes. On the Binning/ROI page, select
- 171 Full screen.

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2.1.2. Prepare a glass slide containing a drop of 5 mg/mL Lucifer Yellow (LY) under a cover slip.

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- 2.1.3. Place the LY slide under a 63x objective, open the laser shutter and perform a serial
- acquisition using the following settings: Pixel Binning: 1x1, Trigger mode: Internal, Exposure

time: Minimal (to be determined).

2.1.4. Adjust the laser power to produce a fluorescence spot of 4 μm in diameter (the image
 should not contain saturated pixels).

2.1.5. To perform a calibration of the laser positioning system, select the following settings in the laser positioning software: **Spot-size diameter**: 10, **Scanning velocity**: 43.200 kHz. Click the **Start image acquisition** button on the right panel. Set **Runs**: 0, **Run delay**: 0, and select the **Run at TTL** option on the **Sequence** page. Click the **Calibrate button** on the Calibration page and calibrate the laser control software according instructions shown in the top-left corner of the acquisition screen.

2.1.6. If the setup uses independent software for camera and laser control, acquire screenshots from the camera acquisition window and send it to the laser control software for a recalculation of the XY coordinates. If the software is installed on different computers, then use a video grabber to import the image into the laser control software. The laser control software will need the information on the XY scaling factors and offsets. For this purpose, determine the coordinates of the top-left, bottom-left and bottom-right corners of the image within the acquisition window of the laser control program. Calculate the scaling factors according to the following equations: X factor = (X bottom-right – X bottom-left)/2048, X offset = X bottom-left, Y factor = (Y top-left – Y bottom-left)/2048, Y offset = Y bottom-left.

2.1.7. At the end of the calibration procedure, create a rectangular region of interest (ROI) of 267x460 pixels, move it to the center of the screen and click the **Start sequence** button.

2.1.8. Return to the following camera control software settings: **Binning**: 2x2, **Trigger mode**: External exposure, **Exposure time**: Minimal (to be determined).

2.2 Autofluorescence correction mode

NOTE: The resting level of [Glu] in the environment of active synaptic terminals is typically below 100 nM $^{14,30-34}$. Accordingly, any sensor of glutamate, especially a low affinity sensor like iGlu_u will be rather dim in the absence of synaptic glutamate release. Nevertheless, some iGlu_u fluorescence can even be detected at rest, but must be distinguished from the tissue autofluorescence. The 473 nm illumination elicits both iGlu_u fluorescence and autofluorescence (**Figure 1A–C**). The latter occupies a wide range of wavelengths, while iGlu_u fluorescence is limited to 480–580 nm (with a maximum at 510 nm). The correction for autofluorescence is based on the acquisition of two images with different high-pass filters.

2.2.1. Prepare brain slices in advance as described elsewhere³⁵. Keep bran slices ready.

2.2.2. Transfer the slices into the recording chamber, submerging them into oxygenized artificial cerebrospinalfluid (ACSF) containing 125 mM NaCl, 3 mM KCl, 1.25 mM NaH₂PO₄, 25 mM NaHCO₃, 2 mM CaCl₂, 1 mM MgCl₂, 10 mM glucose (pH 7.3, 303 mOsm/L), supplemented

with 0.5 mM sodium pyruvate, 2.8 mM sodium ascorbate and 0.005 mM glutathione. Use a flow rate of 1–2 mL/min. Keep the bath temperature at 28–30 °C.

2.2.3. Locate the dorsal striatum under a 20x water immersion objective. Fix the slices with a nylon grid on a platinum harp to minimize the tissue movement. Switch to the 63x /NA 1.0 water immersion objective. Select filters reflecting light at 473 nm (dichroic mirror) and passing light with wavelengths >510 nm (emission filter).

2.2.4. Synchronize illumination, stimulation and image acquisition using an AD/DA board with the respective control software. Set the trigger program to control acquisition with a laser exposure time of 180 ms and an image acquisition time of 160 ms. Use the laser positioning device to send the laser beam to a predetermined number of points and define the settings for **Scanning velocity** and **Spot size**.

2.2.5. Acquire an image of both autofluorescence and $iGlu_u$ -positive structures using a high pass filter at 510 nm ("yellow image").

2.2.6. Acquire an image with autofluorescence alone using a high-pass filter at 600 nm ("red image").

2.2.7. Scale the red and yellow images, using the mean intensities of the 10 brightest and the 10 darkest pixels to define the range. Perform a subtraction "yellow minus red image" and rescale the subtracted image to generate a standard 8-bit tif file for convenient visualization of the bouton of interest (**Figure 1D**). It contains the bright pixels from the iGlu $_u$ -positive structures, grey pixels from the background and dark pixels from structures with autofluorescence.

NOTE: With the given equipment the mean resting fluorescence (F) will be below 700 A.U.

2.3 Bouton search mode

NOTE: $iGlu_u$ -positive pixels may belong to functionally different elements of the axonal tree, such as axon branches of different order, sites of bifurcation, varicosities after vesicle depletion or fully active varicosities. However, it is almost impossible to identify functional synaptic terminals just by visual inspection. Therefore, each glutamatergic synaptic terminal needs identification by its responsiveness to electrical depolarization. Sites that do not respond to stimulation have to be discarded. The physiological means to induce glutamate release from corticostriatal axons is to elicit an action potential. This can either be achieved by using a channel rhodopsin of appropriate spectral characteristics or by electrical stimulation of an axon visualized by $iGlu_u$ itself. To avoid accidental opsin activation, we preferred the latter approastep

2.3.1. Use the micropipette puller (in a four-step mode) to produce stimulation pipettes from borosilicate glass capillaries. The internal tip diameter should be about 1 μ m. When filled with ACSF, the electrode resistance should be about 10 M Ω .

2.3.2. To induce the action potential-dependent release of glutamate from a set of synaptic boutons attached to the same axon, use 63x magnification, the 510 nm emission filter and the subtracted image to place a glass stimulation electrode next to a fluorescent varicosity. Avoid the proximity of additional axons.

2.3.3 Turn on the stimulator to deliver depolarizing current pulses to the stimulation pipette. Use current intensities around 2 μ A (no more than 10 μ A).

2.3.4. Turn on the multi-channel bath application system where one channel delivers the standard bath solution and the other channels deliver the necessary blockers of ion channels, transporters or membrane receptors. Control the flow at the site of recording and then switch to the tetrodotoxin (TTX) channel (bath solution plus 1 μ M TTX). After 2-3 min, stimulate the bouton of interest again, but now in the absence of action potential generation. The release is directly due to calcium influx through voltage-depended calcium channels.

2.3.5. By turning the intensity key on the stimulator, adjust the stimulation current to obtain responses similar to those elicited via action potentials. Typically, the stimulation current would be around 6 μ A for a 0.5 ms depolarization.

2.3.6. For basic testing of the preparation, apply 0.5 mM CdCl₂. The presence of this calcium channel blocker glutamate release completely prevents the synaptic glutamate release thereby validating the calcium dependence of the directly evoked glutamate signal.

NOTE: The following general recommendation may help to increase the success rate of single synapse experiments in the striatum. To select glutamatergic synaptic terminals with intact release machinery, a varicosity should: (i) Have a smooth spindle-like shape; (ii) Not be associated with an axon bifurcation; (iii) Be brighter than other structures on the "yellow image"; (iv) Reside in the striatal neuropil rather than in the fiber tracts; (v) Reside on a very thin axon branch; (vi) Not reside within the deeper parts of the slice.

