Human Neural Organoids for Studying Brain Cancer and Neurodegenerative Diseases

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Abstract

The lack of relevant in vitro neural models is an important obstacle on medical progress for neuropathologies. Establishment of relevant cellular models is crucial both to better understand the pathological mechanisms of these diseases and identify new therapeutic targets and strategies. To be pertinent, an in vitro model must reproduce the pathological features of a human disease. However, in the context of neurodegenerative disease, a relevant in vitro model should provide neural cell replacement as a valuable therapeutic opportunity.

Such a model would not only allow screening of therapeutic molecules but also can be used to optimize neural protocol differentiation [for example, in the context of transplantation in Parkinson's disease (PD)]. This study describes two in vitro protocols of 1) human glioblastoma development within a human neural organoid (NO) and 2) neuron dopaminergic (DA) differentiation generating a three-dimensional (3D) organoid. For this purpose, a well-standardized protocol was established that allows the production of size-calibrated neurospheres derived from human embryonic stem cell (hESC) differentiation. The first model can be used to reveal molecular and cellular events occurring during in glioblastoma development within the neural organoid, while the DA organoid not only represents a suitable source of DA neurons for cell therapy in Parkinson's disease but also can be used for drug testing.

Video Link

The video component of this article can be found at https://www.jove.com/video/59682/

Introduction

The World Health Organization (WHO) classifies astrocytomas as low grade (grade I to II) or high grade (grade III and IV). Glioblastoma multiforme (GBM) is an astrocytoma grade IV, the most lethal of primary brain tumors, that is resistant to all current forms of treatments1. Despite standard-of-care therapy including neurosurgery, chemotherapy, and radiotherapy, GBM remains fatal and the 15-month overall survival rate has not dramatically changed over the past 15 years2. To make significant progress in understanding GBM pathogenesis, the use of relevant models is key. So far, the study of GBM has relied on cell lines, rodent organotypic slices, and xenotransplantation of patient-derived cells into mice or transgenic mice developing spontaneous tumors3,4. Although these models have been useful to study brain metastasis and tumor aggressiveness, they are restricted by differences among species, and resulting conclusions may be incorrectly translated to human tissues. Moreover, existing models with human cells are also limited by the absence of host tissue/tumor interactions5,6. Experimental models are critical for the translation from basic science to therapeutic targets. Therefore, describing a protocol to produce in vitro human neural organoids co-cultured with GBM-initiating cells (GICs) can provide a relevant system that mimics morphological and functional features of GBM development. This system reproduces some in vivo features of GBM developmentsuch as diffuse migration of invading cells and necrosis areas, and it highlights gene expression relevant to tumor biology. As previously revealed, some critical microRNAs are induced during GIC development within 3D nervous tissue5,6.

PD is a major neurodegenerative disorder and associated with the degeneration of multiple neuronal subtypes7. Even if a progressive onset of symptoms (e.g., bradykininesia, asymmetric rest tremor, rigidity and posture instability) characterizes the disease, its exact etiology is not clearly established. Indeed, many studies have highlighted evidence that major risk factors can result from a combination of genetic and environmental factors. Parkinsonian symptoms are associated with the bilateral degeneration of dopaminergic neurons in the substantia nigra (SN), leading
This study describes two protocols using human neural organoids, with one specifically oriented towards a midbrain-like phenotype enriched with TH-positive cells.

**Protocol**

This protocol follows the guidelines of University of Geneva’s human research ethics committee.

**1. Maintenance and culture of undifferentiated human embryonic stem cells (hESCs)**

1. Perform maintenance and expansion of hESCs on feeder-free conditions by pre-coating dishes with a specific extracellular matrix.
   1. Thaw 300 µL of extracellular matrix at 4 °C (typical range concentration 18-22 mg/mL, keep on ice) and gently mix with 15 mL of cold DMEM medium to avoid a premature gelation of the extracellular matrix. Add 7.5 mL of the extracellular matrix to both T150 flasks.
   2. Incubate the dishes coated with extracellular matrix at 37 °C for at least 1 h (maximum overnight).
   3. Remove the medium and seed hESCs to a density of 6.5 x 10^5 cell/cm².

