

Analysis of Synapses in Cerebral Organoids

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Abstract

Cerebral organoids are an emerging cutting-edge technology to model human brain development and neurodevelopmental disorders, for which mouse models exhibit significant limitations. In the human brain, synaptic connections define neural circuits, and synaptic deficits account for various neurodevelopmental disorders. Thus, harnessing the full power of cerebral organoids for human brain modeling requires the ability to visualize and analyze synapses in cerebral organoids. Previously, we devised an optimized method to generate human cerebral organoids, and showed that optimal organoids express mature-neuron markers, including synaptic proteins and neurotransmitter receptors and transporters. Here, we give evidence for synaptogenesis in cerebral organoids, via microscopical visualization of synapses. We also describe multiple approaches to quantitatively analyze synapses in cerebral organoids. Collectively, our work provides sufficient evidence for the possibility of modeling synaptogenesis and synaptic disorders in cerebral organoids, and may help advance the use of cerebral organoids in molecular neuroscience and studies of neurodevelopmental disorders such as autism.

Keywords

cerebral organoids, human brain, stem cells, synapses, neurodevelopmental disorders, autism, synaptic deficits

Introduction

Cerebral organoids are stem-cell derived, three-dimensional (3D) neural-tissue-like structures, that recapitulate early (ca. first trimester) *in vivo* developing brain tissues. While an *in vitro* system, cerebral organoids perhaps provide the only portal to studies of human-specific brain development and disease in 3D. Cerebral organoids recapitulate lamination and organization of the human cortex¹, diversity of the cell types observed *in vivo* (such as neuronal subtypes, glial cells, and neuroepithelial and neuroprogenitor cells^{2,3}), and even the gene expression profiles observed during human fetal neocortical development as revealed by single-cell RNA-sequencing analyses⁴. The organoid technology is still in its rather early stages, but the promise it poses to the study of the human-specific developmental and disease neuroscience is enormous, which has encouraged ongoing efforts to further improve the technology and reduce its limitations, including attempts for circuitry reconstruction^{5–8}, in order to enhance the organoids' utility to study some aspects of human brain development or developmental disorders.

Human cerebral organoids enabled detection of disease-relevant phenotypes and characterization of disease mechanisms that could not be detected in traditional models^{1,9}, including *in vivo* mouse models or the human two-dimensional model (stem-cell-derived neurons). Some disease-associated gene mutations that are responsible for

severe phenotypes and devastating neurodevelopmental conditions in humans were found to exhibit no phenotype, or a marginal, barely significant, phenotype in the mouse brain. For example, mouse models for microcephaly-associated mutations (as in *CDK5RAP2* (CDK5 (cyclin-dependent kinase 5) related activator protein 2 or CDK5 regulatory subunit-associated protein 2)) failed to show the severe microcephalic phenotype associated with these mutations in human patients^{1,9,10}. Organoids were also used to investigate the pathological mechanisms of Zika virus infection and its relationship to dysregulation of neurogenesis and neuroinflammatory pathways and to microcephaly in humans^{11–14}. Moreover, human forebrain organoids derived from induced pluripotent stem cells (iPSCs) of patients carrying a mutation

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in *Disrupted-in-schizophrenia 1 (DISC1)*, a gene associated with multiple neuropsychiatric disorders such as schizophrenia, affective disorders and autism spectrum disorders (ASDs)^{15,16}, revealed a role of this gene in cell cycle progression in radial glial cells and neurogenesis¹⁷. Arguably, cerebral organoids were also reported to recapitulate a neurodegenerative disease model. For example, cerebral organoids derived from a familial Alzheimer's disease (FAD) patient's iPSCs showed aggregation of amyloid- β and hyperphosphorylated tau¹⁸. Moreover, cerebral organoids harboring a frontotemporal dementia (FTD)-associated tau mutation showed evidence of a CDK5-dependent tau phosphorylation and a decrease in synaptophysin levels¹⁹.

Thus far, cerebral organoids have mostly been used for studies that are focused on functions of the radial glial and neural progenitor cells (including the aspects of proliferation and neurogenesis)^{13,20–22,17} or on non-synaptic aspects of neuronal biology such as neuronal differentiation, neuronal migration, or gene expression and transcriptional pathways analyses^{7,8,23}. However, synaptic function is at the very center of neuronal functions and is probably the most important neuronal function^{24–26}. Moreover, analyzing synapses is essential to understanding the disease mechanisms of neurodevelopmental disorders that are caused by deficits in synapses^{27–30}. Despite its immense importance, thorough synapse analyses have not yet been performed in cerebral organoids. Scarcely, some Ca²⁺ imaging experiments and electrophysiological analyses were performed on organoids^{1,5}. While electrophysiological approaches, despite being highly tedious, provide important insights into synaptic connectivity, synaptic microscopical analyses (such as described here) complement the electrophysiological approaches by directly visualizing synapses and determining synapse counts^{31,32}.

