Trophoblast organoids with physiological polarity model placental structure and function

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ABSTRACT

Human trophoblast organoids (TOs) are a three-dimensional *ex vivo* culture model that can be used to study various aspects of placental development, physiology, and pathology. Previously, we showed that TOs derived from full-term human placental tissue could be used as models of trophoblast innate immune signaling and teratogenic virus infections. Here, we developed a method to culture TOs under conditions that recapitulate the cellular orientation of chorionic villi *in vivo*, with the multi-nucleated syncytiotrophoblast (STB) localized to the outer surface of organoids and the proliferative cytotrophoblasts (CTBs) located on the inner surface. We show that standard TOs containing the STB layer inside the organoid (STB^{in}) develop into organoids containing the STB on the outer surface (STB^{out}) when cultured in suspension with gentle agitation. STB^{out} organoids secrete higher levels of select STB-associated hormones and cytokines, including human chorionic gonadotropin (hCG) and interferon (IFN)-λ2. Using membrane capacitance measurements, we also show that the outermost surface of STB^{out} organoids contain large syncytia comprised of >50 nuclei compared to STB^{in} organoids that contain small syncytia (<10 nuclei) and mononuclear cells. The growth of TOs under conditions that mimic the cellular orientation of chorionic villi *in vivo* thus allows for the study of a variety of aspects of placental biology under physiological conditions.

INTRODUCTION

Three-dimensional organoid culture models from tissue-derived stem cells have emerged as important *ex vivo* systems to study a variety of aspects of the physiological and pathological states of their tissues of origin. Established organoid models often preserve key features of their source organs, including tissue organization and composition, expression signatures, immune responses, and secretion profiles. Importantly, organoid cultures can be propagated long-term and can often be cryopreserved, and thus have the capacity to serve as powerful *in vitro* tools even in the absence of access to new donor tissue. Over the past several years, trophoblast organoids (TOs)

derived from human placentas at different gestational stages have emerged as models by which

to study trophoblast development and biology, congenital infections, and innate immune

defenses[1-4]. We have shown that TOs can be derived and cultured from full-term human placental

tissue and used to model trophoblast immunity and teratogenic viral infections[1].

In tissue-derived TOs models, trophoblast stem/progenitor cells are isolated from

placental chorionic villi by serial dissociation with digest solution followed by mechanical

disruption (in the case of full-term tissue), then are embedded within an extracellular matrix (ECM,

such as Corning Matrigel) 'domes'. The domes containing isolated trophoblast stem/progenitor

cells are then submerged in growth factor cocktail-reconstituted growth media to support

stem/progenitor cell proliferation and differentiation and promote their self-organization into

mature organoid units. TOs differentiate to contain all trophoblast subtypes present in the human

placenta, including proliferative cytotrophoblasts (CTBs), which differentiate into the

multinucleated non-proliferative syncytiotrophoblast (STB), and invasive extravillous trophoblasts

(EVTs). Human chorionic villi are covered by an outermost STB layer and an inner CTB layer that

fuses to replenish the outer STB during pregnancy. However, TOs cultured as three-dimensional

organoids embedded in ECM develop with the opposite polarity and mature organoids contain an

inward-facing STB (STB^{in}) and an outward-facing CTB[1,3]. This inverse polarity limits the utility of

TOs for studies that require access to the STB layer. For example, STB^{in} TOs may not recapitulate

the vertical transmission route of teratogenic infections, the transport of nutrients and antibodies

across the STB, or the directionality of hormones and other factors that are critical for

communication to maternal tissues and cells.

To overcome the limitation of existing TO models, we developed a suspension culture

method to reverse the polarity of TOs such that the STB layer is outward facing (STB^{out}). Similar

approaches have been developed and applied to a variety of epithelial-derived organoid models[5-

9]. We show that this culture method not only reverses the polarity of STB^{in} TOs but enhances the

secretion of hormones and cytokines associated with the STB. Furthermore, we performed patch

clamping of STB^{in} and STB^{out} TOs to measure the size of cells comprising the outermost layer of

these organoids and found that STB^{out} organoids are covered by large syncytia (>50 nuclei),

whereas STB^{in} TOs contain smaller syncytia (<10 nuclei) and are largely composed of

mononuclear cells. The STB^{out} TO culture model described here thus better reflects the

physiological and pathological processes of the human placenta, which can facilitate studies to

define the underlying mechanisms of normal and diseased placental conditions.

RESULTS

Generating STB^{out} trophoblast organoids

Like epithelial-derived organoids, TOs cultured within ECM domes have an inward facing

apical surface[5,6]. However, the polarity of epithelial-derived organoids can be reversed by

culturing of mature organoids under suspension culture conditions, which can occur within ~24

hrs of initiating these cultures[5-8]. Given this, we developed a TO culturing approach that involved

the culturing of organoids for 7 days in Matrigel domes to promote their differentiation and

maturation, then the release of organoids from Matrigel. Once released, organoids were cultured

for an additional period of 24-48 hrs in suspension with gentle agitation (schematic, **Figure 1A**).

