Modelling Epithelial Barrier Function And Mucociliary Clearance Using Small Airway Epithelium Cells

Anusha Gupta, Hannah Grice, Chloe Whiting, Oliver Birch, Fiona Leslie, Megan Webster Newcells Biotech Limited, The Biosphere, Drayman's Way, Newcastle Helix, Newcastle upon Tyne. UK NE4 5BX



- The small airways form a first line defence mechanism against inhaled particles and facilitate gas exchange by distributing air to the alveoli. The small airways play a crucial role in disease and are often the initial site of inflammation and obstruction in conditions such as bronchiolitis obliterans, asthma, and chronic obstructive pulmonary disease.
- Our ability to study small airway physiology in vivo is limited due to their small size and location deep within the lungs, making them difficult to visualize and access using noninvasive imaging techniques. At Newcells Biotech, we have developed a complex, multicellular system to enable the study of small airway physiology and disease in vitro.
- Primary human basal cells are expanded in culture flasks prior to seeding onto semi-permeable inserts under submerged conditions. On reaching confluency, the cells are 'airlifted', i.e. apical medium is removed, to allow air-liquid interface (ALI) culture conditions. Promoting epithelial polarisation and cell differentiation to form a heterogeneous population of epithelial cells representative of the small airways.





Days post-airlift

(2A) Gene expression (relative to maximal expression) of key junctional genes CDH1 (E-Cadherin) and TJP1 (ZO-1), throughout the differentiation period, from day 0 (day of air-lift) to day 42, indicating presence of epithelial junctions, in-line with literature data.

(2B) Representative images showing expression, and localisation of ZO1 and E-Cadherin proteins at epithelial cell: cell junctions.



(A) Gene expression data shows relative expression of epithelial KRT5, SCGB1A1. specific genes, MUC5B/MUC5AC, and FOXJ1 throughout the 28-day differentiation period, with changes in expression reflecting differentiation of basal, club, goblet and ciliated cells, respectively.

(B) Presence of key epithelial cell types, basal, club, goblet and ciliated cells, at 28-days post-airlift is confirmed by immunocytochemical detection of KRT5, CC10, MUC5B/MUC5AC acetylated-Tubulin and (AcT) respectively.

(A) Exposure of Small Airway Epithelial Cells (SAEC's) to increasing concentration of the TNF α reduces TEER, in-line with cytokine induced breakdown of the epithelial barrier.

(B) No significant increase in LDH release or decrease in ATP- or MTT- activity (formazan formation), was observed following exposure of SAEC's to TNF α , suggesting TNF α induced breakdown of the epithelial barrier was not observed as a result of cellular toxicity.

(C) Puromycin, an antibiotic with known toxic side effects in the lung, was used as a positive control. A decrease in TEER is triggered with increasing concentrations of Puromycin, indicating a loss in barrier integrity.

(D) High increase in LDH release as well as a decrease in ATP- and MTT- activity was observed following exposure of SAEC's to Puromycin, indicating drug induced cellular toxicity.

In Development

Newcells Biotech's SAEC model



1. Modelling Idiopathic Pulmonary Fibrosis (IPF) in Small Airway **Epithelial Cells (SAEC's)**

On treating differentiating SAECs with variations of cytokine mixes for 12 days, an increase in epithelial damage was observed as seen through a reduction in (1A) TEER and increase in (1B) FITC dextran permeability. The cytokine induced damage to the epithelial barrier was not caused due to cytotoxicity as (1D) LDH release was not increased.

2. Ciliary Beat Frequency

On treating differentiating SAECs with variations of cytokine mixes for 10 days (1C), a decrease in ciliary beat frequency was observed compared to the control.



Model Features:

- 3 validated primary basal cell donors
- Fully differentiated and multicellular •
- Optimised under Air-Liquid Interface conditions

Available Analytical Techniques and Data Outputs:

- Epithelial barrier integrity: TEER and FITC-dextran flux measurements.
- Epithelial damage studies: cytokine-induced barrier breakdown.
- ✓ Gene and protein expression: qPCR and immunocytochemistry.
- Biomarker analysis: ELISA based detection of secreted factors.
- Cellular toxicity assessment: LDH release and cellular metabolism.





Also available: Fibroblast-to-Myofibroblast Transition (FMT) assay.



