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Title: Automated Production of Human Induced Pluripotent Stem Cell-Derived Cortical and Dopaminergic Neurons with Integrated Live-Cell Monitoring

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# **Author Questionnaire**

- **1. Microscopy**: Does your protocol involve video microscopy, such as filming a complex dissection or microinjection technique? **N**
- 2. Software: Does the part of your protocol being filmed demonstrate software usage? Y
- **3. Filming location:** Will the filming need to take place in multiple locations (greater than walking distance)? **N**

Protocol Length

Number of Shots: 55

# Introduction

### 1. Introductory Interview Statements

### **REQUIRED:**

- 1.1. <u>Ashutosh Dhingra</u>: The overall goal of this study is to show how to implement automated procedures for culturing and differentiating –human iPS cells to neuronal lineages as well as their automated imaging [1].
  - 1.1.1. INTERVIEW: Named talent says the statement above in an interview-style shot, looking slightly off-camera

## **REQUIRED:**

- 1.2. <u>Ashutosh Dhingra</u>: Standardized automated procedures -allow low experimental variation while ensuring high phenotypic reproducibility. In addition, this system can be adapted for the development of new cell culture protocols [1].
  - 1.2.1. INTERVIEW: Named talent says the statement above in an interview-style shot, looking slightly off-camera

#### Introduction of Demonstrator on Camera

- 1.3. <u>Ashutosh Dhingra</u>: Demonstrating the procedures will be <u>Joachim Taeger</u> and <u>Elisangela Bressan</u>, post docs from our lab [1][2].
  - 1.3.1. INTERVIEW: Author saying the above
  - 1.3.2. The named demonstrator(s) looks up from workbench or desk or microscope and acknowledges the camera

## **Protocol**

- 2. New Culture Plate and Tip Loading and Decontamination
  - 2.1. In the graphic user interface of the automated system, click **Resource-Instrument** process view [1] and select the resource-instrument process [2].
    - 2.1.1. WIDE: Talent clicking button, with monitor visible in frame
    - 2.1.2. SCREEN: screensho 1: 00:00-00:02
  - Click Run Instrument process and run the RunHepaHood and Reloading processes [1-TXT].
    - 2.2.1. SCREEN: screensho\_1: 00:03-00:09 **TEXT: Monitor material placement to avoid transport errors**
  - 2.3. Select the resources to be loaded [1], open the door [2], and load the cell culture plates and disposable tips in the appropriate positions as indicated by the pop-up images [3].
    - 2.3.1. SCREEN: screenshot 3: 00:00-00:10
    - 2.3.2. Talent opening door
    - 2.3.3. Talent loading plate(s) and/or tips
    - 2.3.4. SCREEN: screenshot\_3: 00:11-00:23
  - 2.4. Then decontaminate the door with 70% ethanol before closing [1-TXT] and run the **Decontamination** process. The system will be sterilized by UV radiation for 30 minutes [2].
    - 2.4.1. Talent spraying and/or wiping door **TEXT: Decontaminate door after every** handling
    - 2.4.2. SCREEN: screensho 4: 00:00-00:08

## 3. Culture Medium and/or Dissociation Reagent Filling

- 3.1. To load the plates with culture medium or dissociation reagent, execute the **RunHepaHood** step [1] and open the door of the liquid handling station [2].
  - 3.1.1. WIDE: Talent executing step, with monitor visible in frame
  - 3.1.2. Talent opening door
- 3.2. After decontamination with 70% ethanol [1], place the reservoirs on the deck to the assigned position [2] and de-lid them [3].
  - 3.2.1. Talent spraying/wiping reservoirs with ethanol
  - 3.2.2. Talent placing reservoir(s) onto deck
  - 3.2.3. Talent removing lid(s)
- 3.3. Then run the **InitHepaHood** resource-instrument process to power down the HEPA hood [1].
  - 3.3.1. SCREEN: screenshot 2: 00:00-00:11