Unlike epithelial organoids in which polarity reversal can be distinguished based on brightfield

microscopy alone[5,6], we were unable to clearly distinguish between TOs grown in suspension

(STB^{out}) and those cultured in Matrigel domes (STB^{in}) based on brightfield microscopy alone

(**Figure 1B**). To directly compare the architecture and viability of STB^{in} versus STB^{out} TOs, we

performed H&E staining of cryo-sections in parallel to immunostaining these sections for a marker

of CTBs (ITGA6) and the STB (CGBs). Similar to first-trimester TOs[3], we found that STB^{in}

organoids formed intracellular "mini-cavities", with ITGA6-positive CTBs lining the outer surface

of the organoids and the CGB-positive STB distributing across these intra-organoid cavities,

which varied in size and quantity between organoids (**Figure 1C, top rows**). Similarly, we found

that STB^{out} organoids also formed mini-cavities, but in the case of these organoids, the ITGA6-

positive CTBs were largely localized to the intracellular compartment, with the CGB-positive STB

on the outer surface (**Figure 1C, bottom rows**). A schematic of the general architecture of these

organoids is shown in Figure 1C, left. In some cases, we did observe greater aggregation of

and/or fusion between organoids in STB^{out} TOs grown in suspension (**Figure S1A**). However, this

aggregation could be avoided by limiting the number of organoids seeded into each well while in

suspension culture (to <100 organoids) and dissociating aggregates by manual pipetting should

aggregation occurs during suspension culture.

87   To determine the kinetics of the formation of STB^{out} TOs, we cultured organoids for

between ~21-48hrs and performed three-dimensional confocal microscopy using syndecan-1

(SDC-1), a cell surface proteoglycan that localizes to the apical surface of the STB, and a pan-

trophoblast cytokeratin (cytokeratn-19). We found that the STB began to appear at the outer

surface of TOs cultured in suspicion by ~20hrs post-incubation but required 48hrs in culture to

reach maximal exterior localization (**Figure 1D, 1E**). We did not observe any increased toxicity

during this time as assessed by light microscopy, H&E staining, or lactate dehydroganse levels in

media (**Figure 1B**, **1C**, and **Figure S1B**).

Figure 1: Development of STBout trophoblast organoids. (A), Schematic of the workflow to generate STBout trophoblast organoids (TOs) from STBin TOs collected from Matrigel domes. **(B),** Brightfield images of STBin (top) or STBout (bottom) TOs at the end of their culture period. Scale, 25μm. **(C),** Left, schematic of STBin (top) or STBout (bottom) TOs representing the cellular orientation of cytotrophoblasts (CTBs, in green), extravillous trophoblasts (in blue), and the syncytiotrophoblast (in red). All schematics created using Biorender. Middle, hematoxylin and eosin (H&E) histological staining of cryosections of STBin and STBout TOs as noted at left. Right, immunostaining for ITGA6 (green) and CGBs (red) in cryosections of matched same TOs. DAPI-stained nuclei are shown in grey. Scale, 15μm **(D),** Confocal micrographs of TOs cultured as STBin (left panels) or in suspension to generate STBout for 21hrs (middle) or 48hrs (right) and immunostaining for SDC-1 (green) and cytokeratin-19 (red). DAPI-stained nuclei are in blue, Top panels were captured at the outermost surface of organoids (surface) and bottom panels were

captured at the innermost layers (inner). Orthogonal views (Orth) are shown at bottom. **(E),** Image
analysis of the extent of surface immunostaining for SDC-1 as assessed by image analysis of
SDC-1 intensity on the surfaces of TOs three-dimensional whole organoid images (shown as a
percent of STB^{in} TOs) in STB^{out} TOs cultures for 21hrs (light blue) or 48hrs (dark blue). Data are
shown as mean $\pm$ standard deviation with significance determined by a student's t-test (** p<0.01).
Symbols represent unique fields of organoids from individual replicates.
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Figure S1. Evaluation of STB^{in} and STB^{out} TOs. (A), Left, schematic of STB^{in} (top) or STB^{out}
(bottom) TOs demonstrating the aggregation that can occurs in STB^{out} TOs that results from fusion
of the STB and/or adhesion between organoid units. At right, brightfield images of STB^{in} (top) or
STB^{out} (bottom) TOs demonstrating the extent of aggregation that can occur. Scale, 150μm (top)
and 125μm (bottom). All schematics created using Biorender**. (B),** Levels of lactate
dehydrogenase (LDH) in conditioned medium from STB^{out} TOs cultured for ~48hrs. Data are
shown as 490nm absorbance normalized to total protein. Data are shown as mean $\pm$ standard
deviation with significance determined by a student's t-test (ns, not significant). Symbols represent
unique fields of organoids from individual replicates.
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Evaluation of the efficiency of generating the STB^{out} TO model at the single organoid level