3. Visualization of glutamate release and clearance

3.1. Recording mode

NOTE: After the subtraction of the autofluorescence (step 2.2) and a few initial tests for responsiveness of the bouton of interest (Step 2.3), data acquisition can begin. The responses to electrical activation of glutamate release from single axon terminals can be observed in the image directly (Figure 2), without applying further analysis tools. However, it proved to be very convenient to immediately extract some basic indicators of synapse performance (Figure 3). This information is needed to make decisions on subsequent course of experiment, such as selection of particular terminal types, rapid assessment of HD-related alterations, the number and frequency of trials or drugs to be applied with the superfusion system. It might also be necessary to deal with eventually appearing artifacts. First, the standard settings for data

recording is described.

3.1.1. Using the microscope XY drives, place the tested iGlu $_{u}$ -positive bouton close to the viewfield center. Stop the image acquisition with the **Abort acquisition button**. On the last acquired picture, determine the XY position of the resting bouton center by clicking on it with the left mouse button. The XY coordinates of the set cursor will be shown on the bottom status panel of the acquisition window.

3.1.2. Using the calibration data (step 2.1.6), calculate the coordinates of the site where the laser beam should be sent for the excitation of the $iGlu_u$ fluorescence. Use the following equations: X laser = X camera * X factor + X offset, Y laser = Y offset – Y camera * Y factor. While performing this recalculation, pay attention to the vertical or horizontal flip settings of the camera.

3.1.3. Create a one-point sequence in the laser control software using the calculated coordinates. For this purpose, select **Point** in the **Add to sequence** box on the **Sequence page** of the laser control software. Select 10 µs for the delay to trigger onset and 180 ms for the laser pulse time. Move the mouse to the calculated coordinates and click left button.

328 3.1.4. Select the following settings in the laser control software: Runs: 0, Run delay: 0, 329 Sequence: Run at TTL. Then click Start sequence.

3.1.5. In the camera control software, select the following settings. On the Binning/ROI page, set Image Area: Custom, Pixel Binning: 2x2, Height: 20, Width: 20, Left: X-coordinate of the resting iGluu-positive spot minus 10 px, Bottom: Y-coordinate of the resting iGluu-positive spot minus 10 px, Acquisition Mode: Kinetic Series, Kinetic Series Length: 400, Exposure Time: 0.0003744s (minimal value). With such settings, the acquisition rate will be 2.48 kHz.

3.1.6. Select **Trigger mode**: External. Click **Take signal** in the camera control software. Initiate the experimental protocol laid down for the trigger device.

3.1.7. Implement the experimental protocol **Trial** with the following time line: 0 ms – start trial, 1 ms – start laser illumination, 20 ms – start image acquisition with camera, 70 ms – start electrical stimulation 1, 120 – start electrical stimulation 2, 181 ms – end trial (laser and camera off). Thus, during one trial the camera acquires 400 frames with a frequency of 2.48 kHz. See steps 2.3.3 and 2.3.5. for details on electrical stimulation.

3.1.8. To allow for sufficient recovery of presynaptic vesicle pools, apply the protocol **Trial** with a repetition frequency of 0.1 Hz or lower.

349 3.2 Off-line construction of the glutamate transient and rapid assessment of glutamate release and clearance for the identification of pathological synapses

3.2.1. Turn on the evaluation routines. Here, we use an in-house-written software SynBout v.

3.2. (author: Anton Dvorzhak). The following steps are needed to construct a $\Delta F/F$ transient from the pixels with elevated iGlu, fluorescence as in the video of **Figure 2.**

3.2.2 To determinate the bouton size, calculate the mean and standard deviation (SD) of the ROI fluorescence intensity at rest (F), before the onset of stimulation. Determine and box the area occupied by pixels with F>mean + 3 SD (**Figure 3A**). Determine a virtual diameter (in μ m) assuming a circular form of the supra-threshold area.

3.2.3. Determine ΔF as the difference between the peak intensity value and F. Plot the stimulating current and $\Delta F/F$ against time (**Figure 3B**). Calculate the SD of $\Delta F/F$ at rest (before the onset of stimulation) and the *Peak amplitude*. Use pixel with peak amplitude more than 3 SD of $\Delta F/F$ to perform monoexponential fitting for the decay from peak. Determine the time constant of decay *TauD* of $\Delta F/F$.

3.2.4. To estimate the *Maximal amplitude* at a given synapse, select the pixel with the highest $\Delta F/F$ value. It is typically located within or next to the boundaries of the resting bouton $iGlu_u$ fluorescence. The *Maximal amplitude* would be the best indicator of the glutamate load presented to the clearance machinery of a single synapse.

3.2.5. To estimate the spatial extension of the iGlu_u signal, determine the diameter of the area of all supra-threshold pixels combined to form a virtual circle. The respective diameter is called *Spread*. The term *Peak spread* then refers to the peak value of the averaged spread transient (**Figure 3C**, difference between dotted red lines).

3.3 Corrections for possible artifacts

NOTE: Electrical depolarization is associated with an outward flow of the intra-pipette solution. This may result in a small tissue displacement. To discriminate between stimulation-induced changes of $iGlu_u$ and out-of-focus shifts of the image, one could use the following approach to identify and to eliminate displacement artifacts (**Figure 4**).

3.3.1. Analyze the spatial characteristics of the suprathreshold pixels derived from a ROI with signs of displacement (Figure 4F–H).

3.3.2. Find pixels outside the initially determined boundaries of the bouton at rest (**Figure 4F–H**, boxed in red).

3.3.3. Identify eventually existing negative pixel intensity values (**Figure 41, J**). Any $\Delta F/F$ in the negative direction with an amplitude larger than the pre-stimulation mean \pm 3 SD should be regarded as artifact, and the record should be discarded.

3.3.4. To avoid out-of-focus artifacts, place the stimulation electrode on the antero-lateral side of the synaptic bouton, use biphasic electrical stimulation and minimize the stimulation intensity/duration. NOTE: Out-of-focus artifacts can also be derived from the camera. If the latter is not optimally fixed to the microscope it can produce oscillations, most likely due to small vibrations of the camera cooling system. Such vibrations create a "yin and yang" pattern in the $\Delta F/F$ images rotating with a frequency of 50 Hz. The vibrations can be mathematically subtracted from the fluorescence signal, but the final quality of the measurements would then critically depend on the recording time.

3.3.5. Fix the camera to the microscope such that vibrations are absent. If the latter remain, place a rubber gasket between the camera and microscope adapter.

NOTE: One cannot neglect the possibility that the striatal neuropil around the synapse of interest contains glutamatergic varicosities that do not express iGlu_u and therefore remain invisible. In the case of their co-activation (more likely in the absence of TTX) the transients obtained from the boutons of interest might be affected by spill-over. The same applies to iGlu_u-expressing terminals that were out of focus. A characteristic phenomenon would in this case be that fluorescence transients originate from a much wider area. As this response is eliminated by TTX, one can interpret it as an unspecific response induced by unwarranted multi-fiber activation.

3.3.6. To correct for this unspecific multi-fiber response, follow the procedure outlined in **Figure** 5. Use the fluorescence signal of pixels at the viewfield boundaries. To minimize background response, minimize stimulation intensity and duration or use TTX.

