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Ballistic Labeling of Pyramidal Neurons in Brain Slices and in Primary Cell Culture --Manuscript Draft--

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1 TITLE:

2 Ballistic Labeling of Pyramidal Neurons in Brain Slices and in Primary Cell Culture

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KEYWORDS:

ballistic labeling, dendritic spine, pyramidal neuron, hippocampus, confocal imaging, rat

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SUMMARY:

We present a protocol to label and analyze pyramidal neurons, which is critical for evaluating potential morphological alterations in neurons and dendritic spines that may underlie neurochemical and behavioral abnormalities.

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LONG ABSTRACT:

It has been reported that the size and shape of dendritic spines is related to their structural plasticity. To identify the morphological structure of pyramidal neurons and dendritic spines, a ballistic labeling technique can be utilized. In the present protocol, pyramidal neurons are labeled with DilC18(3) dye and analyzed using neuronal reconstruction software to assess neuronal morphology and dendritic spines. To investigate neuronal structure, dendritic branching analysis and Sholl analysis are performed, allowing researchers to draw inferences about dendritic branching complexity and neuronal arbor complexity, respectively. The evaluation of dendritic spines is conducted using an automatic assisted classification algorithm integral to the reconstruction software, which classifies spines into four categories (i.e., thin, mushroom, stubby, filopodia). Furthermore, an additional three parameters (i.e., length, head diameter, and volume) are also chosen to assess alterations in dendritic spine morphology. To validate the potential of wide application of the ballistic labeling technique, pyramidal neurons from in vitro cell culture were successfully labeled. Overall, the ballistic labeling method is unique and useful for visualizing neurons in different brain regions in rats, which in combination with sophisticated reconstruction software, allows researchers to elucidate the possible mechanisms underlying neurocognitive dysfunction.

INTRODUCTION:

In 2000, Gan et al. described a rapid labeling technique for individual neurons and glia in the nervous system that combined various lipophilic dyes, allowing for the simultaneous labeling of many brain cells with different colors^{1,2}. More recently, a ballistic labeling technique was described by Seabold et al.³ that introduced fluorescent dyes (Dil) into the neurons of brain slices. A versatile staining technique, ballistic labeling is appreciated for its ability to be utilized in multiple animal species and across a wide range of ages. Furthermore, it can be combined with immunostaining to identify subpopulations of brain cells³. Compared with traditional techniques (e.g., Golgi-Cox silver impregnation, microinjection)⁴, ballistic labeling affords an opportunity to more clearly distinguish morphological characteristics, including dendritic spines, a feature that is critical for drawing inferences about neuronal complexity and synaptic connectivity⁵.

Excitatory pyramidal neurons are characterized by a single, large apical dendrite, multiple shorter basal dendrites, and thousands of dendritic spines⁶. Pyramidal neurons are found in multiple brain regions related to higher order cognitive processing, including the prefrontal cortex (PFC) and hippocampus. In the PFC, pyramidal neurons are observed in layers II/III and layer V, with each exhibiting unique morphology. Specifically, pyramidal neurons in layer II/III of the PFC have a shorter apical dendrite and less branching than pyramidal neurons in layer V⁶. Within the hippocampus, pyramidal neurons are located in both the CA1 and CA3 regions, with each displaying distinct morphologies. Specifically, pyramidal neurons in the CA1 region exhibit a more distinctive apical dendrite, with branching occurring farther from the soma, relative to the CA3 region⁶.

Dendritic spines on pyramidal neurons in both the PFC and hippocampus are the primary site of excitatory synapses⁷. Morphological characteristics of dendritic spines, which are classically characterized into three primary categories (i.e., thin, stubby, or mushroom8), have been related to the size of the excitatory synapse⁹. Thin spines, characterized by a long, thin neck, small bulbous head, and smaller postsynaptic densities, are more unstable and develop weaker connections. However, mushroom spines, which have a larger dendritic spine head, are recognized for forming stronger synaptic connections, an effect resulting from their larger size. In sharp contrast, stubby spines are devoid of a spine neck, exhibiting an approximately equal head and neck volume ratio⁸. Within the hippocampus, branched spines may also be observed, whereby the spine has multiple heads that emerge from the same dendritic spine neck¹⁰. Therefore, the morphological changes of dendritic spines could reflect functionality and structural capacity. Furthermore, studies have demonstrated that the size and shape of dendritic spines relates to their structural plasticity, leading to the idea that small spines are involved in learning and attention, whereas larger, more stable spines, are involved in long-term processes, including memory¹¹. Additionally, the distribution of dendritic spines along the dendrite may be associated with synaptic connectivity^{5,12}.