To evaluate the efficiency of the STB^{out} model, and to determine whether contact between

organoids was required for STB^{out} TO formation, we developed a U-bottom ultra-low attachment

96-well plate-based culture format in which individual TO unit were cultured in suspension in an

individual well. To do this, we collected ~ 10 mature TO units per donor from Matrigel domes, with

lines derived from three unique placentas for a total of ~40 organoids, in Matrigel domes. Then,

individual organoids were cleared from traces of Matrigel using a needle under a tissue culture

microscope and placed into a well of a U-bottom plate for 48 hr suspension culturing (**schematic,**

Figure 2A, **2B, top**). Individual organoids were then fixed and immunostained for ITGA6 and

SDC-1 to determine the orientation of the STB. We retrieved 37 single suspension cultured

organoid units after immunostaining and found that of these organoids, 34 exhibited >50% exterior

SDC-1 immunolocalization compared to matched STB[in] organoids, in which only 1 of 37 organoids

exhibited this orientation (**Figure 2C**).

Figure 2. Single organoid culturing to generate STB[out] TOs. (A), Schematic depicting the
workflow to generate STB[out] TOs as single organoids in a 96-well plate format. (B), Top, brightfield
images of STBout TOs cultured as single organoids. Scale bar at top, 400µm Bottom,
representative images of three independent STBout TO wells immunostained for the STB marker
SDC-1 (in green). DAPI-stained nuclei are shown in grey. Scale bar in bottom, 30µm. **(C),** A total
of 37 organoids from three independent TO lines were cultured as single organoids and then fixed
an immunostained for SDC-1 as shown in (B). Organoids with >50% SDC-1 immunolocalization
on the exterior surface of the organoid (SDC-1 exterior, in green) were quantified and shown as
a percentage of total organoids.

There-dimensional imaging to define cell populations and their localization in STB[in] and

STB[out] TOs

To further define the localization of key trophoblast cell populations between STB[in] and

STB[out] TOs, we performed whole organoid immunostaining followed by three-dimensional

confocal microscopy. SDC-1 localized to   interior min-cavities in STB[in] TOs (**Figure 3A,**

Supplemental Movie 1). In contrast, SDC-1 almost exclusively localized to the outermost

surfaces of STB[out] TOs (**Figure 3A**, **Supplemental Movie 2**). Similarly, CGBs localized to the

inner-most surfaces of STB[in] TOs and were surrounded by outer layers of ITGA6-positive CTBs

(**Figure 3B, Supplemental Movie 3**). In contrast, CGB localization was on the exterior of STB[out]

TOs, with ITGA6-positive CTBs comprising the interior of organoids (**Figure 3B, Supplemental**

Movie 4).

The STB is a primary producer of hormones required for pregnancy, including human

chorionic gonadotropin (hCG) which is comprised of two subunits, CGBs and CGA. We and others

have shown that STB[in] TOs recapitulate this secretion[1,3]. To determine if there were differences

in the secretion of hCG between STB[in] and STB[out] TOs, we performed Luminex assay of

conditioned medium from both STB[in] and STB[out] TOs wells. To control for variability in organoid

size and quantity, we also collected and quantified total protein used to normalize these values.

We found that there were significantly higher levels of hCG in media collected from STB[out] TOs

compared to STB[in] TOs (~10000pg/mL versus ~7000pg/mL, respectively) (**Figure 3C**), consistent

with the enhanced localization of the STB to the outer surfaces of TOs.

TOs can self-differentiate to contain small amounts of HLA-G[+] EVTs[1,2]. To determine if

this also occurred or was altered by STB[out] suspension culture conditions, we performed

immunostaining for HLA-G in STB[in] and STB[out] TOs. Consistent with our previous study, we found

that STB[in] TOs differentiated to contain <10% HLA-G[+] EVTs (**Figure 3D, 3E**). In contrast, we were

unable to detect any HLA-G^{+} cells in STB[out] TOs, suggesting that suspension culture condition

reduces spontaneous EVT differentiation. Although rates of EVT differentiation can be promoted

by altering composition of culture media[1,2], this process requires an extended culture period of >3-

4 weeks, which is beyond the time frame possible to culture STB[out] TOs in suspension. Thus, it

remains unclear if these STB[out] organoids can also be cultured to promote EVT differentiation.