Therefore, in this study, we attempted to visualize synaptogenesis in cerebral organoids, via confocal microscopy. We have tested numerous approaches for analyzing synapses, and concisely report here multiple assays that were shown to be compatible with cerebral organoids. We provide ample procedural details and technical tips to allow researchers to successfully reproduce these assays. Thus, we expect this study to help advance the field of study of neurodevelopmental disorders and synaptogenesis in the human cerebral organoid model.

Materials and Methods

Chemicals and Reagents

Materials used in generation of organoids were previously described in details³³. The following reagents were used for preparing the organoids for synapse analysis: Tissue-Tek® O.C.T. Compound (Cat # 25608-930; Sakura, Alphen aan den Rijn, The Netherlands), Tissue Path™ Disposable Base Molds (Cat # 22-363-553; Fisher Scientific, Hampton, NH, USA), Goat serum (Cat # 16210064; Invitrogen, Carlsbad, CA, USA), Vectashield 4',6-diamidino-2-phenylindole (DAPI)

mounting medium (Cat # H-1200; Vector Labs, Burlingame, CA, USA), Adhesive glass slides (Cat # I6172PLUS, Thermo Fisher, Waltham, MA, USA), Coverslips (Cat # 102222; Thermo Fisher), Kimwipes (Cat # 06-666; Fisher Scientific), and paraformaldehyde stock (32%) solution (Cat # 15714; Electron Microscopy Sciences (EMS), Hatfield, PA, USA).

Paraformaldehyde solution for tissue fixation (4% paraformaldehyde solution, pH 7.4) can be prepared by 1:8 dilution of the 32% stock solution in PBS pH 7.4; 4% paraformaldehyde solution is to be stored in amber-colored or light-proof containers at 4°C for 1–2 weeks.

The following solutions were used: Blocking Solution (5–10% goat serum, 0.1% BSA, 0.3% Triton X-100 in PBS), and Antibody Vehicle Solution (AVS) (1–2% goat serum, 0.1% Triton X-100 in PBS).

Primary Antibodies. The following primary antibodies were used: Synapsin 1 (SYN1) antibody, rabbit polyclonal antiserum (Cat # 106002; Synaptic Systems, Göttingen, Germany) was used at the dilution of 1:1000. Microtubule-Associated Protein 2 (MAP2) antibody, chicken polyclonal IgY (Cat # 188006; Synaptic Systems) was used at the dilution of 1:1000. Glutamate NMDA Receptor Subunit 1 (GluNR1) antibody, mouse monoclonal IgG (Cat # 114011; Synaptic Systems) was used at the dilution of 1:500. Vesicular Glutamate Transporter 1 (VGLUT1) antibody, Guinea pig polyclonal antiserum (Cat # AB5905; Millipore, Burlington, MA, USA) was used at the dilution of 1:1000.

Secondary Antibodies. Secondary antibodies conjugated with Alexa Fluor-488, -546 or -633 (Molecular Probes, Invitrogen) were used for confocal microscopy, as follows: Goat anti-rabbit IgG secondary antibody—Alexa Fluor-488 (Cat # A-11008; Invitrogen); Goat anti-mouse IgG secondary antibody—Alexa Fluor-546 (Cat # A-11030; Invitrogen); Goat anti-chicken IgY secondary antibody—Alexa Fluor-546 (Cat # A-11040; Invitrogen); Goat anti-Guinea pig IgG secondary antibody—Alexa Fluor-633 (Cat # A-21105; Invitrogen). Secondary antibodies were generally used at the dilution of 1:1000, except the anti-Guinea pig—Alexa Fluor-633 secondary antibodies, which were used at the dilution of 1:500.

Equipment

The following equipment was used: Liquid nitrogen tank and liquid nitrogen bucket, ImmEdge™ Hydrophobic Barrier Pen (Cat # 101098-065; VWR, Radnor, PA, USA), temperature-controlled microtome cryostat (Leica Research Cryostat, Cat # CM3050 S; Leica Biosystems, Buffalo Grove, IL, USA), Confocal microscope (Nikon, or Zeiss), NIS-Elements™ software (Nikon, Tokyo, Japan) and sterile forceps.

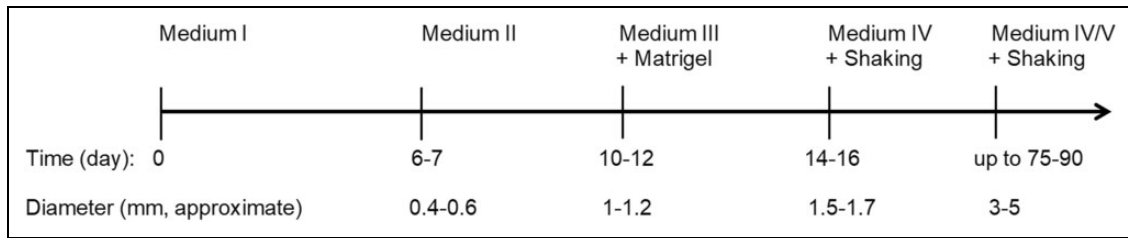


Fig 1. Overview of the cerebral organoid development procedure. Human ESCs were induced into cerebral organoids using the indicated multi-step procedure, where different media and treatments were introduced at each stage of the protocol. Timeline and approximate size measurements at each stage are also indicated.