REPRESENTATIVE RESULTS:

Identification of two types of corticostriatal glutamatergic varicosities

IT and PT afferents originate in layer 2/3 and 5, respectively, and exhibit differential ramification and termination patterns in the ipsilateral and contralateral (IT terminals only) striatum. Still little is known about the properties of glutamate release and clearance under repetitive activation conditions as observed during the initiation of movements, but it is well documented that the respective glutamate-releasing varicosities differ in size²². Applying a size criterion, it was found that IT and PT terminals exhibit contrasting forms of short-term plasticity²⁴. At stimulus intervals of 50 ms, the smaller IT terminals were prone to paired pulse depression (PPD) while the larger PT terminals showed paired pulse facilitation. This difference was also observed at shorter intervals (20 ms) and throughout a series of 6 pulses. **Figure 3** and **Figure 6** illustrate these experiments where synaptic glutamate release was elicited via the action potential mechanism at physiological Ca²⁺/Mg²⁺ concentration.

Identification of dysfunctional synapses in mice with advanced Huntington's disease

Neurodegenerative diseases such as Alzheimer's, Parkinson's and Huntington's disease are characterized by an ever progressing loss of glutamatergic synapses³⁶. Novel therapies aim to impede or even reverse this fatal progression. What exactly triggers the disappearance of a synapse, and when, is largely unknown. Further insight can be expected from studies that offer criteria for (a) the vital identification of a particular class of glutamatergic synapses and (b) the

detection of dysfunctional versus normally performing contacts. Here it will be shown how the *TauD* values obtained from PT-type of terminals were used to estimate the fraction of dysfunctional synapses in Q175 heterozygotes with an identified motor phenotype.

Prior to the single synapse imaging experiments, the mice were submitted to a rapid but rather robust test for alterations in their exploratory behaviors. This test is called "step-over test". The animal was placed into the center of a Petri dish (of 185 mm diameter and 28 mm wall height). The test was recorded with a video camera. Using offline analysis, one can determine the time between the take-off of the experimenter's hand and the moment when the animal has all 4 feet out of the dish. Plotting the data from over 100 WT and Q175 HET at ages between 12 and 18 months suggests that mice with a step-over latency of >300 s can be diagnosed as hypokinetic. **Figure 7A** illustrates a significant positive correlation between the results obtained for the total path run in the open field and the step-over latency.

Single synapse iGlu $_u$ imaging showed that these symptomatic HD mice exhibited a deficit in the speed of juxtasynaptic glutamate decay as reflected in the TauD values from single (or first in a sequence) stimuli (**Figure 7B,C**). In WT, such prolongation was only observed after the application of a selective non-transportable inhibitor of glutamate uptake - DL-threo- β -benzyloxyaspartic acid (TBOA, **Figure 7D,E**). This suggests a role of astrocytic glutamate transporters in the regulation of synaptic glutamate clearance. Changes in the diffusion of Glu in the perisynaptic space have not been found²⁴. But, of course, much more work is needed to actually identify the cause of slowed glutamate clearance in HD as well as in other forms of parkinsonism. Apart from changes in the astrocyte proximity⁹ and reduced slc1A2 expression³⁷, one may as well consider disease-related instability of EAAT2 in the plasma membrane of the perisynaptic astrocyte processes (PAPs). This might be a result of changes in the EAAT2 interactome. Indeed, recent mass spectroscopy experiments in the lab point to a loss of EAAT2-dystrophin interaction in striatal astrocytes (Hirschberg, Dvorzhak, Kirchner, Mertins and Grantyn, unpublished).

Very little is known regarding the timeline of synaptic dysfunction with progression of HD, but it is very likely that healthy synapses co-exist with already impaired ones. In searching for a classification criterion, we examined the *TauD* data from different mice. For this purpose, the amplitude and *TauD* values from 3 consecutive paired trials were normalized to the first response (see **Figure 6F** for an experimental scheme), and the probability of occurrence of a given *TauD* value was compared in age-matched WT versus Q175 HET (**Figure 6G**). It was found that in WT *TauD* never exceeded 15 ms, while in symptomatic Q175 HET, 40% of the synapses exhibited *TauD* values between 16 and 58 ms, despite a tendency for a reduction in the amount of released glutamate (**Figure 6H,I**). *TauD* might then be regarded as a biomarker for dysfunctional synapses in HD and further be used to verify functional recovery in experiments targeting astrocytic glutamate transport.

FIGURE LEGENDS:

Figure 1: Identification of iGlu_ν-positive varicosities. (A) Fluorescence image obtained with a

510 nm high pass filter ("yellow image"). (**B**) Same view field acquired with a 600 nm high pass filter ("red image"). Note that that the spot marked with black arrowhead has disappeared in (B). Overlay of (A) and (B). White arrow = autofluorescence, black arrow = iGlu_u-positive varicosity. (**D**) Image obtained by subtraction of (B) from (A). The autofluorescent spots are dark and the iGlu_u+ spot is bright.

Figure 2: Movie still from a slow-motion video (slowdown factor 1240x). Upper row: Images from WT (left), Q175 HET (middle) and HOM (right). Lower row: Respective iGlu_u transients from the pixel with the highest glutamate elevation ($Maximal\ \Delta F/F$). The red cursor indicates the point on the transient corresponding to the image above the trace. note prolonged elevation of iGlu_u fluorescence (red pixels and $\Delta F/F$ transients). Modified and reprinted with permission from Dvorzhak et al. ²⁴

Figure 3: Extraction of functional indicators from single synapse images of the genetically encoded ultrafast Glu sensor iGlu_u in corticostriatal neurons. (A, D) Example of a PT (A) and IT (D) bouton with the respective iGlu_u fluorescence at rest (left) and at the peak of an AP-mediated iGlu_u response (right). (B, C, E, F) iGlu_u responses recorded from the bouton shown in (A, D); Experiment in 2 mM Ca²⁺ and 1 mM Mg²⁺. (B) Simultaneous recording of the stimulation current (upper trace) and mean intensity of supra-threshold pixels (bottom trace). Same time scale for all traces. Peak amplitudes (between dotted red horizontal lines) and a monoexponential function fitted to the decay from this peak (red overlay). The corresponding TauD (τ) values are shown next to the fitting curves. (E, F) Plot of spread against time. Peak spread: difference between dotted red horizontal lines.

Figure 4: Characteristics of a glutamate-induced iGlu_u transient (A–E) as opposed to a displacement artifact (F–J). (A, B) and (F, G) show the absolute fluorescence intensity at rest (A, F) and after stimulation (B, G) in arbitrary units (au). (C, H) Fluorescence change in percent of the resting fluorescence prior to stimulation (Δ F/F%). (D, I) Superposition of the iGlu_u transients from all pixels in arbitrary units. (E, J) Superposition of the iGlu_u transients from all pixels in Δ F/F %. In the case of synaptic glutamate release, the pixels next to the resting terminal exhibit a fluorescence increase after stimulation, whereas in the case of out-of-focus shifts the brightest pixel at rest merely change their position in the ROI, without an accompanying increase in the over-all fluorescence intensity of the view-field. A displacement artifact can also be recognized by the appearance of negative Δ F/F signals (J).

Figure 5: Correction for unspecific $iGlu_u$ response. (A–D) Example of paired synaptic response contaminated by an unspecific background response. (E–H) Same after correction. (A, E) Before stimulation. (B, F) During stimulation. (C, G) After stimulation. (D, H) Corresponding superimposed intensity transients (in $\Delta F/F$) from all pixels of the ROI. The timepoint of acquisition of the images is marked by corresponding small letters over the arrowheads. Note that in panel C the background response is very widespread and slowly decaying.