 Thus, the present methodological paper has three goals: 1) Present our protocol for ballistic labeling, which has been utilized with a success rate (i.e., neurons meeting selection criteria and appropriate for analysis) of 83.3%^{5,12,13} and across multiple brain regions (i.e., PFC, nucleus accumbens, hippocampus); 2) Demonstrate the generalizability of the technique and its

application to neurons grown in vitro; 3) Detail the methodology utilized in neuronal reconstruction software and the inferences that can be drawn from such data.

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PROTOCOL:

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All animal protocols were reviewed and approved by the Animal Care and Use Committee at the University of South Carolina (federal assurance number: D16-00028).

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1. Preparation of Dil/Tungsten bead tubing

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99 1.1. Dissolve 100 mg of polyvinylpyrrolidone (PVP) with 10 mL of ddH₂O. Vortex the PVP solution lightly.

101

102 1.2. Fill the tubing with the PVP solution (see **Table of Materials**) and leave it for 20 min. Then, expel the PVP solution through the other end of the tubing using a 10 mL syringe.

104

1.3. Combine 170 mg of tungsten microcarrier beads with 250 μL of methylene chloride.
 Vortex the tungsten bead suspension thoroughly.

107

108 1.4. Combine 6 mg of lipophilic DilC18(3) dye with 300 µL of methylene chloride. Vortex the DilC18(3) dye solution thoroughly.

110

111 NOTE: Perform steps 1.3–1.4 in a fume hood.

112

113 1.5. Pipette 250 μL of the tungsten bead suspension onto a glass slide. Wait for the suspension to air dry (~3 min).

115

1.6. Add 300 μ L of DilC18(3) dye solution on the top of the tungsten bead suspension layer.

Mix the DilC18(3) dye solution and tungsten bead suspension thoroughly with a pipette tip and allow the mixture to air dry (~3 min).

119

120 1.7. Once dried, use a razor blade to split the mixture into two 1.5 mL centrifuge tubes. Fill the tubes with water.

122

123 1.8. Sonicate in a water bath (Amp I, 100%) until homogenous (~5 min). Ensure that the tip of the sonicator lies directly on the tubes with the bead suspension.

125

126 1.9. Combine the two 1.5 mL of homogenized mixture into a 15 mL conical tube. Sonicate the mixture for another 3 min.

128

129 1.10. Draw the tungsten beads-DilC18(3) dye mixture into the PVP-coated tubing using a 10 mL syringe. Feed the tubing into the preparation station (see **Table of Materials**).

132 1.11. Rotate for 1 min on the preparation station. Carefully remove all the water from the tubing using a 10 mL syringe.

134

135 1.12. Turn on the nitrogen gas and adjust the nitrogen flow to approximately 0.5 Liters per minute (LPM), rotate the tubing in the preparation station, and dry with nitrogen for 30 min.

137

1.13. Remove the tubing from the preparation station and cut into 13 mm lengths (matching the loading size of the cartridge) using a tubing cutter. Keep the 13 mm lengths in scintillation vials in the dark.

141

2. Preparation of brain sections

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NOTE: Adult male F344/N rats were pair housed in a controlled environment under a 12/12 light:dark cycle with ad libitum access to food and water. All animals were cared for using guidelines established by the National Institutes of Health in the Guide for the Care and Use of Laboratory Animals.

148

149 2.1. Deeply anesthetize rats using 5% sevoflurane.

150

2.2. Proceed to the following step when the rats are not responsive to noxious stimuli and reflexes are absent.

153

2.3. Secure the rat in a supine position inside a chemical fume hood.

155

2.4. Make an incision through the skin along the thoracic midline. Separate the diaphragm and
 open the chest with scissors. Insert a 20 G × 25 mm needle into the left ventricle.

157 158

2.5. Immediately cut the right atrium with scissors. Perfuse 50 mL of 100 mM PBS with 5 mL/min of flow rate. Perfuse 100 mL of 4% paraformaldehyde buffered in 100 mM PBS.

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2.6. Remove the entire rat brain right after perfusion.

163

2.7. Postfix the entire brain for 10 min with 4% paraformaldehyde.

165166

NOTE: Do not postfix in 4% paraformaldehyde for more than 30 min, because it will affect the labeling.

167 168

2.8. Cut 500 μm thick coronal sections using a rat brain matrix (see **Table of Materials**). Make
 the first cut and keep the blade in place. Make a second cut using a second blade and vertically
 remove the first blade, keeping the tissue on the blade surface.

