# High-content screening-based phenotypic analysis of 3D organotypic bronchial tissues <u>Diego Marescotti</u>, David Bovard, Moran Morelli, Karsta Luettich, Stefan Frentzel, Manuel Peitsch, Julia Hoeng PMI R&D, Philip Morris Products S.A., Quai Jeanrenaud 5, 2000 Neuchâtel, Switzerland (Part of Philip Morris

#### Introduction

Increasing requirements for more physiologically relevant in vitro systems have led to the development of more complex cellular models, which range from 2D co-cultures to 3D organotypic tissues with one or more cell types. To further evolve our High Content Screening (HCS) assay portfolio, we sought to establish HCS methods of 3D bronchial tissues to perform:

- **1. Tissue quality control**: to verify differentiated phenotype and tissue homogeneity.
- 2. Experimental endpoints: to evaluate phenotypic changes upon tissue treatment.

#### Objectives

When grown to confluence on a Transwell<sup>™</sup> system and cultured at the air-liquid interface (ALI), normal human bronchial epithelial (NHBE) cells form a polarized, pseudostratified epithelium composed of basal, ciliated, and goblet cells (Figure 1).

We investigated phenotypic markers that are known to characterize this 3D model. Cell type-specific markers, such as  $\beta$ -Tubulin 4, Muc5AC, and p63, were selected to identify ciliated, goblet, and basal cells, respectively. In addition, ZO-1, a member of the family of tight junction proteins, was included.



Figure 1. a) Representation of the bronchial pseudostratified epithelium. b) Tissue cross-section after staining with hematoxylin, eosin, and Alcian blue. c) Tissue cross-section after immunohistochemical detection of p63.

As a proof of concept, an *in vitro* model of IL13-driven goblet cell hyperplasia was evaluated. To induce goblet cell hyperplasia, 3D bronchial tissues were treated with IL13 (basolaterally) for two weeks. The resulting phenotype was examined by Muc5AC-specific antibody staining.

### Methods

**Immunostaining** *Fixation:* 4% (w/v) paraformaldehyde for 20 min  $\rightarrow$  *Blocking:* 0.5% Triton X-100 5% normal goat serum (NGS) + 2% bovine serum albumin (BSA) for 60'  $\rightarrow$  Staining: PBS with 1% normal goat serum +1% BSA overnight.

Antibodies: β-Tubulin 4 AF647 (Abcam), Muc5AC AF555 (Abcam), ZO-1 AF594 (Invitrogen), p63 (Abcam) in combination with  $\alpha$ -goat anti-mouse AF488 (Invitrogen). Hoechst 33342 was used at 1:2000 dilution. Images were acquired with the CellInsight<sup>™</sup> CX7 platform (Thermo Fisher) and analyzed with Studio<sup>™</sup> software (Thermo Fisher).



- (Figure 2b).

 $\rightarrow$  Selected phenotypic markers were used to identify the optimal focal plane for their imaging.  $\rightarrow$  Nuclear compartment was then imaged by applying a specific offset.

- (Figure 3b).

#### Phenotypic marker identification and quantification

- 4a).
- investigate
- same tissue.



# PMI SCIENCE PHILIP MORRIS INTERNATIONAL

International group of companies)

# **Method Development**

#### Autofocus and imaging strategy

HCS-based methods of 2D cell cultures rely on the nuclear staining for focal plane and for cell identification (Figure 2a).

Same approach proved to be challenging when applied to more complex 3D cultures due to their increased cellular complexity

Spatial distribution of the selected markers along the z-axis appeared to be more consistent (Figure 1b).





Nuclear area identification and quantification Nuclear staining was analyzed using the spot detection method, a flexible image analysis algorithm that can be utilized to provide generic spot analysis (Figure 3a).

Evaluation of the area of the field of view effectively covered by the tissue is enabled, allowing identification and quantification of fields where the tissue is damaged or detached as well as those where the camera reaches the border of the inserts

 $\rightarrow$  The obtained values are then used for normalizing the values obtained from the quantification of the specific markers.

Similar to the nuclear signal detection, the expression of specific phenotypic markers was also investigated by applying a spot detection-based mask, which enables the evaluation of their expression and the distribution (Figure

One of the advantages of high-content  $\frac{1}{2}$ imaging, similar to other fluorescencebased methods, is the possibility to  $\overline{3}$ markers several simultaneously (Figure 4b).

 $\rightarrow$  Such a combination enables gathering  $\stackrel{O}{\rightarrow}$ biological insights while decreasing  $\breve{\underline{\Theta}}$ cost, as more markers can be  $\overline{\circ}$ investigated simultaneously in the  $\frac{2}{3}$ 

Figure 4. a) Raw single channel image, composite image of Hoechst-stained tissue and spot detection mask image of ZO-1,  $\beta$ -Tubulin 4 and Muc5AC staining. b) Single channel and composite images of p63 and Muc5AC multiplex staining.





## Conclusions

This approach demonstrates that high-content imaging can be used to evaluate more complex 3D in vitro models. In fact, it can be exploited either as a tool, for fast and accurate quality control of the tissue, evaluating both the expression and the distribution or maturation markers, or as an experimental endpoint in studies looking at specific phenotypic changes upon treatment.

## Applications

Philip Morris International R&D is the sole source of funding and sponsor of this project.



**Declaration of Interest**