Figure 6: Distinct size and size-related differences in the amplitude paired pulse ratio (PPR) of the $iGlu_u$ transient. (A) Simplified scheme of the corticostriatal circuitry²², illustrating the

concept of preferential projection of pyramidal tract (PT) neurons to indirect pathway striatal projection neurons (iSPNs) and intratelencephalic (IT) neurons to direct pathway SPNs (dSPNs), with size-differences between the IT and PT terminals. (B) Bimodal distribution of bouton diameters as determined by the supra-threshold resting fluorescence before stimulation. Boutons with diameter ≥ 0.63 µm were defined as "Large" and assumed to be issued by PT axons. For original images and specimen traces from PT- and IT-type of synapses see Figure 3. (C) Significant positive correlation is seen between the PPR of peak amplitudes and the bouton diameters. Each data point represents the average from the first 3 trials of each bouton. *P<0.05, **P<0.01, ***P<0.001.

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Figure 7: Identification of dysfunctional synapses in Q175 mice with a hypokinetic phenotype.

(A) Results of motor testing on the day of single synapse imaging. Step-over latencies >300 ms were considered as pathological. At the age tested (average 16 months), 17/54 HET exhibited a pronounced phenotype in the step-over test. With regard to hypokinesia, the step-over test seems to be more sensitive than the open field test. Nevertheless, there was a significant correlation between the outcome of the step-over test (time between placement and barrier crossed with all four feet) and the open field test (i.e., total path run in 5 min). Y = -0.01511X + 458,7; P=0.0044 (simple regression). (B) Average evoked single synapse iGlu_u transients normalized to same peak amplitude to illustrate HD-related differences in the clearance of synaptically released glutamate. The respective fitting curves highlight the differences in the duration of the glutamate transients. (C) Quantification of the results from wild-type (grey), HET (red) and HOM (magenta). (D, E) Incubation of WT slices in 100 nM of TFB-TBOA simulated the depression of glutamate clearance observed in HOM. (F) Scheme of synapse activation and data organization. (G) Cumulative histograms of #1 TauD values. (H, I) Plots of normalized #2 responses. Data from 31 WT and 30 HET synapses (all of PT type). In the WT sample all TauDmax values were ≤15 ms. In the HET sample, 40% of synapses exhibited TauDmax values exceeding the 15 ms threshold defined by the longest TauDmax in WT. These graphs emphasize the HD-related differences in the ranges of maximal amplitude (i.e., the value from the pixel with the highest iGlu_u fluorescence increase) and TauDmax (the TauD of the pixel with the highest elevation). TauDmax values exclusively encountered in HET are shown in red, and amplitude values exclusively seen in WT are shown in grey. *P<0.05, **P<0.01, ***P<0.001. The used statistical tests are indicated next to the respective graph.

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DISCUSSION:

The experiments concern a question of general interest – synapse independency and its possible loss in the course of neurodegeneration, and we describe a new approach to identify affected synapses in acute brain slices from aged (>1 year) mice. Taking advantage of the improved kinetic characteristics of the recently introduced glutamate sensor iGlu_u the experiments illuminate the relationship between synaptic glutamate release and uptake in a way that has not been possible before.

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The influence of glutamate clearance on the function and maintenance of synapses is not very well elucidated, although the hypothesis that glutamate-induced excitotoxicity can cause neuron loss and synapse pruning is mentioned in almost any pertinent review on epilepsy,

stroke and neurodegenerative diseases³⁸⁻⁴¹. However, the available evidence is more limited than possibly anticipated from the literature. Confusion is further added by the fact that the selected experimental tools might be insufficient in view of the problems associated with low spatial and temporal resolution when interpreting the results obtained with gross stimulation and recording techniques^{42,43}. This can lead to false negatives discouraging further research, which is more than unfortunate in view of neurologic disorders as severe as Huntington's disease. The present approach ensures better signal discrimination and therefore stronger support of the idea that in HD a significant fraction of corticostriatal synapses exhibits signs of impaired glutamate clearance.

The present results disclosed differences of short-term plasticity within the corticostriatal pathway. Although the latter had been in the focus of numerous studies^{17,44}, it has not been anticipated that the afferents originating from the upper cortical layers would exhibit frequency-dependent depression, in contrast to pyramidal tract afferents originating in layer 5. The latter preferentially showed a frequency-dependent potentiation of release and may therefore be at higher risk for glutamate uptake insufficiency.

Experiments on acute brain slices from adult mice are important for elucidating synaptic alterations at an appropriate age of life. The age and functional state of the preparation is particularly relevant if astrocytes were involved in the mechanism of interest. Here, it is essential that the astrocytes are mature enough to exhibit adult levels of glutamate transporter activity and related chloride homeostasis⁴⁵.

Single synapse assays will pave the road towards a better understanding of HD-related changes of glutamatergic synaptic transmission in the intact brain⁴⁶. It has been shown that single synapse resolution can also be achieved in the intact brain, provided that the synapses of interest are localized in the superficial layers of the cerebral cortex⁴⁷.

Finally, the present experimental approach has the appeal that it can be used by many followers, since the required equipment is still on the low-cost side.

In short, fluorescence signals reporting glutamate release and the juxtasynaptic changes in the concentration of glutamate in the intact brain can provide data unachievable with any electrophysiological method. But as with any new method, this approach has its limitations and disadvantages that have in part been addressed in steps 2.2 and 3.3. The low resting fluorescence of the fast and high affinity iGluu sensor requires a bit of exercise/experience to identify suitable varicosities for further testing. With co-expression of a genetically encoded calcium indicator (GECI) such as XCaMP-R⁴⁸ and at least two coordinated lasers illumination systems, the search of functional axon terminals would become much easier. In any case, it is critical to expose the preparation as little as possible to the exciting light before and during iGluu recording.

The use of a 473 nm laser (instead of the regular whole field elimination) to elicit $iGlu_u$ fluorescence is a prerequisite for obtaining sufficient emission but will also cause bleaching.

Under the given conditions, iGlu $_u$ was fully bleached after 10 stimulation/acquisition trials (10 stimulus pairs at an inter-stimulus interval of 50 ms and a repetition rate of 1/10 s). The maximal total illumination time for steady-state data acquisition with the presently used 1 photon laser system and the described setting was approximately 2 s. The bleaching of iGlu $_u$ is exponential, being very strong during the first 20 ms after illumination start and much slower thereafter. It is therefore advisable to avoid signal acquisition during the first 20 ms of each trial and to acquire not more than a total of 10 response pairs. Responses to single stimuli could be recorded with exposure times of 60-80 ms instead of the presently used 180 ms.

Another critical issue is the expression level of $iGlu_u$. In the pioneering study of Helassa et al.²⁷, the viral constructs were applied by electroporation to cultured neurons²⁷, which produces higher expression levels of the sensor and presumably also less bleaching especially if a two-photon laser scanning device can be used instead of a one-photon microscope. Dürst et al.⁴⁹ report routine acquisition of postsynaptic expression of $iGlu_u$ in CA1 pyramidal neurons in ~100 trials (each exposed for only 80 ms). However, cultured brain tissue is not an option when the experiments aim at clarifying astrocyte-dependent synaptic functions in the aged brain. A CaMKII-Cre-dependent expression of $iGlu_u$ using a stronger promoter may provide preparations with stronger expression in fewer cells, thereby increasing the resolution of the method and allowing for the acquisition of more trials per synapse.

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DISCLOSURES:

The authors have nothing to disclose.