172

2.9. Place the brain slices in a 24 well plate with 1 mL of 100 mM PBS in each well. Repeat this process until all slices have been cut.

3 .	Ballistic labeling and visualization of brain sections
3.1.	Remove the PBS from each targeted well.
3.2. applio	Load the cartridge with a piece of Dil/Tungsten bead tubing and place it into the rator.
<mark>3.3.</mark> heliur	Put a piece of filter paper between two mesh screens. Connect the applicator to the n hose. Adjust the output pressure of helium to 90 pounds per square inch (psi).
3.4. 1.5 cr	Place the applicator vertically by hand on the center of the targeted well at a distance of n between the sample and the mesh screen. Fire the Dil/Tungsten beads tubing.
NOTE	: Be sure to remove all PBS from the targeted wells before shooting.
3.5. Dil/Tu	Load the cartridge with the next Dil/Tungsten bead tubing. Continuously fire the ingsten beads from the tubing on the remaining slices.
3.6. Do no	Fill the 24 well plate with 100 mM of PBS. Wash with 500 μ L of fresh 100 mM of PBS 3x. It let the slices flip over while washing with PBS.
3.7.	Add 500 μL of fresh 100 mM PBS and keep the slices at 4 °C in the dark for 3 h.
3.8.	Transfer the brain slices onto glass slides using a fine brush.
NOTE	: Three brain sections can be transferred onto each glass slide.
3.9. 50 mr	Immediately add 1 mL of antifade mounting medium onto each section. Place a 22 mm \times m coverslip over the brain sections. Dry the glass slides in the dark for 2 days.
3.10.	Turn on the confocal microscope system and switch to a 60× objective.
•	Set the confocal microscope system to have a magnification of $60\times$ (A/1.4, oil) and a Z-interval of 0.15 µm (pinhole size 30 µm, back-projected pinhole radius 167 nm). Use a 543 avelength to acquire images of the neurons of interest.
<mark>3.12.</mark> morp	Obtain Z-stack images for the targeted neuron type based on brain region boundaries and hological characteristics of neurons.
NOTE	: Acquire at least three images from each animal.

4. Use of the methodology with cell culture

Isolate primary cortical neurons from F344/N rats at postnatal day one using previously 219 220 reported methodology¹⁴.

221

222 Culture primary cortical neurons in a 35 mm glass bottom dish for one week. Change half of the culture medium with fresh neuron growth medium at third day after isolation. Wash the 223 224 glass bottom dish 2x with 1 mL of 100 mM PBS.

225

226 **4.3.** Fix with 4% paraformaldehyde for 15 min at room temperature. Repeat steps 3.2–3.6 to 227 ballistically label the cells.

228

229 Wash with 1 mL of 100 mM PBS 3x. Add 500 µL of fresh 100 mM of PBS and keep it at 4 **4.4.** 230 °C in the dark for 3 h.

231

232 4.5. Add 200 μL of antifade mounting medium.

233

4.6. Obtain Z-stack images for each targeted neuron using the parameters in step 3.10. 234

235 236

5. Neuronal analysis and dendritic spine quantification

237

238 5.1. Blind neurons using code numbers to prevent experimenter bias.

239

240 5.2. Establish selection criteria for neurons based on the brain region of interest.

241

242 NOTE: Selection criteria for neurons include continuous dendritic staining, low background, no 243 dye clusters inside of the cells, minimal diffusion of the Dil dye into the extracellular space, 244 correct morphology of pyramidal neurons (Figure 1).

245

5.3. Open neuronal reconstruction software (See the attached **Video 1**). 246

247

Load an image file by clicking on the "File Folder" image in the upper left-hand corner. 248 5.4.

249

250 5.5. Click "Soma" and mark the soma of the neurons on the image.

Click "Tree" and select "User-Guided".

251 252

5.6.

253

254 5.7. Trace all dendritic branches of interest.

255

NOTE: For pyramidal neurons, which are characterized by one apical dendrite and multiple basilar 256 257 dendrites, only the apical dendrite is traced. Make sure all connecting branches are attached to one another. 258

259

260 5.8. Click "Spine".

261

5.9. Define Detection Parameters and click "Detect All". 262

263

- NOTE: For brain slices, the consistent parameters utilized across brain regions are: 2.0 (Outer
- Range), 0.3 (Minimum Height), 100% (Detector Sensitivity), and 10 (Minimum Count). For cell
- culture, it is necessary to increase the Detector Sensitivity and decrease the Minimum Count.