Figure 3: Confocal microscopy for distinct trophoblast markers in STB^in and STB^out TOs.
(A), Cross-sections of STB^in (top) of STB^out (bottom) TOs immunostained for SDC-1 (green), and cytokeratin-19 (red). DAPI-stained nuclei are in blue. At bottom and right are orthogonal views of three-dimensional stacked images. Movies demonstrating image reconstruction and sectioning are in Supplemental Movies 1 and 2. **(B),** Confocal micrographs of STB^in (top) or STB^out (bottom) TOs immunostained for ITGA6 (in green) or CGBs (in red). DAPI-stained nuclei are shown in grey. Right are individual channels, bottom is orthogonal views of three-dimensional stacked images. Movies demonstrating image reconstruction and sectioning are in Supplemental Movies 3 and 4. Scale bar, 30μm. **(C),** Levels of human chorionic gonadotropin (hCG) (shown as pg/mL normalized to total protein of wells from which CM was collected) in conditioned medium collected from STB^in or STB^out TOs wellsas determined by Luminex. **(D),** Confocal micrographs of STB^in (top) or STB^out (bottom) TOs immunostained for ITGA6 (in blue), SDC-1 (in green), and HLA-G (in red). DAPI-stained nuclei are shown in grey. At right are individual channels. Scale bar, 30μm **(E),** Quantification of the percentage of HLA-G$^+$ positive cells in STB^in versus STB^out TOs. Individual points represent unique fields used for image analysis. A minimum of 15 organoids was used for analysis. Significance determined by a student's t-test (** $p<0.01$).

Profiling of cytokine and chemokine secretion in STB^in and STB^out organoids

In addition to hormones, the STB also secretes cytokines required to facilitate the establishment of tolerance and/or to defend the fetus from pathogen infection, such as the release of the antiviral type III interferons (IFNs) IFN-λs[11]. We showed previously that TOs recapitulate this secretion and release a number of these cytokines, including IL-6 and IFN-λ2[1]. To determine if STB^out TOs maintain this cytokine secretion or induce unique cytokines and chemokines compared to STB^in TOs, we performed multiplex Luminex profiling of 73 cytokines and chemokines, a subset of which we previously showed were released from STB^in TOs[1]. We did not observe any secretion of cytokines and chemokines in STB^out TOs that were not also secreted from STB^in TOs (**Figure 4A**). However, we found that STB^out TOs secreted higher levels of two factors, IFN-λ2 (>7-fold increase) and IL-6 (>4fold increase), (**Figure 4B-D**). In contrast, other analytes such as GRO-α/MIF were secreted at similar levels between both STB^in and STB^out TOs (**Figure 4A, 4E**). One chemokine, CXCL1, which has been associated with decidual stromal cell responses to trophoblasts[12] and which uses SDC-1 as a co-receptor[13], was significantly reduced in STB^out TOs (>12-fold reduction) (**Figure 4B, 4F**). Collectively, these suggest that STB orientation in TOs has implications on immune secretion profiles.

Figure 4: Levels of STB-associated cytokines and chemokines in STB^{in} and STB^{out} TOs.
(A), Heatmap depicting the levels of cytokines and chemokines in conditioned medium (CM)
isolated from three independent codes of TOs derivedfrom unique placental tissue. Three
replicate wells of each condition are shown. Analytes with detected values at or above 500ng/mL
are shown in red and analytes with no to little detection are shown in blue, as shown in scale at
bottom. **(B),** Heatmap of cytokines and chemokines released from STB^{out} TOs. Data are shown
as a fold-change from STB^{in} organoids (blue is increased and grey is decreased levels). Red
asterisks designate factors increased in STB^{out} TOs by >2-fold or decreased >2-fold. Data are
shown from twelve independent CM preparations from at least 3 codes, with average shown at
right. **(C-F),** Levels of IFN-λ2 (C), IL-6 (D), MIF (E), or CXCL1 (F) from CM collected from STB^{in}
or STB^{out} organoids as determined by Luminex assays. All data are shown as pg/mL and
normalized to total protein of wells from which CM was collected. All data are also shown as mean

227 ± standard deviation with significance determined by a student's t-test (****, p<0.0001, ns, not
228 signfiicant). Symbols represent unique media samples collected from replicate experiments from
229 at least three unique codes.
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Membrane capacitance measurements confirms the presence of large syncytia on the

exterior surface of STB^{out} TOs

Cell fusion dramatically increases the surface area of the fused cell. As cell surface area

is proportional to its membrane capacitance (Cm)[14], patch clamp, a quantitative

electrophysiological technique[15,16], can be used to evaluate cell size. We therefore utilized patch

clamping to calculate the size of cells/syncytia comprising the exterior cellular surface of STB^{in}

versus STB^{out} TOs (schematic, **Figure 5A**). When a small voltage step (10 mV) was applied to

TO lines derived from three unique placentas, the capacitive current from STB^{out} TOs showed

much slower decay than the capacitive current from STB^{in} TOs (**Figure 5B**). The average Cm in

STB^{in} were 0.238 nF, 0.076 nF and 0.104 nF for code 1, code 2 and code 3, respectively. Whereas

the average Cm in STB^{out} TOs were 2.812 nF, 1.603 nF and 1.899 nF, respectively for the three

codes. (**Figure 5C**). Interestingly, the Cm of the surface trophoblasts in STB^{out} TOs exhibited a

Gaussian distribution in all lines tested (**Figure 5D-F**). In stark contrast to the broader distribution

of the Cm from STB^{in} TOs centered at 0.113 nF, 0.051 nF and 0.087 nF for code 1, code 2 and

code 3, respectively, the Cm from the STB^{out} TOs was largely centered at 3.350 nF, 1.230 nF and