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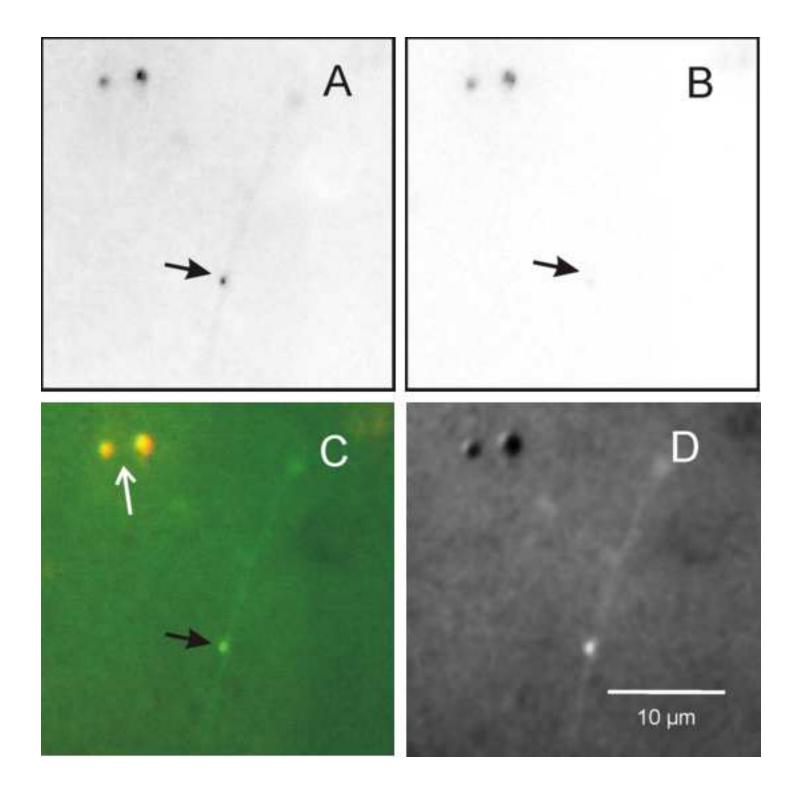
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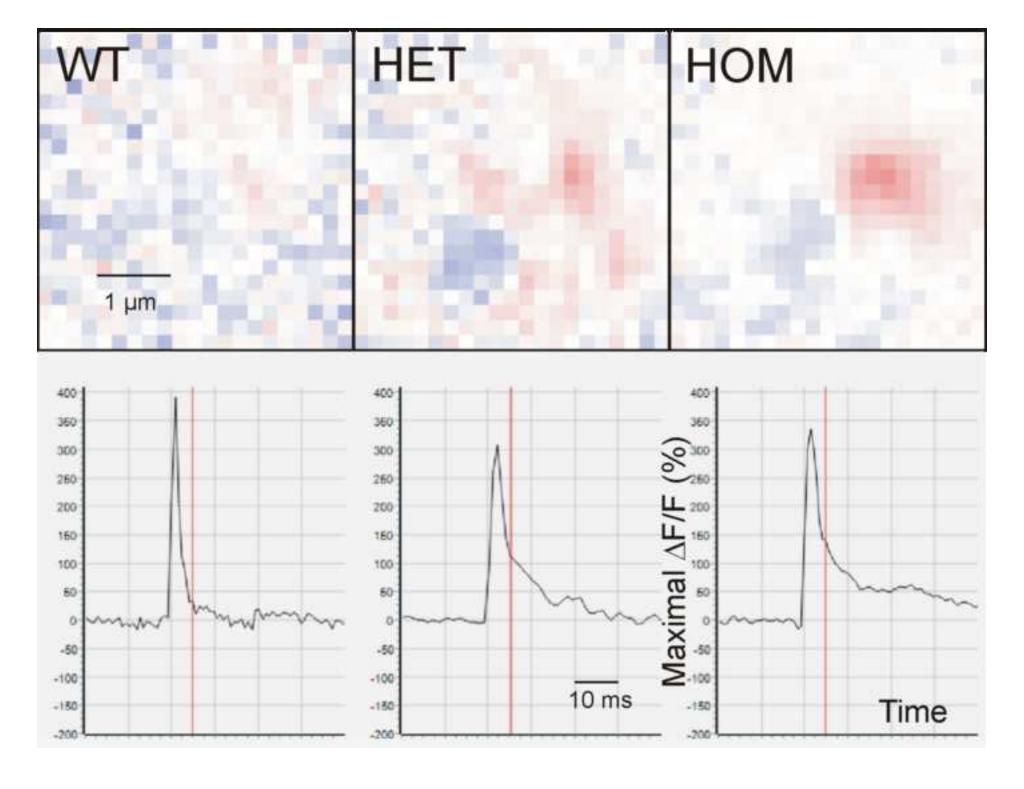
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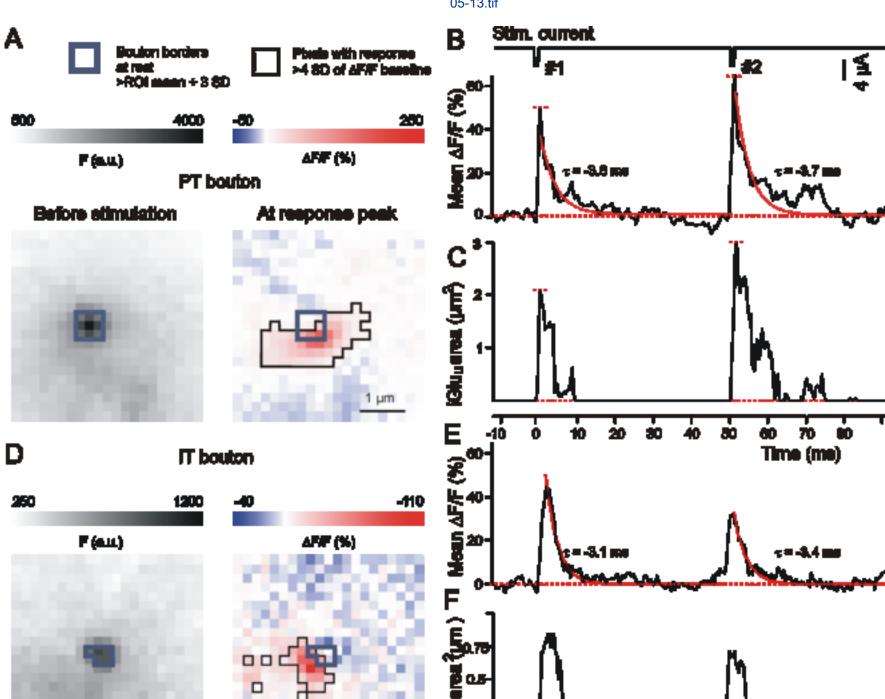
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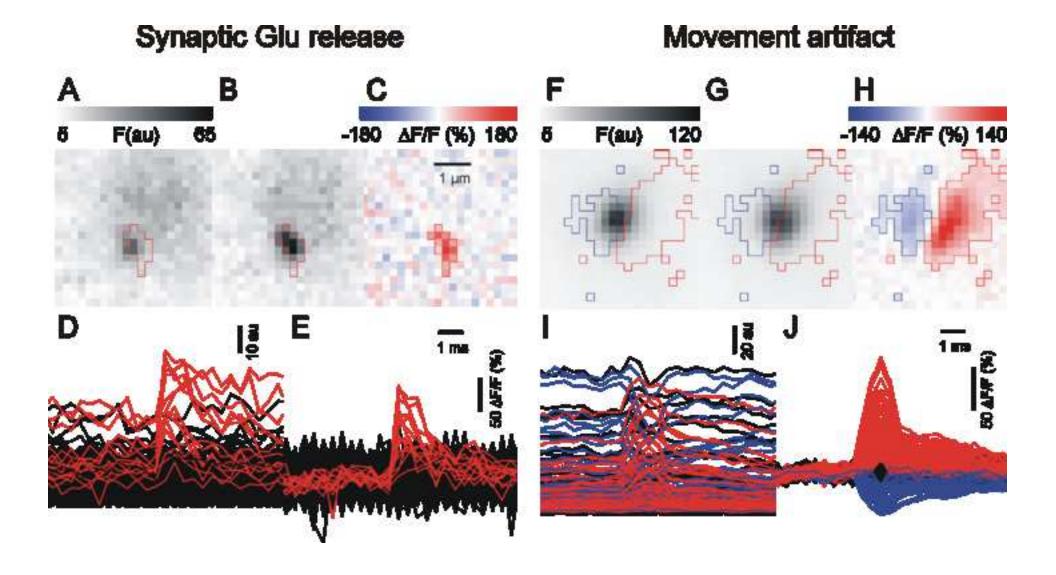
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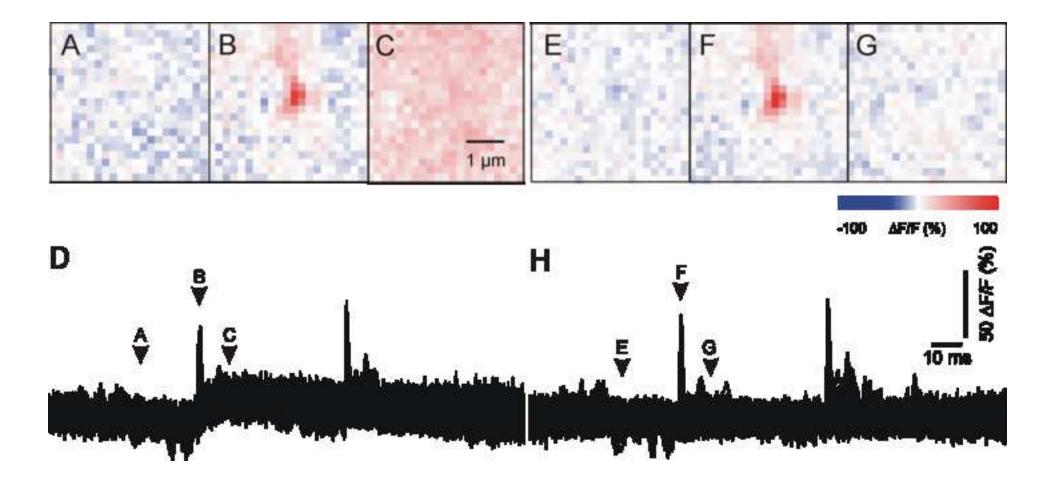