2. Maintain H1 (hESC cell line) in hESC medium and 1% penicillin/streptomycin.

3. Pass the cells with enzymatic procedure: add 7.5 mL of enzymatic solution to a T75 cm² flask and incubate at 37 °C for 1-2 min.

4. From days 11-13, culture the spheres in B27 medium supplemented with 0.5 µM BMP inhibitor.

5. From days 13-21, culture the spheres in B27 medium supplemented with 10 ng/mL glial derived neurotrophic factor (GDNF), 10 ng/mL brain derived neurotrophic factor (BDNF) and 1 µM of γ-secretase inhibitor. GDNF and BDNF promote neuronal and glial differentiation.

6. Change the medium every 2-3 days (usually on Monday, Wednesday and Friday), for a following 3 weeks of differentiation.

**2. hESC-derived neural organoids for GBM studies**

1. 24 h before starting the 3D culture, replace the hESC medium with a serum-free medium supplemented with 10 µM ROCK inhibitor (both components are necessary to support cell survival and spontaneous neurosphere formation during the aggregation phase in a microwell plate). The cells should be at 60% confluency. The next day (day 0), detach hESC colonies as single cells: remove the medium, then rinse with PBS without Ca²⁺/Mg²⁺, add 5 mL of enzymatic dissolution solution, and incubate at 37 °C for 1-2 min.

2. Collect the cells in serum-free medium with 10 µM of ROCK inhibitor and centrifuge the cells at 300 x g for 5 min. To allow for better survival, re-plate cells at the desired density onto extracellular matrix-coated dishes, in the same medium containing Rock-associated protein kinase (ROCK) inhibitor (10 µM) for 24 h.

3. Pass the cells in serum-free medium with 10 µM of ROCK inhibitor and centrifuge the cells at 300 x g for 5 min. Remove the supernatant and count the cells in 10 mL of serum-free medium supplemented with 10 µM of ROCK inhibitor.

4. Prepare 28.2 x 10^6 cells in 12.5 mL of serum-free medium supplemented with 10 µM ROCK inhibitor. Dispense 1000 cells/microwell. Centrifuge the cells at 300 x g for 5 min and place the plate in the incubator at 37 °C overnight (maximum 36 h). For example: to obtain 30 human neural organoids, use one T150 flask at 70%-80% of confluence (about 30 million cells).

5. The next day (day 1), collect the spheres (with a P1000) and place them in a 6 well plate. In each well, add 2 mL of B27 medium and DMEM-F12 GlutaMAX and Neurobasal medium (mix at 1:1), supplemented with 1% B27 supplements and 1% non-essential amino acids (NEAA). To promote fast neural induction, supplement the medium with dual-SMAD inhibition cocktail, composed of 10 µM TGFβ/Act/Inhibitor and 0.5 µM bone morphogenetic protein (BMP) inhibitor. From this step forward, the spheres are cultured in rotation (60 rpm, orbital shaker).

6. Change the medium every 2-3 days: bend the plate and let the spheres fall down for 5 min, remove half of the medium (2 mL), and add 2 mL of fresh B27 medium supplemented with growth factors and inhibitors. Do not centrifuge the spheres.

7. Perform neural induction according to the following time course:
   1. From days 1-4, culture the spheres in B27 medium supplemented with dual-SMAD. The dual-SMAD inhibition cocktail (10 µM TGFβ/Act/Inhibitor and 0.5 µM BMP inhibitor) promote the neural induction.
   2. From days 4-11, promote proliferation of hESC-derived neural rosettes (into the spheres), by adding 10 ng/mL epidermal growth factor (EGF) and 10 ng/mL basic fibroblast factor (bFGF) to the B27 medium supplemented with dual-SMAD cocktail.
      NOTE: On day 11, most of the cells should be positive for Nestin.
   3. From days 11-13, culture the spheres in B27 medium supplemented with 0.5 µM BMP inhibitor.
   4. From days 13-21, culture the spheres in B27 medium supplemented with 10 ng/mL glial derived neurotrophic factor (GDNF), 10 ng/mL brain derived neurotrophic factor (BDNF) and 1 µM of γ-secretase inhibitor. GDNF and BDNF promote neuronal and glial differentiation. The γ-secretase inhibitor allows for greater neural maturation.
   5. On day 21, plate the spheres (about 1,000 spheres) on a hydrophilic polytetrafluoroethylene (PTFE) membrane (6 mm diameter, 0.4 µm) deposited on a culture plate insert designed for 6 well plate. Stop any rotation from this step. The presence of rosettes, observed with a bright-field microscope, indicate the initiation of neural differentiation. The neural rosettes can be observed 2-3 days after plating spheres on the PTFE membrane.
   6. Add 1 mL of B27 medium supplemented with growth factors and inhibitors (as followed) to each well underneath the membrane insert, every 2-3 days (usually on Monday, Wednesday and Friday), for a following 3 weeks of differentiation.
   7. From days 21-25, cultivate human neural organoids in the same neural maturation medium (Cf. step 2.7.4).
   8. From days 25-28, only complement B27 medium with 1 µM γ-secretase inhibitor.
   9. From days 28-39, stop adding the γ-secretase inhibitor and continue human neural organoid culture in B27 medium only.
3. Isolation and cultivation of glioblastoma-initiating cells (GICs)