#### 4. Automated Method Execution

- 4.1. To execute an automated culture method, in the calendar view of the graphic user interface [1], click **Add process step** and select the cell line to be used in the experiment [2].
  - 4.1.1. WIDE: Talent selecting calendar view, with monitor visible in frame
  - 4.1.2. SCREEN: screensho\_5: 00:00-00:07
- 4.2. Use the wizard to select the project and mark the batch to be used. Click the right-facing arrowhead and navigate to the next page of the wizard to select the process step to be executed [1-TXT].
  - 4.2.1. SCREEN: screenshot\_5: 00:07-00:14 **TEXT: Batch should be empty or contain** culture or assay plates

- 4.3. In the last page of the wizard, schedule the experiment and set the appropriate **Parameter Details** variables necessary for running the method. Then click **OK** [1].
  - 4.3.1. SCREEN: screensho 5: 00:14-00:24
- 5. Human Induced Pluripotent Stem Cell (hiPSC) Seeding
  - 5.1. After loading a 50-milliliter tube with the human iPS (eye-P-S-) cell suspension [1], load coated culture plates for receiving cells on the shelf [2-TXT] and execute the Seeding of plates from tubes method [3].
    - 5.1.1. WIDE: Talent loading tube into system, with monitor visible in frame
    - 5.1.2. Talent loading plate(s) onto shelf **TEXT: Decontaminate and close doors after loading**
    - 5.1.3. SCREEN: screenshot 6: 00:12-00:20
  - 5.2. To count the cells in the brightfield imaging cytometer, automatically prepare a 384-well counting plate with cell suspension on the deck [1] and use the robotic arm to transfer the 384-well counting plate from the deck to the brightfield imaging cytometer [2-TXT].
    - 5.2.1. Author NOTE: automatic function so not shot.
    - 5.2.2. Arm transferring plate to cytometer **TEXT: cytometer automatically determines the cell number per milliliter**
  - 5.3. **[1 TXT]**.
  - 5.4. After counting, the system will return the plate to its original position [1] and transfer a coated culture or assay plate from the shelf to the pipetting deck [2].
    - 5.4.1. Arm returning plate *Videographer: Important step*
    - 5.4.2. Arm selecting/transferring plate to pipetting deck *Videographer: Important* step

- 5.5. The system will remove coating and add fresh medium [2-TXT] . The system will then seed the cells in the user-defined number and volume suitable for the plate [1].
  - 5.5.1. Cells being seeded
  - 5.5.2. The coating is removed and add fresh medium **TEXT: Remove coating before** cell seeding as necessary. NOTE: Please move this step before 5.5.1
- 5.6. After seeding, the plate will be moved to the on-deck shaker for 10 seconds at 500 revolutions per minute for cell distribution [1] before being transferred to the carbon dioxide incubator [2].
  - 5.6.1. Shot of plate on shaker
  - 5.6.2. Plate being placed into incubator

#### 6. Automated Confluence Assessment

- 6.1. To assess the automated confluence, execute the **Check Confluency** method [1] and select a batch that contains at least one culture plate and no assay plates [2].
  - 6.1.1. WIDE: Talent executing method, with monitor visible in frame
  - 6.1.2. SCREEN: screenshot 8: 00:06-00:12
- 6.2. In the **Parameter Details** section, enter iPSCf\_2020 for image acquisition and the imaging analysis settings [1].
  - 6.2.1. SCREEN: screenshot 8: 00:16-00:29 Video Editor: can speed up
- 6.3. The system will then transfer the first plate from the incubator to the brightfield imaging cytometer [1] and image the cell confluence [2-TXT].
  - 6.3.1. Plate being transferred with robotic arm *Videographer: Important step*
  - 6.3.2. Cells being imaged *Videographer: Important step* **TEXT: Transfer plate back to** incubator after imaging