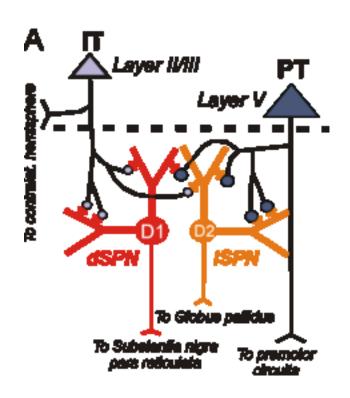


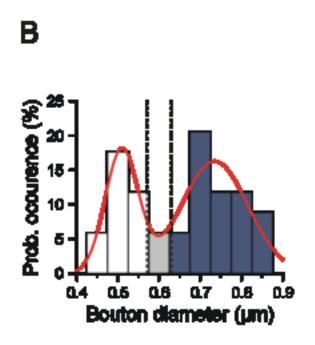


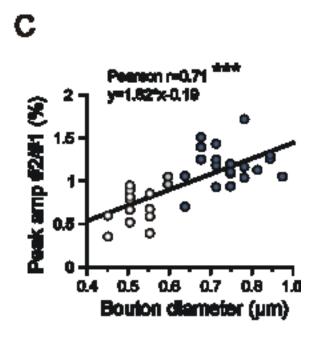
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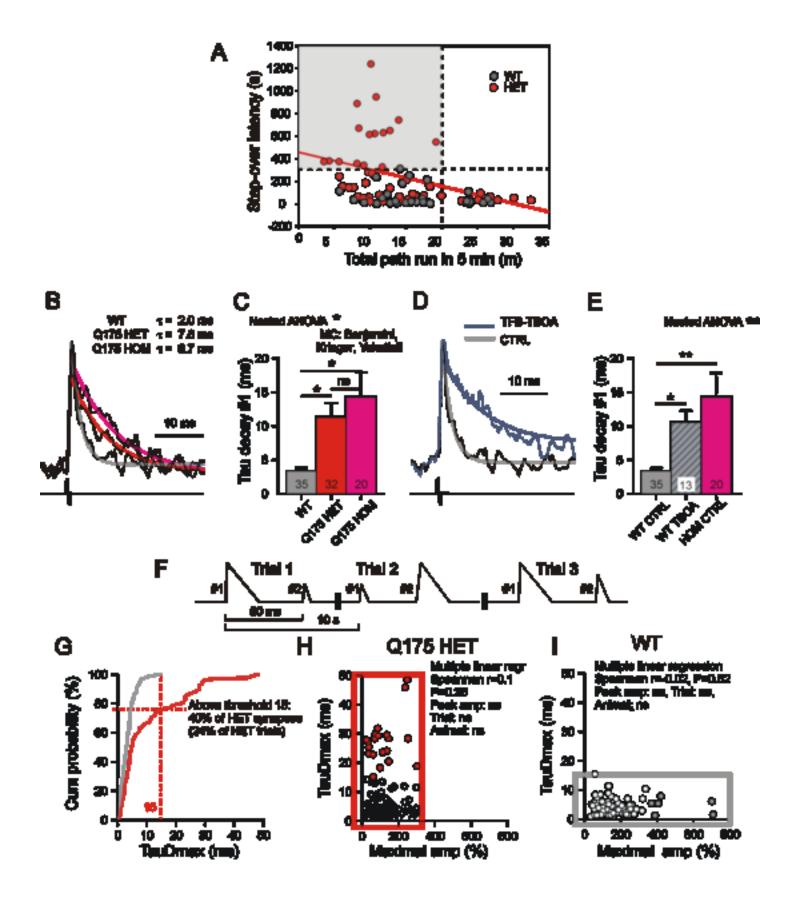












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Stereo microsope	Precision Stereo Zoom Binocular Microscope	WPI
Stereotaxic frame	Digital Lab New Standard stereotaxic frame	Stoelting
High speed drill equipment	Foredom K1070 cromoter Kit	Stoelting
Injection system	Quintessential Stereotaxic Injector (QSI)	Stoelting
Hamilton syringe 5 μl	75RN Syr (26s/51/2)	Hamilton
Laser positioning system	UGA-40	OptoElectronic
Blue laser for iGluu excitation	473 nm laser	OptoElectronic
Dichroic mirror for 473 nm	Dichroic	OptoElectronic
1P upright microscope	Axioskop 2 FS Plus	Carl Zeiss
Objective 63x/1.0	W Plan-Apochromat	Carl Zeiss
4x objective	Achroplan 4x/0,10	Carl Zeiss
Dichroic mirror for iGluu		Omega optical
Emission filter for iGluu		Omega optical
Dichroic mirror		Omega optical
Emission filter for autofluorescence subtr.		Omega optical
sCMOS camera	ZYLA 4.2MP Plus	Andor
Acqusition software	Solis	Andor
AD/DA converter	InstruTECH LIH8+8	HEKA Elektronik
Aquisition software	TIDA5.25	HEKA Elektronik
Electrode positioning system	Micromanipulator	Sutter Instrument
Electrical stimulator		Charite workshops
Slicer	Vibrotome	Leica
Brown/Flaming-type puller	P-1000	Sutter Instr
Glass tubes for injection pipettes		WPI
Glass tubes forstimulation pipettes		WPI
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iGlu <i>u</i> plasmid	pCI-syn-iGIu _u	Addgene
Q175 mice	Z-Q175-KI	Jackson Lab

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STIM-26
VT1200 S
SU-P1000
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Response to the Editor and the Reviewers

We are truly grateful to the Editor and the reviewers for the thorough attention to our manuscript. We have addressed all concerns. In the following the issues will be commented in the order of appearance in the reviews.