1. Isolate GICs by fragmenting a high grade human GBM biopsy. Transfer the piece of tumor in a beaker containing 0.25% trypsin in 0.1 mM EDTA (4:1) and slowly stir at 37 °C for 30-60 min (depending on tumor size).

2. Plate the dissociated cells in 75 cm² tissue culture flasks plated at 2,500-5,000 cells per cm² in GIC medium: DMEM/F-12 medium (1:1) containing 1% N2, 1% B27, and 1% G5 supplements (to favor GIC survival), supplemented with bFGF and EGF (both at 10 ng/mL, to promote stemness) and 1% of penicillin/streptomycin.

3. Once the GIC is well-established and growing, remove the N2 and G5 supplements from the GIC medium.

4. One day before adding the cells onto the organoid, dissociate the GICs and count them.

5. Rinse the microwell plate with 2 mL of GIC medium and centrifuge the plate at the maximum speed to remove bubbles (1000 x g). Place the GICs at 1,000 cells to obtain one gliomasphere per microwell. Incubate overnight at 37 °C (Figure 1C). This step is key and allows for well-calibrated GICs (an example of necrotic and over-sized GICs is shown in Figure 2A,C).

6. To initiate GBM invasion, add one gliomasphere on top of the neural tissue with a large bore pipet tip (Figure 1F). Carefully place the 6 well plate back in the incubator.

4. hESC-derived dopaminergic organoids for PD studies

1. Day 0: Amplify hESCs in 2D culture up to 60% confluency (day 0), then replace stem cell media used to maintain pluripotency features of hESCs with a serum-free medium. Start neural induction by supplementing culture medium with 0.5 µM BMP inhibitor and 10 µM TGFβ/Activin/Nodal inhibitor (dual-SMAD inhibition cocktail), then add 10 µM ROCK inhibitor for 24 h to increase the survival rate of cells during passage.

2. Day 1: Prepare the microwell plate with 2.5 mL per well of serum-free medium supplemented with 0.5 µM BMP inhibitor, 10 µM TGFβ/Activin/Nodal inhibitor, and 10 µM ROCK inhibitor. To specify cells towards the ventral pattern of the neural tube, add 100 ng/mL Sonic Hedgehog (SHH), 100 ng/mL fibroblast growth factor 8 (FGF8), and 2 µM smoothened agonist. Centrifuge the plate (only with medium and without cells) at 1200 x g for 5 min to remove air bubbles from the microwells.

3. Adjust the medium volume to obtain a cell suspension allowing to form neurospheres containing 1000 cells per microwell (for example, the microwell plate used here contains 4,700 microwells per well). So, prepare 4.7 million cells in 2.5 mL of medium and add it to the previous 2.5 mL of medium already placed in the plate.

4. In order to correctly distribute the cells in each microwell, gently shake the plate, and centrifuge the microwell plate 300 x g for 5 min. Incubate the plate at 37 °C for 24 h to generate spheres.

3. Day 2: Gently flush the microwells with the medium and collect then transfer the spheres in tissue-treated six-well plate. Replace medium with Neurobasal medium supplemented with 1% B27, 1x NEAA, 2 mM L-glutamine, and 1% of penicillin/ streptomycin. Add 0.5 mM dibutyryl cAMP (to favor maturation), 20 nM inhibitor of histone deacetylase (for cell cycle exit), 1 µM γ-secretase inhibitor and growth inhibitor, which activates the Wnt/β-catenin pathway. Maintain GSK-3β inhibitor in the medium up to day 13. Split into two new tissue-treated 6 well plates to reduce both sphere density per well and avoid spheres aggregation.