267

268 5.10. Classify spines by selecting "Classify All".

269

NOTE: Dendritic spines are classified using an algorithm integral to the neuronal reconstruction software¹⁵.

272

5.11. Save tracing by selecting the **Disk Image** in the upper left-hand corner.

274

275 5.12. Perform neuronal and dendritic spine morphological analyses.

276

5.12.1. Open neuronal reconstruction quantitative analysis software (See attached **Video 2**).

278

5.12.2. Load image by clicking "File" and "Open Data File".

280

5.12.3. Click "Analysis" and "Branched Structure Analysis" to analyze neuronal morphology and dendritic spine morphology.

283

5.12.4. For neuronal morphology, click "Tree Totals" and select the box for "Dendrite Totals".

285

5.12.5. For dendritic spine morphology, click "Spines" and then select the box for "Spine Details".

287

5.12.6. Save output as a text (.txt) file by right-clicking the output table and selecting "Save to File".

290

291 5.12.7. Click "Analyze" and "Sholl Analysis".

292

293 5.12.8. Set the **Starting Radius** to 10 μm and the **Radius Increment** to 10 μm.

294

5.12.9. Click the box "Dendrites" and click "Display".

296

5.12.10. Save output as a text (.txt) file by right-clicking the **Output Table** and selecting "**Save to** File".

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6. Data analysis

301

302 6.1. Analyze the neuronal morphology (i.e., dendritic branching complexity) data.

- 6.1.1. Add the number of dendrites at each branch order and divide by the total number of
- dendrites. Multiply by 100 to calculate the relative frequencies for the number of dendrites at
- 306 each branch order.

6.2. Analyze the Sholl analysis data to examine neuronal arbor complexity and dendritic spine connectivity.

6.2.1. Calculate the mean and standard error of the mean for the number of intersections at each radius.

6.2.2. Sum the number of spines dependent upon spine type (i.e., thin, stubby, mushroom) at each radius and divide by the total number of spines for that spine type. Multiply by 100 to calculate the relative frequencies for the number of spines at each radius.

REPRESENTATIVE RESULTS:

In **Figure 2A**, the typical pyramidal neurons in the hippocampal region in the rat brain sections were identified by ballistic labeling technology, characterized by one large apical dendrite and several smaller basal dendrites around the soma. **Figure 2B** shows the neuron in the neuronal reconstruction quantitative analysis software after the soma was detected, dendritic branches were traced, and spines were detected. Subsequently, the data were analyzed using neuronal reconstruction quantitative analysis software, which provided an opportunity to assess the dendritic branching complexity (**Figure 2C**) and neuronal arbor complexity (**Figure 2D**).

In **Figure 2C**, we utilized the centrifugal branch ordering method, collected from the "Tree Totals" output, to count the number of segments traversed along each dendrite and assigned branch order. The relative frequency of segments at each branch order was examined for branch orders 1–15. When shifts in the distribution of dendritic branches were observed between groups, alterations in dendritic branching complexity could be inferred. Furthermore, a Sholl analysis was conducted as a complementary measure of neuronal arbor complexity, whereby the number of dendritic intersections occurring every 10 μ m from the soma was quantified in each sample section (**Figure 2D**). When shifts in the number of dendritic intersections were observed between groups, alterations in neuronal arbor complexity could be inferred.

Morphological changes in dendritic spines could be assessed using length (μ m), head diameter (μ m), and volume (μ m³), as seen in **Figure 3A–B**. Furthermore, spines were classified using the automatic assisted classification system in the neuronal reconstruction software. The relative frequency of the number of spines between each radius was examined for thin, mushroom, and stubby spines. Given our understanding of which spine types form stronger synaptic connections (i.e., mushroom relative to stubby) and neurotransmitter afferents, shifts in the distribution of spines along the neuron can indicate synaptic connectivity.

Furthermore, we validated the utility of the ballistic labeling technique on a primary pyramidal neuron in cell culture. First, we cultured primary hippocampal neurons at postnatal D1 (day 1) on a cell culture plate coated with poly-L-lysine for 2 weeks or until approximately 70% confluency. Then the samples were fixed with 4% PFA for 15 min and washed 2x with PBS. Beginning at step 3.1 of the present protocol, we ballistically labeled and imaged the primary pyramidal neurons from the hippocampus. Data showed stabilized labeling and identified pyramidal neurons based

on the triangle shape of the soma and large apical dendrite (**Figure 4A**). Utilization of neuronal reconstruction software to investigate dendritic spines of primary pyramidal neurons grown in cell culture offers opportunities similar to those in the rat brain. Example outcomes are illustrated for the distribution of thin dendritic spines (**Figure 4B**) and dendritic spine length measured in µm (**Figure 4C**). However, it is noteworthy that the primary pyramidal neurons grown in cell culture had less dendritic branching, precluding the assessment, at least in this example, of dendritic branching complexity and neuronal arbor complexity.