Dear Dr. Grantyn,

Editorial comments:

Changes to be made by the Author(s):

1. Please take this opportunity to thoroughly proofread the manuscript to ensure that there are no spelling or grammar issues. The JoVE editor will not copy-edit your manuscript and any errors in the submitted revision may be present in the published version. Please use American English throughout.

Done.

2. Please provide an email address for each author.

Done.

- 3. Please rephrase the Short Abstract/Summary to clearly describe the protocol and its applications in complete sentences between 10-50 words: "Here, we present a protocol to ..."

 Done.
- 4. Please remove all commercial language from your manuscript and use generic terms instead. All commercial products should be sufficiently referenced in the Table of Materials and Reagents. For example: Andor Zyla4.2 plus sCMOS camera, Solis, version 4.30.30034.0 (Andor), Solis Control software (all from Rapp OptoElectronics), UGA-40, Rapp OptoElectronic, Andor Zyla4.2 plus, Zeiss 10 Axioskop 2 FS Plus, oftware TIDA 5.24 (both HEKA 28 Elektronik), etc.

 Done, see attached Table 1.
- 5. Please adjust the numbering of the Protocol to follow the JoVE Instructions for Authors. For example, 1 should be followed by 1.1 and then 1.1.1 and 1.1.2 if necessary. Please refrain from using bullets or dashes.

Done.

6. Please revise the protocol text to avoid the use of any personal pronouns in the protocol (e.g., "we", "you", "our" etc.).

Done.

7. Please ensure that all text in the protocol section is written in the imperative tense as if telling someone how to do the technique (e.g., "Do this," "Ensure that," etc.). The actions should be described in the imperative tense in complete sentences wherever possible. Avoid usage of phrases such as "could be," "should be," and "would be" throughout the Protocol. Any text that cannot be written in the imperative tense may be added as a "Note."

Done.

8. The Protocol should contain only action items that direct the reader to do something. Please move the discussion about the protocol to the Discussion.

Done. See addition to Discussion on page 14.

9. The Protocol should be made up almost entirely of discrete steps without large paragraphs of text between sections. Please ensure that individual steps of the protocol should only contain 2-3 actions per step.

Done.

10. Please add more details to your protocol steps. Please ensure you answer the "how" question, i.e., how is the step performed?

Done.

- 11. Software steps must be more explicitly explained ('click', 'select', etc.). Please add more specific details (e.g. button clicks for software actions, numerical values for settings, etc.).
- 12. 1.4: How do you check the depth of anesthesia? How do you sterilize the site of surgery? Do you shave the surgical site?
- 13. 1.6: How is this done?
- 14. 2.1: Please make action steps in imperative tense using complete sentences.
- 15. 2.2, Note: Some of the details of this note can be moved to the discussion.
- 16. 2.2.1: How were the slices made?
- 17. 3: Please write in action steps only, discussion about the action can be included in the discussion section.
- 18. 3.2.1: We cannot have paragraph of texts in the protocol section, please make substeps instead.
- 19. 3.3: Please include how do you perform each step.

Changed as requested.

20. There is a 10-page limit for the Protocol, but there is a 2.75-page limit for filmable content. Please highlight 2.75 pages or less of the Protocol (including headings and spacing) that identifies the essential steps of the protocol for the video, i.e., the steps that should be visualized to tell the most cohesive story of the Protocol.

The 2.7 pages are highlighted in azure.

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- 22. As we are a methods journal, please revise the Discussion to explicitly cover the following in detail in 3-6 paragraphs with citations:
- a) Critical steps within the protocol
- b) Any modifications and troubleshooting of the technique

- c) Any limitations of the technique
- d) The significance with respect to existing methods
- e) Any future applications of the technique.

We have added respective text to the Discussion.

- 23. Please include all the Figure Legends together at the end of the Representative Results in the manuscript text. Please remove the legends from the figure.

 Done.
- 24. Please include a table of the essential supplies, reagents, and equipment. The table should include the name, company, and catalog number of all relevant materials in separate columns in an xls/xlsx file.

Done.

25: Please reword lines 23-32 in the introduction section and 2.2. note: first sentence as it matches with previously published literatures.

Done.

Reviewer #1:

The description of the surgery and a detailed list of reagents and materials is lacking (glass pipettes, drugs, etc), but this can be easily fixed.

Is now added.

I also am asking that figure 6 include raw data of the PPR experiments, not just the summary graphs!

The requested original traces from PT/IT synapses for the summary graphs of Fig. 6 are now added to Fig. 3. A respective reference is included in the text and the legend to Fig. 6.

Why can't regular epifluorescence be used? why only a scanning laser? or am I missing something? please clarify.

It's the balance between sufficient excitation and bleaching. Focal illumination protects the fluorescence out of the region of interest. The use of a focal 1P laser places the present approach somewhere between regular epifluorescence and 2PLSM. It is the method of choice if the signal obtained with a genetically encoded glutamate indicator (GEGI) at rest is too low for detection with 2P optics.

How do you take the optical resolution limit into consideration (I'd imagine you should never have more than 5 pixels per micron)?

With the 63x objective and the Andor Zyla one may have 10 pixels per μ m. It is possible to work with such resolution, but our recommendation is to work with binning 2x2.

Can you please discuss the prospects of this becoming and in vivo (e.g., 2PLSM) technique? We have mentioned in the Discussion that the present approach can be applied to in vivo brain preparations. To explain more details would be a topic for another JoVE paper. Off-line: Single

synapse imaging in intact mice is certainly possible with genetically encoded Ca sensors (Reynolds et al 2018). We hope that the addition of Ca imaging (and illumination with a second laser) will facilitate the search of presynaptic varicosities expressing a fast GEGI. Using the present approach (without a Ca indicator) we have performed *in vivo* recordings from the contralateral premotor cortex. At this point we can already say that iGluu imaging of single glutamatergic terminals appears to be feasible in the superficial layers of the (motor) cortex of anesthetized mice. A 2PLSM has now become available as well, but the experiments are only starting. At this point we can only refer to the exquisite method description of Dürst et al in Nat Prot 2019 who describe experiments in cultures from neonatal hippocampi. However, after becoming aware of the influence of astrocytes on glutamatergic synaptic transmission cultures are not that attractive. All further efforts should go towards single synapse imaging in the intact brain.

p1 l16 - missing space in A25%

Corrected.

p1 l17 - remains no remain

Corrected

p2 l12 - Because not Since

Corrected

p3 I4 missing space in proper name

Corrected

p3 l23 are these adenoviruses or adenoassociated viruses as mentioned below (e.g., l 31)?

Corrected

p3 I 29 do you need to cut the pipette tips after the pull? please say something about the pulling parameters (e.g., is it a single pull?). Also, the exact kind of glass you use (OD, ID and including supplier) should be mentioned. Also what puller (supplier) do you use?

