Development of a new 3D-Human Airway Epithelium/ Whole-blood Co-culture Model Combined with Multi-Analyte Profile (MAP) Analyses for Assessing Drug Effects

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Background



	Cytokines	
h1-related	Th2-related	Treg-related
terferon-γ terleukin-12p40 terleukin-12p70	Interleukin-4 Interleukin-5 Interleukin-13	Interleukin-10
lonocyte/ IØ-related	Others	Interleukin 2
-CSF terleukin-1 ra	FGF-basic GM-CSF	Interleukin-3 Interleukin-7

Cancer antigens

Alpha-Fetoprotein
Cancer Antigen 19-9
Cancer Antigen 125
Carcinoembryonic Antigen
Prostatic Acid Phosphatase
PSA, Free

Th2-related mediators

Th2-related mediators



We kindly acknowledge the grant from the Project Management Agency within the German Aerospace Center (PT-DLR). DLR grant number 01GG0713

Results and Conclusions

In this newly developed co-culture model of human airway epithelial cells in combination with whole blood Betamethasone exhibited its typical, cells, strong pharmacological effect profile on both, the immune and the epithelial cells: It dose-dependently inhibited a variety of pro-inflammatory mediators, being either T helper cell type 1- (Th1), Th2-, or macrophage-associated, such as interferon (IFN)-gamma, interleukin (IL)-12p70, IL-4, -5, -13 and tumor necrosis factor (TNF)-alpha, respectively. In contrast, IL-10 as anti-inflammatory mediator was upregulated after 24h of co-culture. Furthermore, epithelial cells were cultured for another 6 days showing a dosedependent effect on e.g. the monocyte chemotactic protein-1 (MCP-1) and IL-8.

From the data presented here, it is evident that the highly complex, organo-typical co-culture model provides an excellent tool to study in vitro, under in vivo-like conditions not only the pharmacokinetics and pharmacodynamics of inhaled drugs, but also the harmful effects of toxicants that get access to the human lung.



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