1. Place spheres in rotation at 37 °C (60 rpm, orbital shaker) and change half-medium freshly supplemented every 2-3 days.

4. Day 3: To enhance neural induction and convert to neural progenitors with a midbrain identity, supplement the medium with 3 µM GSK-3β inhibitor, which activates the Wnt/β-catenin pathway. Maintain GSK-3β inhibitor in the medium up to day 13. Split into two new tissue-treated 6 well plates to reduce both sphere density per well and avoid spheres aggregation.

NOTE: At Day 8, most of the cells should be positive for Nestin.

5. Day 8: Start the neural maturation: replace neuralization factors SHH, FGF8, smoothened agonist, and dual-SMAD inhibition small molecules with Neurobasal medium supplemented with 1% B27, 1x NEAA, 2 mM L-glutamine, and 1% of penicillin/ streptomycin. Additionally, add regionalization factors SHH, FGF8, smoothened agonist, and dual-SMAD inhibition small molecules.

1. Place spheres in rotation at 37 °C (60 rpm, orbital shaker) and change half-medium freshly supplemented every 2-3 days.

5. Quantification of TH and Nurr1 gene expression for validation of dopaminergic differentiation

1. RNA extraction: On the indicated day of differentiation, lyse 40 neurospheres with 350 µL of RLT buffer (provided in RNA extraction kit) supplemented with 3.5 µL of 2-mercaptoethanol. Extract the RNA from the lysed neurospheres using an RNA extraction kit following the manufacturer's instructions.
2. Quantify total RNA concentrations.
3. Perform reverse transcription of 300 ng of the total RNA extraction using reverse transcription kit for quantitative real-time polymerase chain reaction (qPCR) and follow the manufacturer's instructions.
4. Perform qPCR analysis on real-time PCR detection systems, based on asymmetrical cyanine dye detection. Normalize the data with housekeeping genes: glyceraldehyde-3-phosphate dehydrogenase (GAPDH) and elongation factor 1-alpha (EF1). Sequences of primers are described in Table 1.

6. High pressure liquid chromatography (HPLC) detection

1. Use high pressure liquid chromatography (HPLC) with electrochemical detection to detect the presence of dopamine. Dopamine was extracted by lysing neural organoids in 100 mL of 0.1 N perchloric acid (HClO4) for 15 min at 4 °C with a vigorous vortexing every 5 min. After centrifugation, collect and store the supernatant at -20 °C for dopamine dosage.
2. Use a C-18 column (5 µm, 4.6 mm x 150 mm) to separate the analytes by reversed-phase HPLC in isocratic mode at a flow rate of 1 mL/minute. Detection of dopamine should be carried out using a coulometric detector with the conditioning cell set at a potential of +200 mV.

7. Raw data recording with microelectrode array (MEA) platform

1. Use a dissection microscope to transfer neurospheres to the center of a porous MEA device.
2. Use an amplifier and data acquisition system for electrophysiological recordings. Measure the signal-to-noise ratio (SNR) as the standard deviation of the voltage during a 5 min recording, using the signal as the average peak-to-peak voltage of the spikes recorded in the same 5 min periods.