FIGURE LEGENDS:

Figure 1: Selection criteria utilized for pyramidal neurons in the medial prefrontal cortex labeled using ballistic labeling technology. (A) A representative confocal image (60x) of a well-labeled pyramidal neuron from the medial prefrontal cortex. A single pyramidal neuron with a clear soma and apical dendrite included continuous, bright dendritic staining with low background. (B–C) A representative confocal image (60x) of a pyramidal neuron from the medial prefrontal cortex with light staining at the more distal branches (B) and high background (C). (D) A representative confocal image (60x) of a labeled neuron from the medial prefrontal cortex (based on Bregma coordinates) that has flawed morphological characteristics.

Figure 2: The labeling of pyramidal neurons in the hippocampus (HIP) using ballistic labeling technology and neuroanatomic assessments. (A) Three representative confocal images (60x) of pyramidal neurons labeled by ballistic tungsten beads. (B) The assessment of neuronal morphology: dendritic branch order analysis and Sholl analysis. The traced image of the dendritic spine in which the spine morphology was also identified using dendritic spine analysis software. (C) Branch order analyses utilized to examine the relative frequency of dendritic branches at different branch orders. (D) The numbers of dendritic intersections every 10 μm from the soma were assessed using Sholl analysis as a measure of neuronal arbor complexity. Data are described as relative frequencies of the entire dataset (C) or fit with 95% confidence intervals (D).

Figure 3. The assessment of dendritic spine morphology. (A–B) The distribution of dendritic spines illustrated as a function of spine type (i.e., thin, stubby, mushroom). Additional dendritic spine parameters were also analyzed (i.e., length, volume, head diameter) as an assessment of dendritic spine morphology. Data are illustrated as relative frequencies of the entire dataset.

Figure 4. The labeling of primary cortical neuron in vitro using ballistic labeling technology. (A) Representative confocal images (60x) of primary cortical neurons labeled by ballistic tungsten beads in vitro. Example results similar to those acquired from ballistic labeling in brain slices are shown for the distribution of thin dendritic spines (B) and length (C). Data are illustrated as relative frequencies of the entire dataset.

Video 1: Procedures of neuronal tracing and dendritic spine detection.

Video 2: Procedures of data collection and output for quantitative analysis.

DISCUSSION:

In this protocol, we describe a versatile labeling technique for neurons from both rat brain and those grown in vitro. Furthermore, we report the methodology for utilizing neuronal reconstruction software and neuronal reconstruction quantitative analysis software to assess neuronal morphology and dendritic spines. The assessment of neuronal morphology and dendritic spines provides an opportunity to determine alterations in dendritic branching complexity, neuronal arbor complexity, dendritic spine morphology, and synaptic connectivity.

When conducting the protocol, researchers should pay special attention to a few steps. First, post-fixing in 4% PFA for too long will damage the integrity of the lipophilic membrane and cause the dye to leak outside of the cells. Second, compared to the specificity of ballistic labeling in brain slices, which targets only neurons, labeling of primary cortical neurons in vitro introduces nonspecific labeling of the glia because ballistic labeling in brain slices is not specific for a type of neuron (i.e., pyramidal neuron, medium spiny neuron, granule cell). Thus, Bregma coordinates, morphological assessments, or specific cell markers should be combined with the ballistic labeling method. Third, the thickness of the brain slices can be between 200–500 µm; for best results it should be optimized. Fourth, the efficiency of labeling and dye penetration is related to many factors, such as helium pressure, incubating times after ballistic application, the Dil/Tungsten beads, the distance between the mesh screen and the surface of brain slices, etc. The protocol should be optimized for each study. Fifth, large clumps or clusters of ballistic dye coated tungsten beads during preparation must be avoided, because clumps would not allow individual neurons to be distinguished. We also determined that Dil diffuses throughout individual neurons more completely than DiO in this ballistic methodology.

