Inhaled Budesonide Enhances Effector Gene Expression in Humans: A Randomized Controlled **Trial and Comparison with In Vitro Analysis** R Leigh¹, EM King¹, S. Shah¹, C Dumonceaux¹, CF Rider¹, SL Traves¹, D.M. Slater¹, M.M. Kelly¹, A Miller-Larsson², R Newton¹



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Rationale & Hypothesis

Glucocorticoids act on the glucocorticoid receptor (GR/NR3C1) and, given as inhaled corticosteroids (ICSs), reduce inflammation in asthma by reducing inflammatory gene expression. Mechanisms by which corticosteroids reduce inflammatory gene expression may include the ability of GR to directly bind to key transcription factors, such as NF-κB, and thereby reduce inflammatory gene transcription. However there is an increasing body of data suggesting that corticosteroids may induce the expression of multiple anti-inflammatory or anti-asthma genes to exert their effects. While this statement is supported in vitro, there is a deficit in respect of relevant clinical information in vivo in humans. The current study was therefore designed to test the hypothesis that the ICS, budesonide, will induce the expression of multiple anti-inflammatory genes in vivo in humans.

Methods & Clinical Protocol

• Twelve healthy non-smoking, non-atopic male volunteers with normal lung function were enrolled into this prospective double-blind, placebo-controlled, randomised, two-period cross-over study (Fig. 1). Participant eligibility criteria are listed in **Table 1**.

• A single dose of inhaled placebo or budesonide (1600 μg) was administered by Turbuhaler[®]. Bronchoscopy was performed 5-6 h later, when endobronchial brushings and biopsies were obtained and samples processed for histology and gene expression analysis.

• A549 cells, primary human bronchial epithelial cells (HBE) (from, brushings), airways smooth muscle (ASM), human bronchial fibroblasts (fibroblasts) and human umbilical vein endothelial (endothelial) cells were cultured according to standard procedures.

• Total RNA was extracted using RNeasy kits (Qiagen). RNA was reverse transcribed to cDNA, and SYBR Green PCR was carried out for the indicated genes and GAPDH.

• CEL files from Affymetrix PrimeView microarrays were analysed with Partek Genomic Suite (v6.6). • Serum budesonide concentrations were measured by LC-MS/MS.

Table 1: Eligibility criteria for inclusion of healthy volunteers

- Non-smoker males aged 18-50 years, not on ICS or other corticosteroid treatments
- Negative skin prick test to common aero-allergens
- Normal lung function (FEV₁/FVC ≥ 0.7, $FEV_1 \ge 80\%$)
- Normal airway responsiveness
- $(PC_{20} \text{ methacholine} > 16 \text{ mg/ml})$ • No exposure to corticosteroids in
- the preceding 3 months
- No participation in any other clinical study in the preceding 4 weeks No associated morbidity where
- bronchoscopy was contraindicated



Fig. 1. Study design. Following screening, study participants were randomized to receive either inhaled placebo followed by inhaled budesonide, or inhaled budesonide followed by inhaled placebo. Treatment interventions were separated by a 2-3 week washout period. Randomization and treatment allocation was by research pharmacy personnel, not otherwise associated with the study.

Results



expression of additional genes was examined by real-time PCR. * P < 0.05, ** P < 0.01, *** P < 0.001.









