

Efficacy evaluation of ingredient mixtures using an in vitro model of atopic dermatitis

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Introduction

Dysfunctional Immune Response

Adaptive

Innate

Th.2 Cytokines
IL-4, IL-13

Eos attracting Chemokines

Attracting Chemokines

Policy genes: FLG, LOR, LOEs, S100s, SPRRs
Proteases: SCTE (KLKS), SCCE (KLKT)
Antiproteases: LEKTI (lympho-epithelial Kazal-type related inhibitor)

Stratum granulosum
Tight junction genes: CLDN1

TSLP

Mast cei

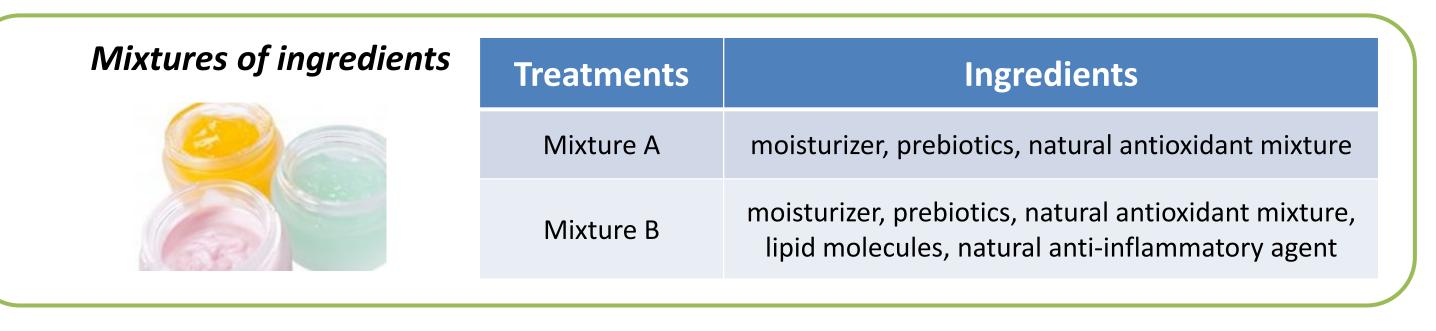
TLRs, CD14, NOD1, NOD2,

Figure 1. AD pathways [1]

Atopic dermatitis (AD) is an inflammatory skin disease, characterized by erythema and pruritus, affecting 2-20% of the general population and predominantly children [1]. AD is associated with the decrease of barrier integrity and functionality, enhanced allergen penetration and skin colonization by microorganisms such as *Staphylococcus aureus*. Thymic stromal lymphopoietin (TSLP) plays an important role in the initiation and maintenance of the allergic immune response in AD, while chemokines and cytokines such as TNF α and IL-8 play an important role in the pathogenesis of the inflammatory skin diseases [2, 3].

The aim of this study was to verify, using an *in vitro* model of atopic dermatitis, the efficacy of two mixtures of "active ingredients" mixture A and mixture B, specifically created to treat atopic babies' skin.

Methods and Materials



Endpoints

Hystology

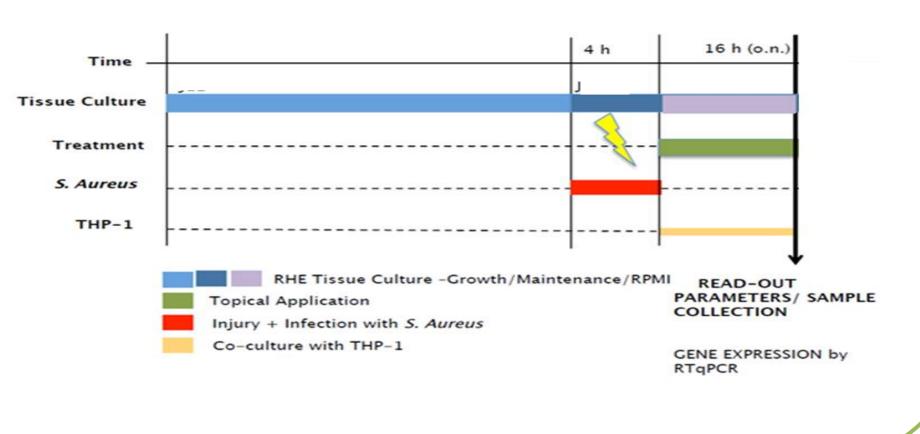
Tissues were fixed in formalin and sections of 5 μ m were obtained. Slides were stained with Haematoxilin and Eosin and analyzed under light microscopy (40x). Morphological modification, alterations at the epithelium surface and thickness tissues were analyzed.