Nevertheless, compared with traditional labeling methods⁴, the ballistic labeling technique makes high resolution confocal imaging possible, allowing for the assessment of neuronal and dendritic spine morphology. Furthermore, neuronal reconstruction software utilizes an algorithm for automatic assisted classification of dendritic spines (i.e., thin, mushroom, stubby), branch ordering measurements, classical Sholl analysis, and measurement of morphological features of dendritic spines, such as length (μ m), head diameter (μ m), and volume (μ m³). The quantification of multiple neuronal parameters affords an opportunity to better understand the mechanisms underlying neurocognitive dysfunction.

Overall, the ballistic labeling method allows visualization of neuronal structures in different brain regions of the rat and in cell culture, which is important to elucidate the possible mechanisms underlying neurocognitive dysfunction. In the present study, we present a method to label pyramidal neurons by a ballistic labeling technique. Furthermore, combined with neuronal reconstruction software, we demonstrated the ability to examine neuronal and dendritic spine morphology in hippocampal pyramidal neurons. Group differences in neuronal and/or dendritic spine morphology provide an opportunity to understand the mechanisms underlying neurocognitive dysfunction.

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DISCLOSURES:

None of the authors have conflicts of interest to declare.

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Figure 1

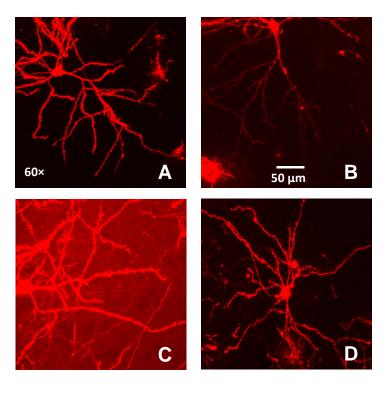


Figure 2

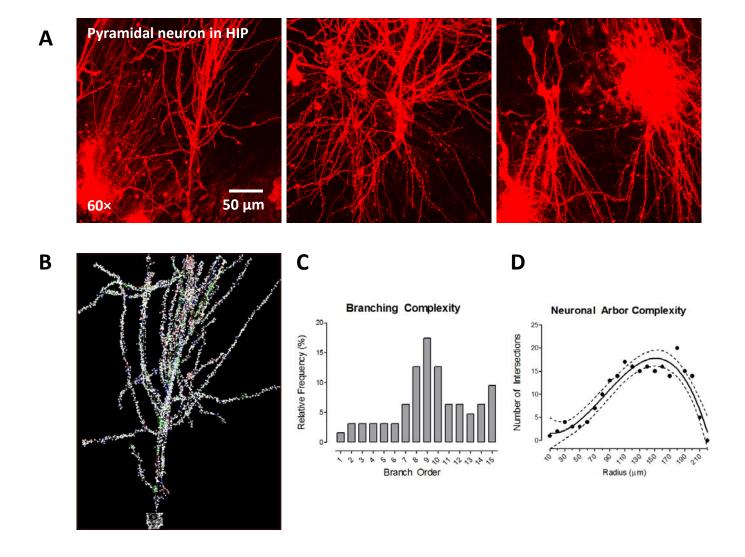


Figure 3

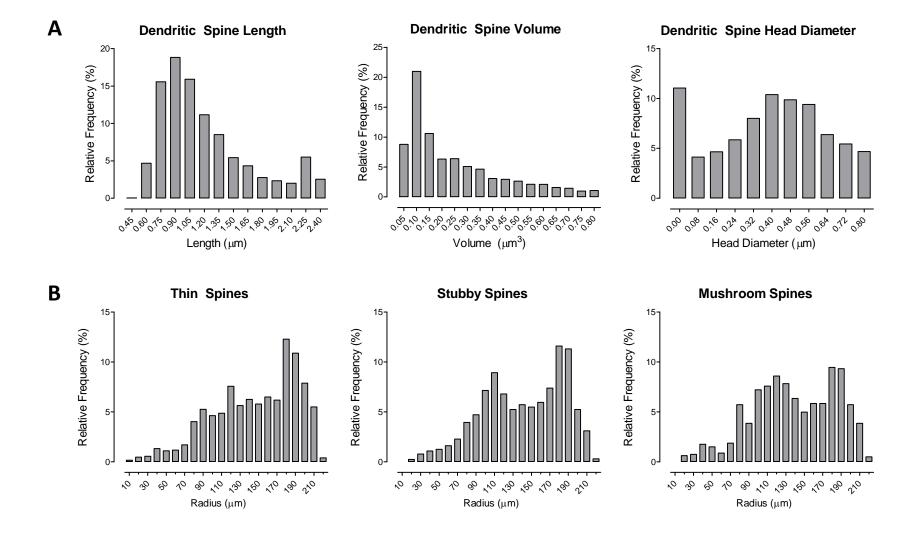
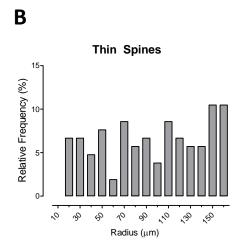
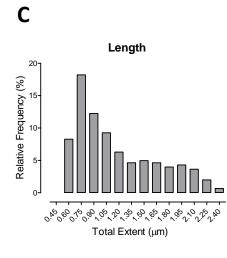