## 7. Medium Change

- 7.1. To change cell culture or assay plate media, execute the **Media Change of Culture Plates** method [1] and select a batch containing only culture plates [2].
  - 7.1.1. WIDE: Talent executing method, with monitor visible in frame
  - 7.1.2. SCREEN: screenshot 9: 00:06-00:13
- 7.2. The system will transfer the plates to the deck [1] and tilt the plates to allow aspiration of the supernatant [2].
  - 7.2.1. Arm moving plate(s) to deck *Videographer: Important step* NOTE: included in shot 6.3.1 take 6).
  - 7.2.2. Tilting plate and/or supernatant being aspirated *Videographer: Important step*
- 7.3. The supernatant will be discarded into the waste collection module [1] and 12 milliliters of fresh medium will be added to each plate [2].
  - 7.3.1. Arm discarding supernatant into waste module and discarding used tips *Videographer: Important step*
  - 7.3.2. Arm adding medium to well(s) *Videographer: Important step*
- 7.4. Then system will then re-lid the plates [1] and return the plates to the cell culture incubator [2].
  - 7.4.1. Arm putting lid onto plate
  - 7.4.2. Arm placing plate into incubator
- 7.5. Alternatively, to change medium of assay plates, execute Media Change of Assay Plates method. [1]
  - 7.5.1. Media change of 96-well assay plates. (shot of 96-well assay plates on the deck with pipette tips). NOTE: Maybe added shot

#### 8. Subcultivation

- 8.1. To subcultivate the cells, execute the **Subcultivation of Adherent Cells** method [1] and select the batch containing the culture plates that need subcultivation [2].
  - 8.1.1. WIDE: Talent selecting method, with monitor visible in frame
  - 8.1.2. SCREEN: screenshot 10: 00:09-00:16
- 8.2. After discarding the medium, the system will wash the cells one time with 8 milliliters of PBS per plate [1] before adding 8 milliliters of 0.5-millimolar EDTA to the cells [2].
  - 8.2.1. Arm adding PBS to cells
  - 8.2.2. Arm adding EDTA to cells
- 8.3. After 8 minutes on the deck with tilting option [1], the EDTA will be replaced with 12 milliliters of fresh medium per plate [2].
  - 8.3.1. Aspirate EDTA and discard
  - 8.3.2. Arm adding medium
- 8.4. The system will shake the plates at 2000 revolutions per minute for 1 minute to dislodge the colonies [1] before triturating the cells with five cycles of pipetting to break up the colonies into 50-80-micron clumps [2].
  - 8.4.1. Plates being shaken OR Plates being transferred to shaker
  - 8.4.2. Cells being triturated
- 8.5. Then cells will then be transferred to a 50-milliliter tube on the deck [1] before seeding them in a 1:7 split ratio [2-TXT]
  - 8.5.1. Arm adding cells to tube
  - 8.5.2. Arm adding cells to plate. **TEXT: Transfer plates to the incubator**

- 9. Automated High Content, High Throughput Imaging
  - 9.1. For automated, high content, high throughput cell imaging, execute the **Imaging** method [1] and select a batch that contains at least one assay plate and no culture plates [2].
    - 9.1.1. WIDE: Talent executing method, with monitor visible in frame
    - 9.1.2. SCREEN: screenshot\_11: 00:08-00:11
  - 9.2. Then system will then transfer an assay plate to the automated confocal microscope for imaging [1-TXT].
    - 9.2.1. Arm transferring plate *Videographer: Important step* **TEXT: Initiate plate** return to incubator after imaging

# **Protocol Script Questions**

**A.** Which steps from the protocol are the most important for viewers to see? 5.4., 6.3., 7.2., 7.3., 9.2.

- **B.** What is the single most difficult aspect of this procedure and what do you do to ensure success?
- 2.2. Reloading of culture plates or other consumables. Please pay attention to proper alignment of the plates/ tips racks while loading. Misplacements can cause transport error and thus abort the method and subsequent steps.