Text added.

p3 l32 are you sure you do not maintain your viruses on ice until the actual injection? why do you leave it at RT?

We rely on experiments of the Charité viral core facility technicians who specifically investigated this issue. Their recommendation is to avoid any freezing of virus preps if they are used within the next 6 months. After virus production we keep part of the virus at 5 °C and take the respective tube out of the fridge just before the mouse is anesthetized. The other part is aliquoted and stored at -80 °C. In that case we do not use ice till the final use after unfreezing. Here we report a working protocol, but cannot discuss the pros and cons of virus storage recommendations.

p3 l37 what stereotaxic frame did you use? (supplier)
Added to Table 1.

p4 I1 what about shaving hairs and sterilization with alcohol and/or betadyne?

Added to Text.

p4 I2 - do you inject manually or with an injection system? Please explain how you monitor the injection rate. Details of how you mount the pipette would also help (i.e., is there use of oil for a pneumatic effect?)

Added to Text.

p4 I4 guarantees instead of secures

Corrected.

p5 l10 what stimulator did you use?

Added to Table 1.

p5 l21 please say which objective you used (Zeiss?).

Added to Table 1.

p 6 II12-14 Please discuss how imaging would be done in conjunction with optogenetics (eg., issue of wavelengths etc). I can foresee two issues: 1. accidentally activating the opsin; and 2. bleaching the fluorescence while activating opsins with light. (btw channelrhodopsin should be without a space).

The use of channelrhodopsin can well be combined with single bouton iGlu*u* imaging if the former is activated by focal stimulation at a distant site, e.g. the contralateral cortex. In the described experiments we stimulate close to the monitored varicosity and therefore refrained from the use of channelrhodopsins. The text has now been changed in accordance.

p 6 l17 please explain above how you pull the stim electrode? what about using theta class or a metal electrode? please discuss.

Done.

p 6 I 22 what concentration of TTX would you use?

Done.

p 9 I 6 how can you be sure that the explanation is that the other terminals do not express the iGluu, perhaps they do but they are simply outside the FoV or depth of field so that you don't see them clearly. I think that is just as likely an explanation as the other fibers not expressing the iGluu. In fact, if you have a spurious signal doesn't that mean that it must be from terminals the DO express iGluu? please explain.

That paragraph needed a complete re-writing, we agree. See new version on p9 Il23-29.

p 9 l 19 change als to also

Done.

p10 | 11 and not andf

Done.

p 10 l 22 missing space in TauDwas

Done.

fig. 4 caption. not clear what is meant by "the brightest pixels at rest are moved". what does "moved" mean in this context?

Text changed to make the point clearer. See p11, Il41-42.

At first sight figure 7E, right (HOM) is misleading - it looks as large an effect as on the left (wt) but you say that it's not significant. Are you sure there isn't a trend? Although, I can't imagine how TBOA could shorten TauD in homozygous Q175s? in any event, if it's not significant, please stress this in the figure legend.

Yes, we agree. We have now removed the confusing results from the TBOA experiments in HOM altogether.

Reviewer #2:

1) The dissociation constant for iGluu is mention to be 2.1 ms (page 3). Glutamate Tau decay in WT slices amounts to about 2.0 ms (Fig. 7).

We have made changes on p3 II9-13 to make this point more clear. The off-rate of 2.1 ms in the Helassa et al 2018 paper refers to their experiments in a stopped-flow device at a temperature of 20 °C. The extrapolated value for 34 °C was 0.68 ms. It can therefore be assumed that the decay time constant of the iGlu fluorescence is determined by diffusion and uptake rather than the kinetic characteristics of the sensor. The mean TauD on WT was 3.4 ms. The example chosen in Fig. 7 is probably one with a particularly fast decay, but still longer than the 0.68 ms estimated for the sensor by Helassa et al 2018.

Does a two-exponential fit provide better approximation?No.

2) The authors refer to astrocyte-mediated glutamate clearance in all parts of the manuscript and show the TFB-TBOA-induced changes data (Fig.7 E). However prolongation of Tau decay might be a result of reduced number of EAATs and/or altering of astrocyte morphology (and, as a result, change of diffusion rate). It is very difficult to separate these changes, but the authors could discuss this point when writing about "glutamate spread" (Fig. 3)?

We admit that the use of the term "clearance" in synaptic physiology is somewhat ambiguous. In general pharmacology and in renal physiology, "the clearance is a pharmacokinetic measurement of the volume of plasma from which a substance is completely removed per unit time; the usual units are mL/min." In most cases this is a virtual indicator. As we do not have exact estimates of the involved volumes, any change in the glutamate concentration (iGlu_u fluorescence) next to the releasing terminal may as well reflect a change both in uptake and in volume of perisynaptic space. But the lack of disease related difference in the "Spread Velocity" (Dvorzhak et al 2019, Fig. 4M). argues against a change in volume/diffusion. Furthermore, a genetically induced recovery of astrocytic uptake EAAT2 function (Hirschberg, Dvorzhak et al, unpublished) resulted in complete

recovery of $iGlu_u$ TauD. We therefore believe that in the P-type corticostriatal synapses glutamate clearance is indeed strongly determined by the rate of glutamate uptake.

To illuminate the complex interrelationships between glutamate uptake and clearance would require qualified modeling and should be subject of a more theoretical paper. For the present communication we prefer to limit ourselves to what is said in the present text, but see the additions on p10, ll33-42.

3) page 7, lines 38-40 "8) ... Note: After 10 trials, iGluu is fully bleached. The maximal illumination time with the 1 photon laser system and the described setting is approximately 2 s." Is the bleaching linear? As the peak amplitude of iGluu probe is proposed to reflect the near-synaptic glutamate concentration, the question arises whether the bleaching would distort the signal. Please discuss.

This is one of the most critical aspects of the method. We have added some text to underline what the method in its present implementation can give or not (p14, ll10-31).

Fig. 6 shows the paired-pulse plasticity at distinct synaptic varicosities. AP-induced glutamate release has a probabilistic nature, i.e. to obtain a representative paired-pulse ratio, several trials have to be applied. Values in Panel C represent values derived from how many trials?

The values in panel C represent the averages of from the first 3 trials in each tested bouton. This information has now been added to the legend.

- 1) Abstract "..single synapse resolution can be achieved in fully mature acute brain slices.." What is a fully mature slice? Please rephrase.

 Done.
- 2) page 3, line 1-3 "Here we describe a new approach which allows one to 1 evaluate single synapse glutamate clearance in its relation to the amount of the released neurotransmitter after expression of the genetically encoded glutamate sensor IGluu in the presynaptic plasma membrane." This sentence sounds as if the release would be dependent on the IGluu expression. Please rewrite.

Done.

- 3) page 4, line 2 "Place the pipette in the cortex..." At which angle? Done.
- 4) page 5, top "...[Glu] levels below 100 nM. Accordingly, any sensor of glutamate, and especially a low affinity sensor like iGluu will be rather dim in the absence of synaptic glutamate release." Is there any fluorescent signal from iGluu at resting conditions?

Yes. See Fig. 3 "Before stimulation" and Chapter 2.2.

Fig. 3B shows that F0 is about 500 a.u. Is this a representative value? How strong (in a.u.) is the background fluorescent after the suggested correction.

This is a representative value. After subtraction of autofluorescence the mean background fluorescence F is typically 400-700 a.u. See new text p6 l17.

5) page 9, line 10. "It als decays more slowly". Please correct. Done.

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