Generation and Microscopy of Cerebral Organoids

Generation of the human cerebral organoids was performed as summarized in Fig 1. Briefly, human cerebral organoids were generated from human embryonic stem cells (ESCs), (H1 or WA-01 cell line) in a stepwise procedure that encompasses five stages: (1) induction of embryoid bodies by seeding the ESCs dispersed in Medium I (contains Knockout serum replacement and basic fibroblast growth factor (bFGF)) on a low-attachment surface for 6–7 days; (2) induction of neuroectoderm for 4–5 days in the presence of Medium II (contains N2-supplement and heparin); (3) embedding the developing organoid in a drop of matrigel and growing it in a stationary culture containing Medium III (contains Neurobasal Medium and B27-supplement) for 4–5 days; (4) transferring the developing organoids to shaking cultures containing Medium IV (contains Neurobasal Medium, B27-supplement and retinoic acid); and (5) if desired, additional “maturing factors” (e.g. ascorbic acid, cyclic adenosine monophosphate (cAMP), brain-derived neurotrophic factor (BDNF), glial cell line-derived neurotrophic factor (GDNF), and transforming growth factor- β (TGF β)) may be added to the organoids from day 60 post-induction of the embryoid bodies. Neuron-rich organoids can be observed within 40–50 days from the initial induction of embryoid bodies. Optimal organoids at the age of \sim 75 days reach a diameter of \sim 3–5 mm. Mature neurons could be reached by day 75–90 from the initial induction. Detailed composition of the various reagents was described previously³³. Organoids analyzed in this study were used on day 80 post-induction. Three independent batches of organoids were analyzed; in each batch, 10–20 organoids were analyzed. From each organoid \sim 10 sections were arbitrarily selected for staining and confocal microscopy.

To prepare the organoids for microscopy and synapse imaging, place the organoid in a sterile Petri dish. With the help of a surgical microscope, use a blunt tool such as blunt-end forceps or a 1000 μ l pipette tip to carefully unwrap the organoid from the matrigel coat. Discard the matrigel wrap. Work carefully and quickly to avoid damaging the organoid structure. Immerse the matrigel-freed organoid in OCT CompoundTM in Tissue Path Disposable Base Molds. Make sure that there is a layer of OCT CompoundTM below and

above the organoid (i.e. the organoid should be completely surrounded by OCT CompoundTM from all directions). If the organoid sinks down to the bottom of the mold, center it to the middle of the OCT CompoundTM using a blunt pipette tip gently. Snap-freeze the OCT CompoundTM-immersed organoid using liquid nitrogen (preferably) or ethanol-dry ice mix. Liquid nitrogen snap-frozen organoids can be stored for long periods of time (6–12 months, or longer) at -80°C .

Before sectioning the organoids, transfer the frozen organoids from -80°C to -20°C for at least 2 h or preferably overnight. We observed that gradually cooling and stabilizing the organoid at -20°C (which is the temperature for cryosectioning) gives better-quality sections that adhere to the slide better. Using a suitable cryostat at approximately -20°C (sometimes we found that -16°C to -17°C temperature may produce better quality sections than slightly lower temperatures), cut organoid sections at the desirable thickness. We found that a section thickness between 8 μm and 15 μm is ideal for on-slide staining and confocal microscopy. Collect the cryostat-cut sections on SuperfrostPlusTM glass slides. The tissue should adhere to the slide; if not, finely adjust the cryostat temperature as noted above, which we found helped reduce this problem. Air-dry the sections at room temperature for \sim 15–20 min. Using a barrier pen, draw a line (hydrophobic barrier) around each section, to help save expensive reagents (such as antibodies) and reduce expenses.

Fix the sections on the slide using a suitable fixative. Testing various fixatives and fixation durations, we found that cold 4% PFA for 8–12 min is optimal. Remove the fixative solution off the section using a pipette, and add \sim 100 μ l PBS on the tissue. Perform this wash process twice, for 5 min each time, at room temperature. Block non-specific binding for 1 h at room temperature, by adding \sim 100 μ l (enough volume to cover the tissue) of Blocking Solution to the section. Incubate the sections with primary antibody (diluted in AVS) overnight at 4°C on a rotating shaker. Optimizing antibody dilution and exposure duration may be performed for the different antibodies. Wash the primary antibody off the sections, with PBS, two to three times, each for 5–20 min (the number of washes and the wash duration may be optimized for each antibody). Incubate the sections with fluorophore-conjugated secondary antibody diluted in AVS for 45–60 min at room temperature in the dark.