Figure 4







Video or Animated Figure

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Name of Material/ Equipment	Company	Catalog Number
20Gx25mm PrecisionGlide needle	BD	305175
24-well cell culture plate	Costar	3562
35 mm Glass Bottom Dishes	MatTek Corporation	P35G-1.5-20-C
Antibiotic-Antimycotic solution	Cellgro	30004CI
B-27 supplement	Life Technologies	17504-044
Barrel liner	BIO-RAD	165-2417
Borax	Sigma	B9876
Boric acid	Sigma	B0252
Cartridge holder	BIO-RAD	165-2426
Confocal imaging software	Nikon	EZ-C1
Confocal microscope	Nikon	TE-2000E
Cover glass	VWR	637-137
DilC18(3)	Fisher Scientific	D282
DMEM/F12 medium	Life Technologies	10565-018
Dumont #5 Forceps	World Precision Instruments	14095
Dumont #7 Forceps	World Precision Instruments	14097
F344 rat	(Harlan Laboratories, Indianapolis, IN)	
Glucose	VWR	101174Y
GlutaMax	Life Technologies	35050-061
HBSS	Sigma	H4641
Helios diffusion screens	BIO-RAD	165-2475
Helios gene gun kit	BIO-RAD	165-2411
Helios gene gun system	BIO-RAD	165-2431
Helium hose assembly	BIO-RAD	165-2412
Iris Forceps	World Precision Instruments	15914
Iris Scissors	World Precision Instruments	500216
Methylene chloride	Fisher Scientific	D150-1
Neurobasal medium	Life Technologies	21103-049
Neurolucida 360 software	mbf bioscience	
Paraformaldehyde	Sigma-Aldrich	158127-500G
Paraformaldehyde	Sigma	P6148
Poly-L-Lysine	Sigma	P9155

Polyvinylpyrrolidone	Fisher Scientific	5295
ProLong Gold antifade reagent	Fisher Scientific	P36930
Rat brain matrix, 300 - 600g, Coronal, 0.5mm	Ted Pella	15047
Sevoflurane	Merritt Veterinary Supply	347075
Sodium Bicarbonate	Life Technologies	25080
SuperFrost Plus Slides	Fisher Scientific	12-550-154%
Syringe kit	BIO-RAD	165-2421
Tefzel tubing	BIO-RAD	165-2441
Trypsin-EDTA	Life Technologies	15400-054
Tubing cutter	BIO-RAD	165-2422
Tubing Prep station	BIO-RAD	165-2418
Tungsten M-25 Microcarrier 1.7 μm	BIO-RAD	165-2269
Vannas Scissors	World Precision Instruments	500086

Comments/Description

100X 50X

version 3.81b

100X 10X

dendritic spine analysis

mounting medium

January 3, 2020

Dr. Vineeta Bajaj,

Review Editor

Journal of Visualized Experiments

One Alewife Center, Suite 200,

Cambridge, MA 02140

Dear Dr. Vineeta Bajaj,

Thank you very much for you and reviewers' recommendations for our manuscript. We revised the manuscript according to the comments provided by you and reviewers. We are now resubmitting this revised manuscript to *JoVE*. Please feel free to let us know if you have any questions.

Sincerely,

Hailong Li, MD PhD

Research Associate

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Tel: (803)777-3568

Editorial comments:

1. The editor has formatted the manuscript to match the journal's style. Please retain and use the attached version for incorporating changes.

Answer: It has been done.

2. Please address all the specific comments marked in the manuscript.

Answer: It has been done.

3. For each step, please ensure you answer the "how" question, i.e., how is the step performed?

Answer: It has been done.

4. Once done please ensure that the highlighting is no more than 2.75 pages including headings and spacings.

Reviewer #1:

Reviewer 1 reply to R1.

I mostly satisfied with the last version of MS, some remaining questions and new comments are follows. To the Editor: may be it is possible to name the software used in the study, at least in Used Equipment chapter? The protocol steps in program are software-specific, and without mention the name whole second part of the protocol are become senseless.