1.622 nF, about 20-30 fold larger than in STB^{in} TOs. It is worth noting that extremely large syncytia

are readily observed on the surfaces of STB^{out} TOs (**Supplemental Figure 2A**). We recorded 5

independent areas of these cells and found that they have unmeasurable cell capacitance

(**Supplemental Figure 2B**). This is likely due to space clamp issues for syncytia with extremely

large surface areas[17]. It is noteworthy to mention that the smallest Cm measured in our study,

which represents the Cm of a single cell, was approximately 30 pF. Syncytialization involves

fusion of plasma membranes, and Cm is directly proportional to the cell surface area. Therefore,

the extend of syncytialization in STB^{in} and STB^{out} TOs can be estimated by dividing the Cm of the

fused cell by the single cell capacitance of 30 pF. By performing the necessary calculations, we

found that the surface trophoblasts observed in the STB[in] samples were predominantly composed

of individual cytotrophoblasts (CTBs) and syncytia with limited fusion (less than 10 nuclei). On the

other hand, the surface trophoblasts in the STB[out] samples are primarily comprised of syncytia

with a greater number of nuclei, exceeding 50 nuclei.

Figure 5: Evaluation of trophoblast fusion on the surface of STB[in] and STB[out] TOs using membrane capacitance measurement. (A), Top, diagram of whole-cell patch clamp to measure membrane capacitance (Cm), which is proportional to cell surface area. Rs: series resistance; Rm: membrane resistance. Bottom**,** representative brightfield images of patch-clamped surface trophoblasts from the TOs growing under STB[in] (left) or STB[out] (right) conditions. **(B),** Representative membrane test traces from STB[in] (dashed lines) and STB[out] (solid lines) TOs to measure cell capacitance from three TO codes derived from independent placental tissues (in blue, red, and green). Current was elicited by a test voltage pulse of 10 mV from a holding potential of 0 mV (top). **(C),** Summary of membrane capacitance measured from STB[in] (grey) and STB[out] (blue, red, and green) TOs derived from independent placental tissues. Two-sided Student's t-test. (n=31 for both STB[in] and STB[out] code 1; n=26 and 28 for STB[in] and STB[out] code 2, respectively; n=22 and 29 for STB[in] and STB[out] code 3, respectively. ***p<0.001, ****p<0.0001). **(D-F)** Distribution of cell capacitance from STB[in] (black) and STB[out] TOs from three TO codes derived from independent placental tissues (in blue (D), red (E), and green (F)). The bars were at the center of each bin (see Methods for details). The data were fitted with Gaussian distribution.

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Figure S2. Patch clamp measurement of large syncytia from the surface of STB^out TOs. (A),
Representative brightfield image of a patch-clamped, extremely large syncytia from trophoblast
organoids (TOs) growing under STB^out conditions. **(B),** Representative membrane test trace from
large STB. Cell capacitance cannot be accurately measured due to the space clamp issue of large
syncytia. Current was elicited by a test voltage pulse of 10 mV from a holding potential of 0 mV
(top).
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DISCUSSION

In this study, we develop a method to culture trophoblast organoids under conditions that

reflect their physiological cellular orientation *in vivo*. This model facilitates access to the STB layer

while also maintaining key features associated with STB^in TOs, including their three-dimensional

morphology, the presence of distinct trophoblast subpopulations, and the secretion of pregnancy

related hormones and immune factors. STB^out TOs have several advantages over STB^in TOs. For

example, STB^out TOs naturally self-reorganize with an STB outward-facing surface and do not

require extensive manipulation to develop this outer layer. In addition, as STB^out TOs are cultured

in suspension, the lack of ECM allows for applications in which this scaffold presents a barrier to

diffusion, such as studies of microbial infections or antibody uptake.

For epithelial organoids grown in ECM domes with basal-out polarity, microinjection can

serve as an option to directly access the enclosed apical surface[18,19]. However, in contrast to

epithelial-derived organoids which often form clear cystic structures, TOs have heterogeneous

mini-cavities, which makes microinjection of these organoids difficult. Additional methods have

been applied to epithelial-derived organoids, such as seeding dissociated organoid fragments

onto Transwell inserts[20,21]. However, this approach compromises the three-dimensional nature of

organoids which may impact their function. The method we describe here avoids several of these

challenges, as STB^{out} TOs maintain their three-dimensional structure and do not require their

disruption to generate. It is unclear whether STB^{in} TOs undergo similar mechanisms of polarity

reversal as do epithelial-derived organoids, which undergo relocalization of junction-associated

proteins to mediate this process, or whether culturing in suspension instead promotes CTB fusion

on the organoid surface. Given that the surface of STB^{out} TOs is covered by very large syncytia,

it is possible that suspension culturing promotes the fusion of CTBs on the organoid surface rather

than inducing a relocalization of the STB from the inner to outer organoid surface. This fusion

could be promoted by factors including low levels of shear stress during suspension culturing,

which has been proposed to enhance rates of CTB fusion[22]. Fluid shear is known to impact myriad

aspects of epithelial cell function, including formation of microvilli[23], suggesting that this shear

likely also impacts rates of CTB fusion.