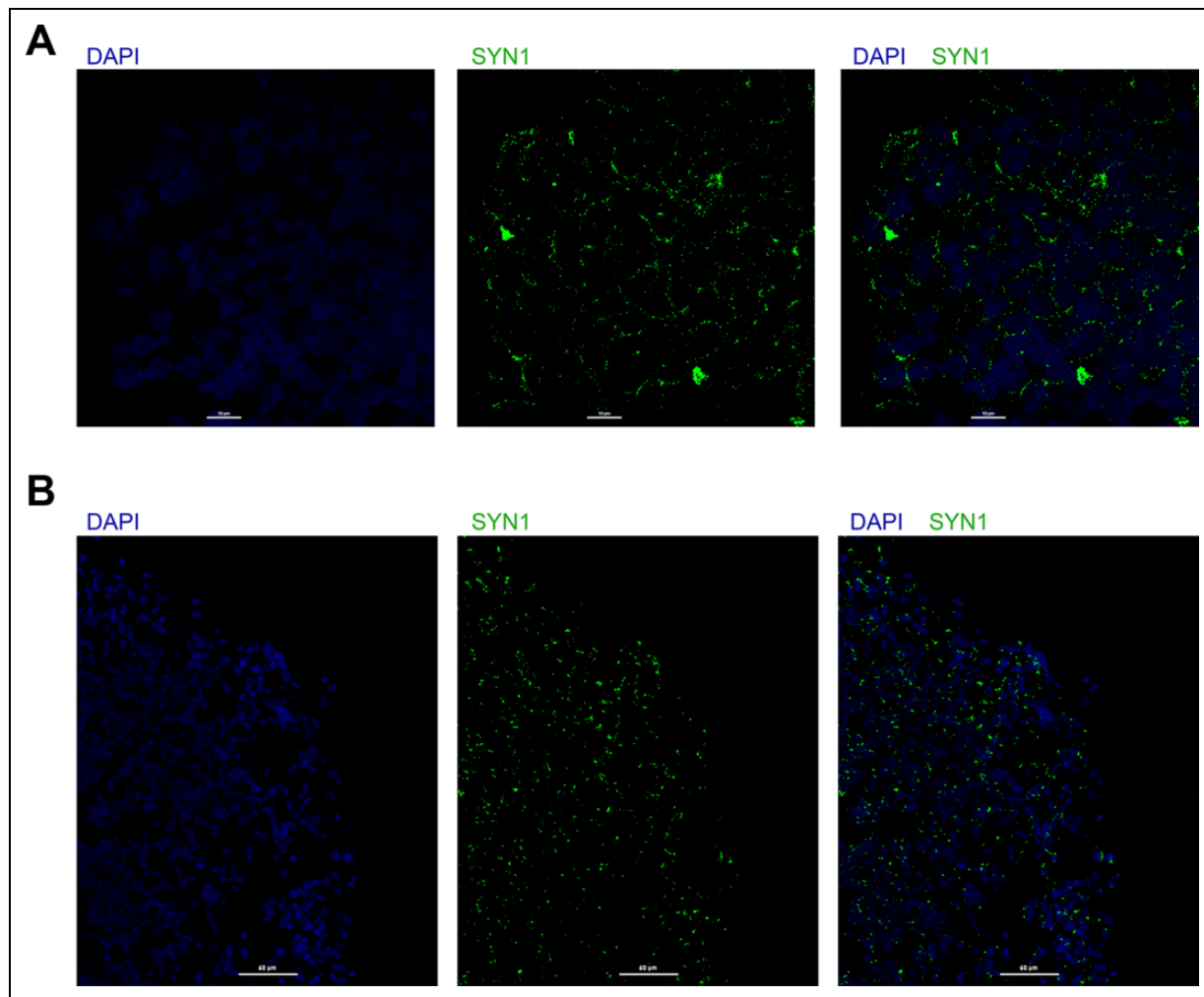


Fig 2. Synapse analysis using single staining for SYN1 or GluNR1. A. Organoid sections were stained for SYN1 (green) and nuclei (DAPI, blue), and imaged using confocal microscopy; scale bar 10 μm . B. Organoid sections were stained for GluNR1 (green) and nuclei (DAPI, blue) and imaged using confocal microscopy; scale bar 60 μm .

Remove the secondary antibody off the sections and wash with PBS for three to four times, each for 15 min (numbers and duration of washes may be optimized depending on the secondary antibody to be used). Carefully remove any remaining moisture on the slide using Kimwipes without touching the sections. Mount the sections in DAPI-containing mounting medium, and cover the slide with the coverslip being cautious to not introduce air bubbles under the coverslip. Proceed to imaging using a suitable confocal microscope with a 100x (or at least 63x) magnification lens for best visualization of synapses, as performed here. Slides can be stored at room temperature in light-proof containers.

Results

Synapse Analysis Using Single Staining for SYN1 or GluNR1

Organoid sections can be stained for SYN1 or GluNR1, via the procedure described above and using primary antibodies against SYN1 or GluNR1, and secondary antibodies

conjugated with a suitable fluorophore, such as Alexa Fluor-488. Punctate SYN1 or GluNR1 staining denotes synapses, as shown in Fig 2.