5. Still have doubts what 10 min postfixation of whole rat brain (not the 300-500 mkm slices) is enough to adequately fix inner parts of the brain, esp. hippocampus. Did you made some experimental optimization of the time required, or just follow the Seabold et al. recommendations?

<u>Answer</u>: Actually, the inner parts of the brain are fixed adequately after the animal is perfused with 100 mL 4%PFA. We have tried different post-fixation times in our study, including 10 minutes, 20 minutes, 30 minutes and 1 hour. The quality of neuronal labeling in the current protocol is decreased when a 1 hour post-fixation time is used. However, there is no significant difference between 10 minutes, 20 minutes, and 30 minutes of post-fixation.

24. Is it possible to be sure you chosen a "typical neuron" for a brain? How you can comment the variability of morphology of pyramids in the same brain, not between-group of animals? For me between-group analysis made from only one arbitrary chosen neuron in brain looks weird. What the typical sample size (number of neurons per brain/structure) for such analysis in other morphological studies, is "gold standard" exists?

<u>Answer</u>: Neurons are randomly selected to prevent experimenter bias. The analysis of multiple neurons per animals violates the independent observation assumption that underlies many statistical approaches. In preclinical studies utilizing biological systems to model disease, many

researchers consider six animals per group to be an adequate sample size (Charan & Kantharia, 2013).

39. I don't see the mentioned paper in References and Discussion. BTW, Discussion is not contained any Refs at all — is it OK by the journal rules? One number for success rate mean nothing without explanation how it was calculated. If I got it correctly, in 83% of brains you were able to find at least 1 neuron? How many section do you did - at least 5 x 500 mkm for hippocampus I guess. All these details must be in MS, to allow the follower of your protocol to correctly estimate the chances of the successful staining and overall successful rate of the method.

<u>Answer</u>: We are able to successfully find, and image, at least 1 neuron in all of our brain slices. The success rate is based on the percentage of neurons meeting selection criteria. We have added these details to the manuscript.

Line 243 3.17 — seems unneeded, as it repeated again in 3.19.

Answer: It has been changed.

Line 245 3.18 — This is not step, it is chapter name for following steps 3.19-20. Or, alternatively this step must be after 3.20.

Answer: It has been changed.

Line 341 and later — Please clarify, which software used for this analysis — the same reconstruction soft, or some table editor.

Answer: It has been explained in video-2.

Line 427 — 'neurons', but only one visible in Fig. Please reconcile. Pages 20-21 in PDF file look strange — is some part of MS missing?

Answer: It has been changed.

Reviewer #4:

Manuscript Summary:

The authors describe methodology to label pyramidal neurons in brain slices and primary culture using ballistic delivery of the fluorescent dye DilC18(3). The authors demonstrate that images obtained from the ballistic method can be analyzed with neuronal reconstruction software to investigate neuronal and dendritic complexity and dendritic spine morphology. Overall, the protocol is well organized and the resulting manuscript would be a worthy addition to the JoVE library. Some minor concerns are noted that would strengthen the protocol.

Major Concerns: None noted.

Minor Concerns:

The first paragraph of the introduction is somewhat confusing as it is not clear if the technique referenced in Seabold et al, is the same or different from the biolistic technique in the present manuscript.

<u>Answer</u>: We have adapted the present protocol from the one originally reported by Seabold et al. We believe the novel aspect of our protocol is the focus on utilizing neuronal reconstruction software to analyze neuronal morphology and dendritic spines. To our knowledge, a protocol has not yet described how to utilize neuronal reconstruction software and analyze output to draw inferences about neuronal complexity and synaptic connectivity.

Protocol

Step 1.1; 1.15: Please clarify the required purity of the water to used (ie., dH20)

Answer: It has been changed.

Step 1.10; 1.13: Please indicate approximately how long it takes for the suspension to air dry

Answer: It has been changed.

Discussion:

The second paragraph discussing the steps to which researchers should pay special attention could be better organized. For example, point 3 begins with slice thickness but then switches to labeling efficiency and dye penetration. Sometimes a point begins with a sentence and sometimes a phrase. I suggest the following format: "When conducting the protocol, researchers should pay special attentions to a few steps. First,... Second,...Third,...Fourth,..."

Answer: It has been changed.

Lines 451-453: The authors state that, "We have also determined that Dil diffuses throughout individual neurons more completely, in comparison to DiO in this ballistic methodology." However, there are no images shown for DiO in the representative results.

Answer: It has been changed.

Fig 2: The term 'HIP' in panel A is not defined.

Answer: It has been changed.