A benefit of TOs is their ability to recapitulate the hormone and cytokine secretion

observed in primary trophoblasts and chorionic villous tissue explants[1], which is not recapitulated

in standard trophoblast cell lines[11]. However, given that STB^{in} TOs are embedded in Matrigel,

many of these STB-associated factors would be secreted into the center of the organoid structure

or perhaps into the surrounding ECM. We found that STB^{out} TOs not only recapitulate the release

of these factors, but that some factors were secreted at significantly higher levels than those

observed in STB^{in} TOs. The mechanistic basis for this is likely two-fold and could include the

increase in syncytia size on the STB^{out} TO surface as well as the direct release of these factors

into the culture media. However, we found that STB^{out} organoids exhibited substantially lower

levels of HLA-G^{+} EVTs, suggesting that there is reduced EVT differentiation, which could impact

EVT-specific secretion profiles. It is not clear whether methods to promote EVT differentiation

previously applied to TOs derived from full-term tissue[1] could also be applied to the STB^{out} TO

system. However, given the extended time to perform this procedure (>3 weeks), it is unlikely that

STB^{out} TOs would be amenable to this process.

We leveraged the power of electrophysiology to define the size of cells/syncytia covering

the surface of STB^{in} and STB^{out} TOs. These studies verified the high efficiency of the STB^{out} TO

system and provided quantitative measurements of the number of nuclei comprising syncytia.

Based on these findings, we estimate that syncytia covering STB^{out} TOs were comprised of at

least 60 nuclei as well as some syncytia that were too large to be measured by patch clamping.

These studies not only confirmed the presence of syncytia on the outer surface of STB^{out} TOs but

provide a strong proof of concept for the application of this approach to quantitatively measure

syncytial size on the surfaces of TOs, which could be applied to a variety of biological questions.

The STB^{out} model we describe does have limitations. We have applied the protocol

described above to multiple lines of TOs derived from unique full-term placental tissue. Although

we would anticipate that this protocol can be adapted to TOs derived from early gestation tissue

as well as from iPSC-derived organoids, it is possible that some of the steps described would

require further optimization for these models. The most significant limitation of this approach is

that STB^{out} organoids cannot be passaged to maintain this physiological polarity phenotype, and

TOs with this orientation must be generated for each experiment. Thus, generation of STB^{out} TOs

is considered a terminal culture approach, like existing EVT differentiation methods[2]. However,

given that STB^{out} polarity is maintained post-culturing, these organoids can be utilized for a studies

post-generation.

The model described here provides an organoid system that recapitulates the cellular

orientation of the human placenta *in vivo* and provides evidence that this system can be used to

model key aspects of STB structure and function. In addition, given that we have developed a

system for single organoid culturing in a multi-well format, this approach could be used for mid-

to-high throughput screening approaches. Collectively, our method described here can be used

model key aspects of placental physiology and development.

MATERIALS AND METHODS

Trophoblast organoid culturing

TO lines used in this study were derived from human full-term placentas as described previously[1].

For passaging and culturing, TOs were plated into Matrigel (Corning 356231) domes, then

submerged with prewarmed complete growth media as described[1]. Cultures were maintained in

a $37^{o}C$ humidified incubator with 5% CO_2. Medium was renewed every 2-3 days. About 5-7 days

after seeding TOs were collected from Matrigel domes, digested in prewarmed TrypLE Express

(Gibco, 12605-028) at $37^{o}C$ for 8 min, then mechanically dissociated into small fragments using

an electronic automatic pipettor and further manually pipetting, if necessary, followed by seeding

into fresh Matrigel domes in 24-well tissue culture plates (Corning 3526). Propagation was

performed at 1:3-6 splitting ratio once every 5-7 days. For the first 4 days after re-seeding, the

complete growth media was supplemented with an additional 5 µM Y-27632 (Sigma, Y0503).

Derivation of STB^{out} TOs by suspension culturing

To generate STB^{out} TOs, mature STB^{in} organoids cultured as described above were first released

from Matrigel domes using cell recovery solution (Corning, 354253) on ice with constant rotating

at high speed (>120 rpm) for 30~60 min, pelleted, washed one time with basal media (Advanced

DMEM/F12 + 1% P/S + 1% L-glutamine + 1% HEPES) and resuspended in complete growth

media supplemented with 5 µM Y-27632. Organoids were then carefully transferred using FBS

pre-coated wide orifice p200 pipette tips (Fisher Scientific, 02-707-134) into an ultra-low

attachment 24-well plate (Corning, 3473). One dome containing ~ 500 organoids units can be

dispensed into up to 5 wells of a 24-well plate with < 100 organoids units per well. TOs were

evenly distributed in the wells prior to culturing in a 5% CO2 37℃ incubator for suspension culture

of 1-2 d. Constant orbital rotating was introduced into suspension culture to improve polarity

reversal efficiency (Thermo Fisher, 88881103). Media was renewed daily, and any aggregates

dissociated using a FBS pre-coated wide orifice p200 pipette tip.