Synapse Analysis Using SYN1 and MAP2 Double Staining

Organoid sections can be stained for MAP2 (stains neuronal dendrites) and SYN1 (stains synapses) via the procedure above, using primary antibodies against SYN1 and MAP2, and secondary antibodies conjugated with Alexa Fluor-488 (SYN1, green) and -546 (MAP2, red). Synapses can be observed as SYN1+ punctae that colocalize with MAP2 (Fig 3).

Synapse Analysis Using SYN1 and PSD95 Double Staining

SYN1 is a presynaptic adaptor protein, while PSD95 is a postsynaptic adaptor present on the excitatory postsynapses. Thus, synapses appear as punctae that are double-positive for both

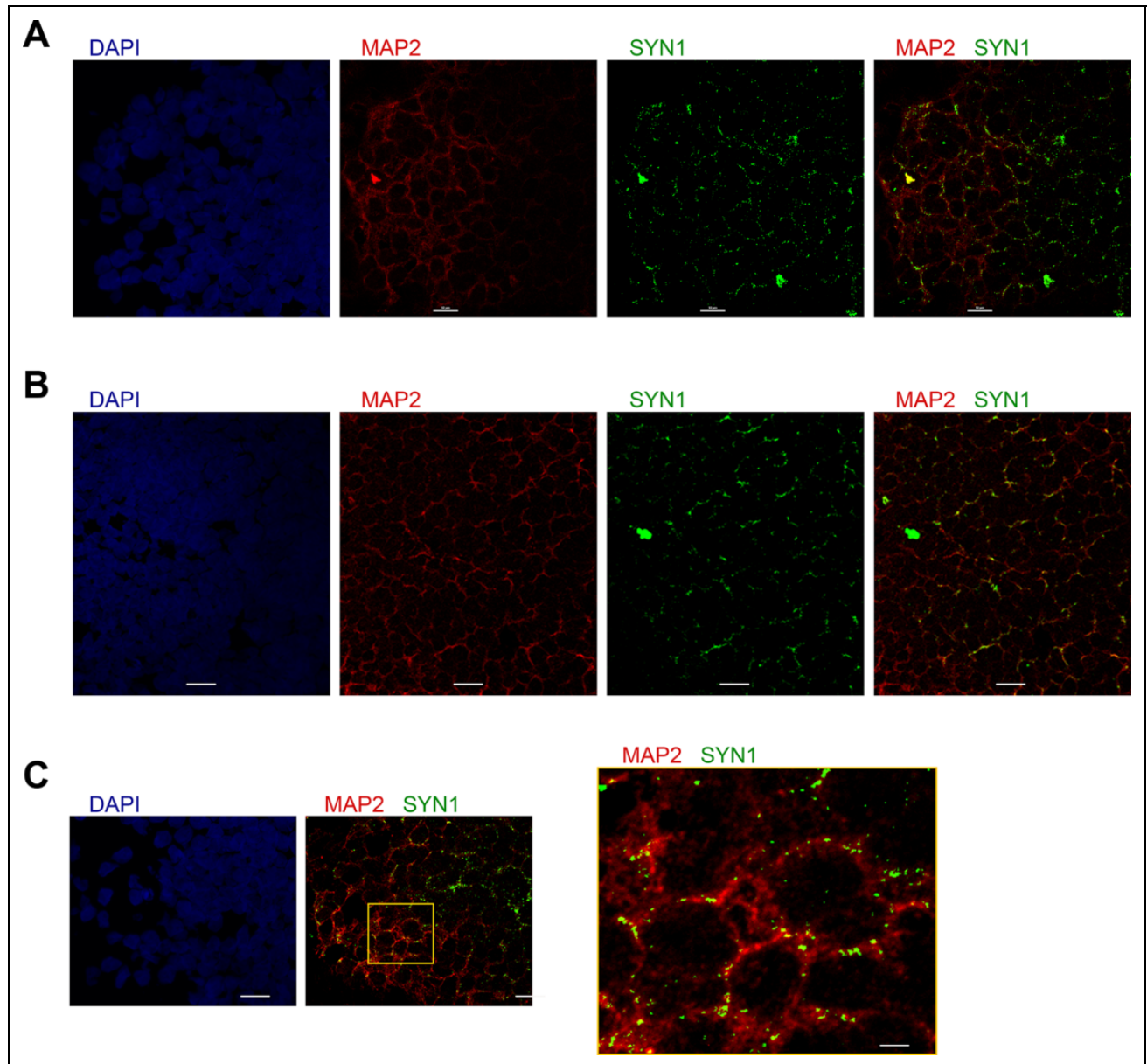


Fig 3. Synapse analysis using double staining for SYN1 and MAP2. A, B. Various organoid sections stained for SYN1 (green), MAP2 (red), and nuclei (DAPI, blue); scale bar 10 μm . C. An organoid section stained for SYN1 and MAP2 (scale bar 10 μm); right panel shows a higher magnification of the box-enclosed area in the middle panel (scale bar 2 μm).