Representative Results

The critical steps of this protocol must be well-identified and handled properly. Therefore, a diagram of culture conditions indicating the time-lapse for each step as well as the compounds used for the differentiation protocol are illustrated in Figure 1A and Figure 3A for NO plus GBM and DA neural organoids, respectively. Figure 1B,C,D,E,F illustrates the cells, spheres, and NO and show the typical morphology for each step. Figure 1G,H,I illustrates immunofluorescence staining with some neural markers.
Figure 1: Human neural organoid (NO) differentiation protocol. (A) Standardized protocol for the generation of NO derived from human embryonic stem cells (hESC). (B) hESCs are maintained on extracellular matrix in hESC medium. (C) Microwell plates were used to generate calibrated neurospheres. At 2 weeks, neurospheres were plated onto the insert containing a PTFE membrane (scale bar = 50 µm). (D) Macroscopic view of NO into the insert in one well of a 6 well plate. During the first days, rosettes were observed (black arrow) (E). (F) Macroscopic view of a NO plus GIC sphere on the top. (G-I) Immunofluorescence analysis of NO plus GIC sphere (EGFR-positive; scale bar = 50 µm) (G) and NO alone, which showed immune reactivity for the neuronal marker βIII-tubulin and slightly positive for nestin; however, synapsin 1 showed a weak signal (H,I) (scale bars = 100 µm and 50 µm, respectively). Please click here to view a larger version of this figure.
Figure 2: Illustration of necrotic spheres and immature NO. The neurospheres (A) and NO (B) can undergo necrosis when they are too numerous in the well or oversized (C) (scale bar = 10 µm). (D) One GIC infected with a tomato reporter help to track tumor cell invasion in NO, scale bar, 10 µm. Example of immature NO with neural tubes (E) and no neural tubes (F) (scale bar = 50 µm). Please click here to view a larger version of this figure.
Figure 3: Standardized protocol for generation of DA neural organoid and electrophysiological and morphological analysis. (A) Standardized protocol for the generation of DA neural organoids. (B) Immunofluorescence analysis of DA neural organoid; TH-immunoreactive cells co-expressing Nurr1, a midbrain specific marker (scale bar = 50 µm). Data are represented as mean ± SEM (n = 3). (C) Graphs represent kinetics of TH and Nurr1 gene expression evaluated by qRT-PCR. (D) Representative HPLC: dopamine peak (arrow) was detected by HPLC from DA neural organoid lysate. (E) Example of raw data recorded with MEA platform. Each spike is displayed by a vertical line (time stamps), whereas the remaining trace is noise. (F) Picture representing a neurosphere deposited on the MEA. (G) Superposition of typical spikes (blue and red curves) detected from the raw data. The black bold curve indicates the average of the corresponding red curves. (H) Raster plot showing the time stamps associated with each spike detected. The different colors highlight the different electrodes. Please click here to view a larger version of this figure.

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<tr>
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<td>GAPDH</td>
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<td>AGGGAGATTCCATGTTGGGT</td>
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Table 1: Primers used in this protocol.

Discussion

One of the most critical aspects of this protocol includes the maintenance of hESC pluripotency during cell culturing and close monitoring of the spheres and neural organoid morphology. hESCs are very sensitive, and every manipulation can lead to early uncontrolled differentiation.
as well as cell death. In order to increase experimental reproducibility and avoid the occurrence of abnormal karyotype events, it is advised to cryopreserve several batches of hESCs at the lowest passage after validation of their chromosome stability. Moreover, it is recommended to thaw a new vial for each experiment and check the behavior of the cells every day. If the spheres are less refractive with abnormal higher size, they will likely start to aggregate and die.

One improvement upon this system is either perfusion or implementing a vascularized system (by adding endothelial cells or within a 3D fluidic microchip)\textsuperscript{12,13}. However, controlling the thickness of the neural organoid (≤300 µm) allows efficient passive perfusion of oxygen and nutrients and prevents necrosis. Another improvement is the introduction of immune cells (microglia). With these limitations in mind, neural organoids plus a GIC system may be a relevant tool for several reasons. First, this system allows drug screening to monitor how a therapeutic compound may affect an organoid or tumor cell. Second, cell-to-cell interactions can be studied, and micro-environmental determinants underlying individual and collective invasions can be visualized and explored\textsuperscript{12,13}.

In the context of Parkinson's disease, a neural organoid enriched in DA neurons can represent a relevant and accurate 3D model to study disease development. In previous studies, Parkinson's patient-derived induced pluripotent stem cells differentiated towards DA neurons have been used to study the affected neuronal subtypes. Of note, some disease-related phenotypes such as the accumulation of α-synuclein and sensitivity to oxidative stress have been observed\textsuperscript{14,15}. Moreover, the neural organoid may be used as a tool to screen therapeutic molecules. However, specific and relevant readouts should be set up to evaluate DA neuron survival and functionality, such as dopamine production and electrophysiological activity. Altogether, this protocol provides two standardized and accurate stem cell-based approaches to generate neural organoids.

Disclosures

The authors declare no competing financial interests.

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