378   Single organoid STB^{out} suspension cultures in 96-well plates

379   To perform single organoid unit suspension culture, each individual mature STB^{in} TO unit was

380   picked out from domes with a sterilized needle (BD, 305125), including removal of any trace

381   Matrigel matrix without compromising organoid integrity using a light microscope. Isolated

382   organoids were then placed into a U-bottom well of an Ultra-low attachment 96-well spheroid

383   microplate (Corning, 4515) for 2d suspension culturing as described above.

385   Collection of conditioned media

386   Conditioned media (CM) was collected from original STB^{in} in domes as described [1]. To harvest

387   CM from STB^{out} TOs in suspension culture, the suspension culture 24-well plate was tilted for ~2

388   min to sediment organoids to one side of the well, then carefully aspirate the supernatant media

389   without disturbing the bottom organoids. Parallel STB^{in} and STB^{out} TOs wells used for CM

390   collection contained approximately same initial number of organoids for following analysis.

392   STB^{in} and STB^{out} TOs total protein extraction and quantification

STB^{in} TOs in Matrigel domes were released and collected as previously described[1], then total

protein was extracted using RIPA buffer containing proteinase inhibitor and sonication at 10

amplitudes, after 10 min incubation on ice, centrifuge at high speed > 12000 g for 10 min, finally

collect the supernatant as organoids lysate. For the STB^{out} TOs total protein extraction, same

protocol was used except skipping the organoids releasing step. Total protein quantifications were

performed using a BCA Protein assay kit (Pierce, 23227) according to the manufacturer's

instructions.

Immunofluorescence microscopy

STB[in] TOs were immunostained as described[1]. For staining of STB[out] TOs in suspension, the same protocol described was used, but the releasing of organoids from Matrigel was omitted. The following antibodies or reagents were used: SDC-1 (Abcam, ab128936), hCG beta (Abcam, ab243581), ITGA6 (Invitrogen, MA5-16884), HLA-G (Abcam, ab52454 and ab283260), cytokeratin-19 (Abcam, ab9221), Alexa Fluor 488 Goat anti-Mouse IgG secondary antibody (Invitrogen, R37120). Alexa Fluor 488 Goat anti-Rabbit IgG secondary antibody (Invitrogen, R37116), Alexa Fluor 488 Goat anti-Rat IgG secondary antibody (Invitrogen, A11006), Alexa Fluor 594 Goat anti-Mouse IgG secondary antibody (Invitrogen, R37121), Alexa Fluor 594 Goat anti-Rabbit IgG secondary antibody (Invitrogen, R37117), Alexa Fluor 633 Goat anti-Mouse IgG secondary antibody (Invitrogen, A21052), Alexa Fluor 647 Goat anti-Rat IgG secondary antibody (Invitrogen, A21247), Images were captured using a Olympus Fluoview FV3000 inverted confocal microscope or a Zeiss 880 Airyscan Fast Inverted confocal microscope and contrast-adjusted in Photoshop or Fiji. Image analysis and generation of three-dimensional movies was performed using Imaris (version 9.2.1, Oxford Instruments).

Histochemistry and immunostaining of organoid frozen sections

Organoid preparation and embedding, cryosectioning, H&E staining, and immunostaining were performed as previously described[10], using an H&E stain kit (Abcam, ab245880) according to the manufacturer's instructions or standard procedure for immunostaining. Briefly, the collected STB[in] and STB[out] TOs were fixed (4% PFA/1×PBS) and permeabilized (0.5% Triton X-100/1×PBS), then submerged in 20% sucrose solution overnight, then finally embedded into the 7.5% gelatin/10% sucrose embedding solution and stored at -80 ℃. Cryosectionion of organoids frozen blocks was performed using a cryotome (Leica, CM1950) at 10 μm thickness. For immunostaining of cryosections, frozen sections were warmed to room temperature, then immunostaining performed

with the antibodies as described above. Images were captured on a Keyence BZ-X810 all-in-one

fluorescence microscope and contrast-adjusted in Photoshop.

Luminex assays

Luminex assays were performed using the following kits according to the manufacturer's

instructions: hCG Human ProcartaPlex Simplex Kit (Invitrogen, EPX010-12388-901), Bio-Plex

Pro Human Inflammation Panel 1 IL-28A / IFN-λ2 (Bio-rad, 171BL022M), Bio-Plex Pro Human

Inflammation Panel 1, 37-Plex (Bio-rad, 171AL001M), and Bio-Plex Pro Human Chemokine Panel,

40-Plex (Bio-rad, 171AK99MR2). Plates were washed using the Bio-Plex wash station (Bio-rad,

30034376) and read on a Bio-Plex 200 system (Bio-rad,   171000205). All samples from both

polarity conditions (STB^{in} and STB^{out}) were tested in duplicate, and each condition was performed

with at least three biological replicates from unique placental tissue. All measurements were

normalized to total protein of either STB^{in} or STB^{out} TOs wells, quantified as described above.