SYN1 and PSD95. The organoid sections can be stained for SYN1 and PSD95 via the procedure above, using primary antibodies against SYN1 and PSD95, and Alexa Fluor-488- or -546-conjugated secondary antibodies. Fig 4A shows synapses in organoid sections that stain for both SYN1 and PSD95. Analysis of synaptic punctae should yield a Fluorescence-Relative distance curve, similar to that shown in Fig 4B, where the puncta shows that the presynapse and the postsynapse are spatially proximal yet separated, with SYN1 occupying the presynaptic localization and PSD95 occupying the postsynaptic localization. We analyzed a large number of synapses (~ 100 synapses) throughout the section, and show a representative example in Fig 4B. This analysis can be performed using automated

microscopy software (NIS-Elements; Nikon), which measures the distance of each fluorophore in a selected puncta relative to a reference point. The analysis confirms the spatial separation of the pre- and post-synaptic markers in the puncta; a bona-fide synapse should show that the presynaptic and the postsynaptic markers are spatially separated (relative distance > 0) and not superimposable (relative distance $= 0$).

Synapse Analysis Using SYN1 and GluNR1 Double Staining

While SYN1 is a presynaptic marker, GluNR1 is a postsynaptic glutamate (NMDA) receptor on excitatory synapses.

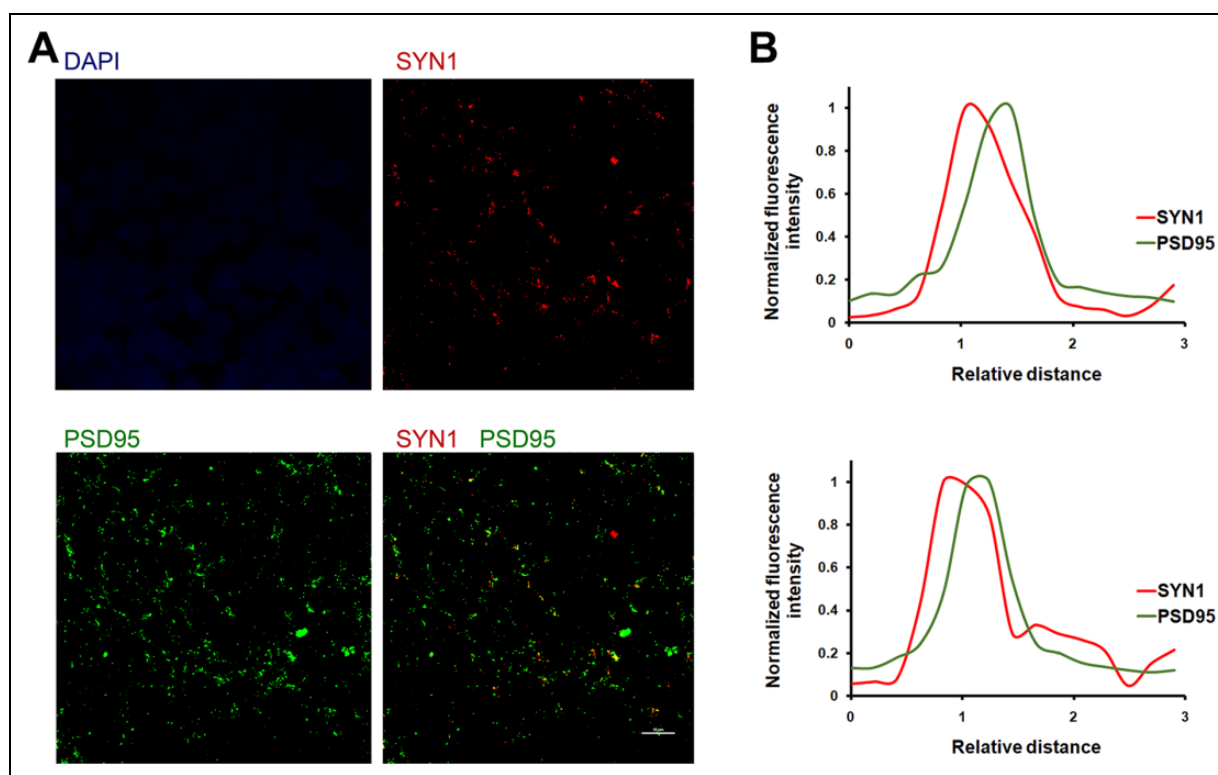


Fig 4. Synapse analysis using double staining for SYN1 and PSD95. A. An organoid section was stained for SYN1 (red), PSD95 (green), and nuclei (DAPI, blue); scale bar 10 μm . B. Fluorescence–Relative distance analysis for two random Syn1+/PSD95+ punctae in (A), showing the spatial separation of SYN1 and PSD95 peaks, denoting their presynaptic and postsynaptic localizations, respectively. Puncta analysis was performed using Nikon's NIS-Elements™ software.

Thus, synapses appear as punctae double positive for both SYN1 and GluNR1. Organoid sections can be stained as usual, using primary antibodies against SYN1 and GluNR1, and Alexa Fluor-488 or -546-conjugated secondary antibodies. Synapses can be observed as SYN1+/GluNR1+ punctae as shown in Fig 5A.