# Results

- 10. Results: Representative Automated System-Produced hiPSC-Derived Cortical and Dopaminergic Neuron Characterization
  - 10.1. The hiPSC cultures should be monitored daily for growth [1] and analyzed for their percentage of confluency in the brightfield imaging cytometer [2].
    - 10.1.1. LAB MEDIA: Figure 3 Video Editor: please sequentially emphasize left and right Figure 3A images
    - 10.1.2. LAB MEDIA: Figure 3 Video Editor: please emphasize Figure 3B
  - 10.2. After passaging and manual or automated system culture [1], the cells exhibit a typical stem cell [2] and pluripotency marker expression [3].
    - 10.2.1. LAB MEDIA: Figure 4A *Video Editor: please sequentially emphasize left and right images*
    - 10.2.2. LAB MEDIA: Figures 4B and 4C *Video Editor: please emphasize OCT4 and SEEA4 images*
    - 10.2.3. LAB MEDIA: Figures 4B and 4C Video Editor: please sequentially emphasize graphs in Figure 4C
  - 10.3. Neurons differentiated in the automated culture system [1] demonstrate a similar morphology and neuronal network organization [2] to neurons cultivated manually [3].
    - 10.3.1. LAB MEDIA: Figure 5A
    - 10.3.2. LAB MEDIA: Figure 5A Video Editor: please emphasize Automated image
    - 10.3.3. LAB MEDIA: Figure 5A Video Editor: please Manual image
  - 10.4. After 6 days of differentiation, automatically differentiated cortical neurons [1] are positive for neuron-specific class three beta-tubulin [2] and upper cortical layer marker expression [3].
    - 10.4.1. LAB MEDIA: Figure 5B
    - 10.4.2. LAB MEDIA: Figure 5B Video Editor: please emphasize TUBB3 image
    - 10.4.3. LAB MEDIA: Figure 5B *Video Editor: please emphasize BRN2 image*
  - 10.5. After 8 days, the cells also express [1] microtubule-associated protein 2 [2], the neural cell adhesion molecule [3], and Synapsin-1 [4], as well as cortical neuron markers [5].

10.5.1. LAB MEDIA: Figure 5C

10.5.2. LAB MEDIA: Figure 5C *Video Editor: please emphasize MAP2 graph* 10.5.3. LAB MEDIA: Figure 5C *Video Editor: please emphasize BRN2 graph* 

10.5.4. LAB MEDIA: Figure 5C Video Editor: please SYN1 graph

10.5.5. LAB MEDIA: Figure 5C Video Editor: please CUX2 and NCAM1 graphs

10.6. Very low or no expression of these markers is observed in hiPSC [1].

10.6.1. LAB MEDIA: Figure 5C

- 10.7. The system can also be used to establish a live-cell automated neurite outgrowth assay [1] that allows neurite lengths to be measured over 11 days of differentiation without manual intervention [2].
  - 10.7.1. LAB MEDIA: Figures 6A and 6B *Video Editor: please sequentially add Figure 6B images under Figure 6A images*
  - 10.7.2. LAB MEDIA: Figure 6C *Video Editor: please emphasize data lines or no animation*
- 10.8. Using the automated culture system to perform medium changes over a 65-day culture period [1] facilitates hiPSC differentiation into midbrain dopaminergic neurons with the expected cellular organization and morphology [2].

10.8.1. LAB MEDIA: Figure 7

10.8.2. LAB MEDIA: Figure 7 Video Editor: please sequentially emphasize Figures 7B, 7C, and 7D

# Conclusion

### 11. Conclusion Interview Statements

- 11.1. <u>Ashutosh Dhingra</u>: Be sure to select the correct batches and plate ids when performing the required steps [1].
  - 11.1.1. INTERVIEW: Named talent says the statement above in an interview-style shot, looking slightly off-camera (6.1., 7.1., 8.1., 9.1.)
- 11.2. <u>Elisangela Bressan</u>: In addition to the neurite-outgrowth assay, other automated phenotypic assays relevant to neurodegeneration, for example, TDP-43 translocations, RNA foci, and fibril uptake, can be explored using this high-throughput, high-content format [1].
  - 11.2.1. INTERVIEW: Named talent says the statement above in an interview-style shot, looking slightly off-camera