Cytotoxicity Assay

Cytotoxicity assays were performed using the CytoTox96$^{®}$ Non-Radioactive Cytotoxicity Assay

kit (Promega, G1780) according to the manufacturer's instructions. All samples were tested in

duplicate, and each condition (STB^{in} or STB^{out}) was performed with at least three biological

replicates.

Coat-seeding of STB^{in} and STB^{out} TOs onto round coverslips for patch clamp

To seed collected original STB^{in} TOs onto the round glass coverslips (VWR, 76305-514) pre-

coated with thin layer of Matrigel (Corning, 356231), each round coverslip was evenly distributed

with ~ 40 μl of Matrigel and carefully transferred into each well of regular 24-well plate to

polymerize in a 37 ℃ incubator for ~ 20 min. Then, organoids were harvested as described above

and evenly dispensed onto the Matrigel pre-coated surface of coverslips to settle down in a 5%

CO2 37 ℃incubator for 3~4 h to ensure that the majority of organoids attach onto the matrix

coating of the coverslip. For the STB[out] TOs coat-seeding, the same protocol described above

was used except omitting the release of organoids from Matrigel domes.

Patch clamp estimation of cell surface area

All results were recorded in whole-cell configurations using an Axopatch 200B amplifier

(Molecular Devices) and the pClamp 10 software package (Molecular Devices). The glass

pipettes were pulled from borosilicate capillaries (Sutter Instruments) and fire-polished using a

microforge (Narishge) to reach a resistance of 2–3 MΩ. The pipette solution (internal) contained

(in mM): 140 CsCl, 1 MgCl$_2$, 10 HEPES, 0.2 EGTA. pH was adjusted to 7.2 by CsOH. The bath

solution contained (in mM): 140 CsCl, 10 HEPES, 1 MgCl$_2$. pH was adjusted to 7.4 by CsOH. All

experiments were at room temperature (22–25°C). All the chemicals for solution preparation were

obtained from Sigma-Aldrich. Once the whole cell configuration was established, a 10-mV voltage

command was delivered to the cell from a holding potential of 0 mV. The corresponding capacitive

current was recorded. Membrane capacitance of the cell was calculated using Clampfit software

(Molecular Devices) based on the following equation, $C_m = \frac{Q}{\Delta V} = \frac{I \times \Delta t}{\Delta V}$, where Cm is the membrane

capacitance, Q is the stored charge across the cell membrane, V is membrane voltage, I is current,

and t is time. For the histogram plot, the bins (x-axis) were set as (pF): 0-20, 20-100, 100-200,

200-500, 500-1000, 1000-2000, 2000-5000 and 5000-10000. The bars on the histogram were set

in the middle of each bin.

Statistics and reproducibility

All experiments reported in this study have been reproduced using a minimum of three

independent organoids lines derived from unique placental tissues. All statistical analyses were

performed using Clampfit (Molecular Devices), Excel, or Prism software (GraphPad). Data are

presented as mean ± SD, unless otherwise stated. Statistical significance was determined as

described in the figure legends. Parametric tests were applied when data were distributed

normally based on D'Agostino-Pearson analyses; otherwise, nonparametric tests were applied.

For all statistical tests, p value <0.05 was considered statistically significant, with specific p values

noted in the figure legends.

Supporting information

Supplemental Movie 1: Three-dimensional image reconstruction of an STB^{in} trophoblast

organoid (shown in Figure 2B, top) immunostained for SDC-1 (in green) and cytokeratin-19 (in

red). DAPI-stained nuclei are shown in blue.

Supplemental Movie 2: Three-dimensional image reconstruction of an STB^{out} trophoblast

organoid (shown in Figure 2B, bottom) immunostained for SDC-1 (in green) and cytokeratin-19

(in red). DAPI-stained nuclei are shown in blue.

Supplemental Movie 3: Three-dimensional image reconstruction of an STB^{in} trophoblast

organoid (shown in Figure 2B, top) immunostained for ITGA6 (in green) and CGBs (in red). DAPI-

stained nuclei are shown in grey.

Supplemental Movie 4: Three-dimensional image reconstruction of an STB^{out} trophoblast

organoid (shown in Figure 2B, top) immunostained for ITGA6 (in green) and CGBs (in red). DAPI-

stained nuclei are shown in grey.

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AUTHOR CONTRIBUTIONS

L.Y. and C.C. conceived the study, developed the methodology, and analyzed the data; P.L. and H.Y. performed patch clamping measurement, and analyzed the data; All authors participated in manuscript writing, review, and editing.

DECLARATION OF INTERESTS

The authors declare no competing interests.

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