Synapse Analysis Using SYN1, VGLUT1 and GluNR1 Triple Staining

Synapses could be visualized using triple staining for SYN1, GluNR1 along with VGLUT1, a glutamate transporter present on the presynapse. Follow the staining Protocol above, using primary antibodies against SYN1, VGLUT1 and GluNR1, and Alexa Fluor-488, -546, and -633-conjugated secondary antibodies. Thus, synapses appear as punctae that are double and triple positive for SYN1, VGLUT1, and GluNR1, as shown in Fig 5B. Analysis of any synaptic puncta should yield a Fluorescence–Relative distance curve, similar to that shown in Fig 5C, where the punctae show that the presynapse and the postsynapse are spatially proximal but separated, with SYN1 and VGLUT1 localized presynaptically and GluNR1 localized postsynaptically.

Discussion

Synaptic function mediates circuitry development in the brain and regulates animal behavior. Deficits in circuitry development underlie devastating neurodevelopmental conditions, such as ASD, intellectual, communication and learning disorders, attention-deficit/hyperactivity disorder (ADHD), and other conditions. Harnessing the power of the organoid model to understand the mechanisms of these conditions requires the ability to analyze synapses in such a promising model. Here, we have described simple, immunofluorescence-based methods that we have tested and optimized to visualize and analyze synapses in sections of human cerebral organoids.

Synapses' proteomic composition is very complex, comprising hundreds of proteins and protein isoforms^{34,35}, and varies significantly between synapse subtypes^{35–37}, and thus synapse analysis is not a simple task. Previous studies on the mouse cortex have attempted to label synapses using various presynaptic proteins that are common to all synapse subtypes as synaptic markers, such as synaptophysin, bassoon, and SYN1, but only SYN1, a phosphoprotein that is largely concentrated at the presynaptic boutons^{38,39}, showed the most robust labeling, characterized by clearly visible punctae with very little nonspecific (non-synaptic) labeling⁴⁰. Quite

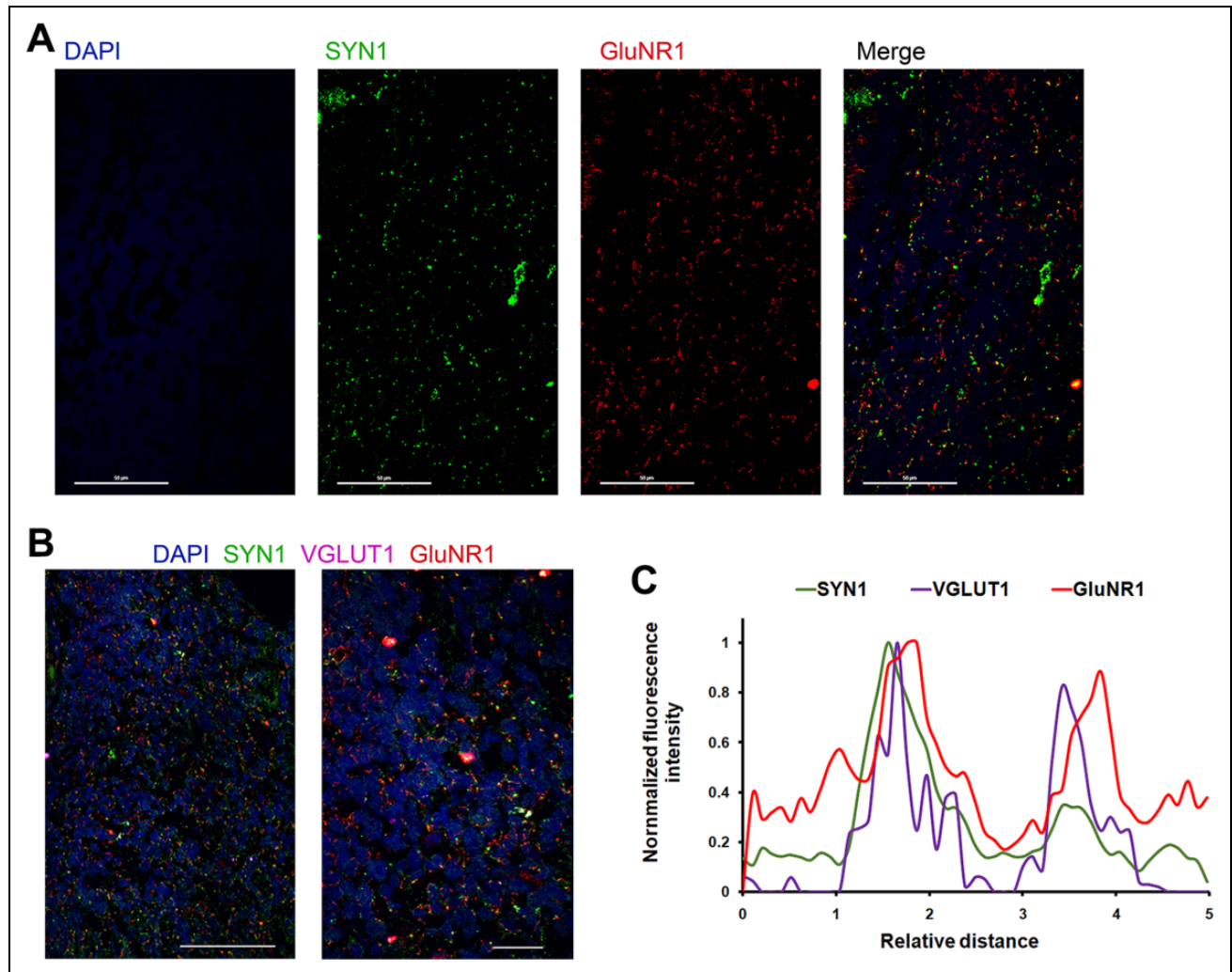


Fig 5. Synapse analysis using double and triple staining for SYN1, GluNR1, and VGLUT1. **A.** An organoid section stained for SYN1 (green), GluNR1 (red), and nuclei (DAPI, blue); scale bar 50 μm . **B.** An organoid section stained for SYN1 (green), GluNR1 (red), VGLUT1 (purple), and nuclei (DAPI, blue); scale bar 50 μm (left panel) or 10 μm (right panel). **C.** Fluorescence–Relative distance curves for two arbitrarily chosen SYN1+/VGLUT1+/GluNR1+ punctae in (B), showing spatial separation of SYN1, or VGLUT1, and GluNR1, and indicating the presynaptic localization of SYN1 or VGLUT1 and the postsynaptic localization of GluNR1. Analysis of the punctae was performed using Nikon’s NIS-ElementsTM software.

intriguingly, comparing the results of electron microscopy (EM) and immunofluorescence analyses in parallel has shown that the vast majority of synapses confirmed by EM stained positively for SYN1, further stressing the reliability of SYN1 immunofluorescence labeling for assaying synapses⁴⁰. Learning from all those studies, we have devised here the confocal microscopical synapse analyses described, using SYN1 staining. SYN1 staining yields a non-cell-body punctate staining pattern characteristic of synapses^{24,40}.

In addition to SYN1 on the presynapse, glutamate receptors decorate the excitatory postsynapses^{34–36}. Glutamate NMDA receptors are of cardinal importance for mediating Ca^{2+} -dependent postsynaptic processes associated with long-term potentiation and depression and learning and memory^{41,42}. Although presynaptic or extrasynaptic

presence of NMDA receptors was suggested by some reports⁴³, presynaptic or extrasynaptic expression of NMDA receptors was shown to be confined to a small subset of neurons in the mouse brain and mostly for the 2B subunit of the receptor (GluN2B)^{44–46}. The overwhelming majority, however, of NMDA receptors are expressed at the excitatory postsynapses and have been used in the field as a reliable postsynaptic marker in mouse brain sections or neuron culture^{47–59}. While multiple NMDA receptor subtypes exist, GluNR1 is the most important subunit as it is required for activity and is expressed mostly postsynaptically⁶⁰, which subunit we have used in our assays here. GluNR1 staining in organoid sections yielded a punctate staining pattern as expected for synapses. Moreover, the spatial analysis of the SYN1/VGLUT1/GluNR1 synaptic punctae in Fig 5C clearly

shows GluNR1's postsynaptic localization and distance from the presynaptic proteins (SYN1 or VGLUT1); notice that SYN1 and VGLUT1 peaks are spatially superimposable or very close to one another, whereas the GluNR1 peak is significantly farther (denoting postsynaptic localization).

Furthermore, to add more confidence to SYN1 or GluNR1 single-staining-based immunofluorescence visualization of synapses, we have devised double and triple labeling involving SYN1 along with another presynaptic marker such as VGLUT1, or postsynaptic markers such as PSD95 and GluNR1. Colocalization of the presynaptic marker with the postsynaptic marker and the Fluorescence-Relative distance curves of those synapses collectively confirm the synaptic nature of the immunopositive punctae visualized microscopically.

Traditionally, besides immunofluorescence, synapses have been analyzed using EM. EM is probably the most accurate assay of visualizing synapses; however, EM approaches may have some limitations, such as being a low-throughput method, limiting the possibility of using it to compare a large number of experimental conditions^{61,62}. We believe that the immunofluorescence- and confocal microscopy-based approaches described here will complement other approaches to visualize synapses in organoids representing multiple genetic or pharmacological conditions, and the conditions that seem promising via this immunofluorescence microscopical approach could be further validated and analyzed for synaptic connectivity and neurotransmitter release via electrophysiological approaches^{31,32}. Overall, the work described here will enhance the utility of the human cerebral organoid model in studies of circuitry and molecular neuroscience and to investigate synaptic deficits that account for certain neurodevelopmental disorders.

Ethical Approval

Ethical approval is not applicable to this article.

Statement of Human and Animal Rights

This article does not contain any studies with human or animal subjects.

Statement of Informed Consent

This article involves no human subjects; and informed consent is not applicable.

Declaration of Conflicting Interests

The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

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