Gene expression

Tissues were collected in lysis buffer for RNA extraction, cDNA retrotranscription and gene expression analysis. TSLP, TNF α and IL-8 were assessed. RHE in co-culture with THP1 negative control was used as calibrator control (RHE+THP1=1).

In vitro model of atopic dermatitis

3D Reconstituted Human Epidermis (RHE) model from young donors (age < 6 years) co-cultured with THP-1 human monocytic cells and infected with *S. aureus* after a slight mechanical abrasion on RHE surface [4]. The test items were prepared in PBS and topically applied on the surface of tissues for 16h as described into the following scheme:



Results

Hystology

Results reported in figure 2 show that:

- Mix A treatment was effective in maintaining a conservative morphology. It is possible to notice only a slight swelling on the surface.
- Mix B treatment has preserved tissue morphology in the viable layers: only few signs of toxicity are visible. The structure, thickness and number of stratum corneum layers is fully preserved.

The histology is comparable to a tissue without any treatment (naïve tissue); the thickness of the mix B treated tissue is 125 μ m comparable to a naïve tissue [4] as reported in the following table:

Treatments	Thickness tissue (µm)
Naïve tissue	~ 120/130
Mixture A	~ 90
Mixture B	~ 125

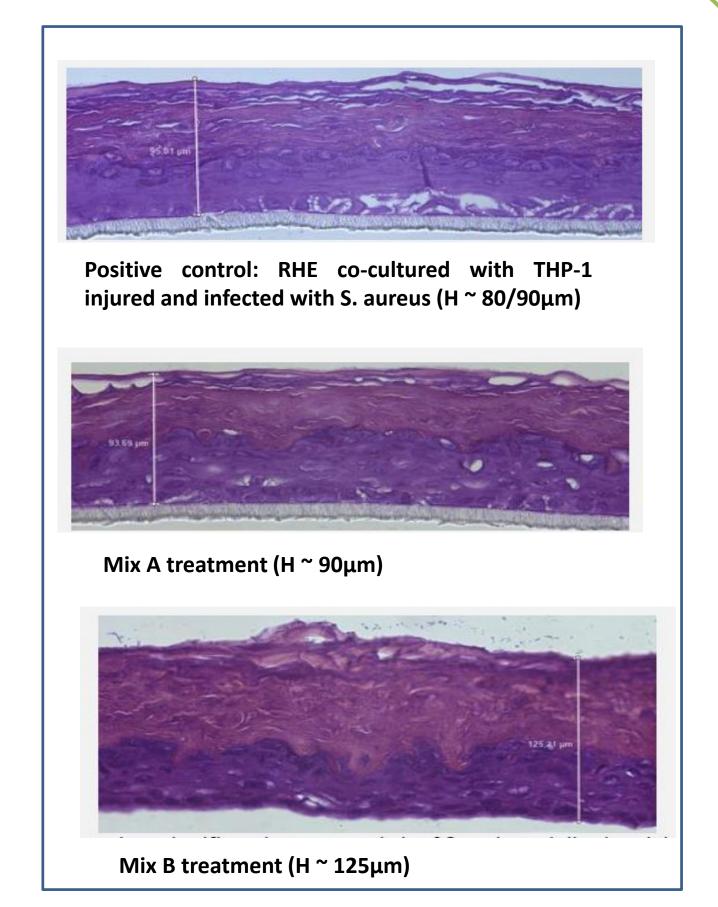
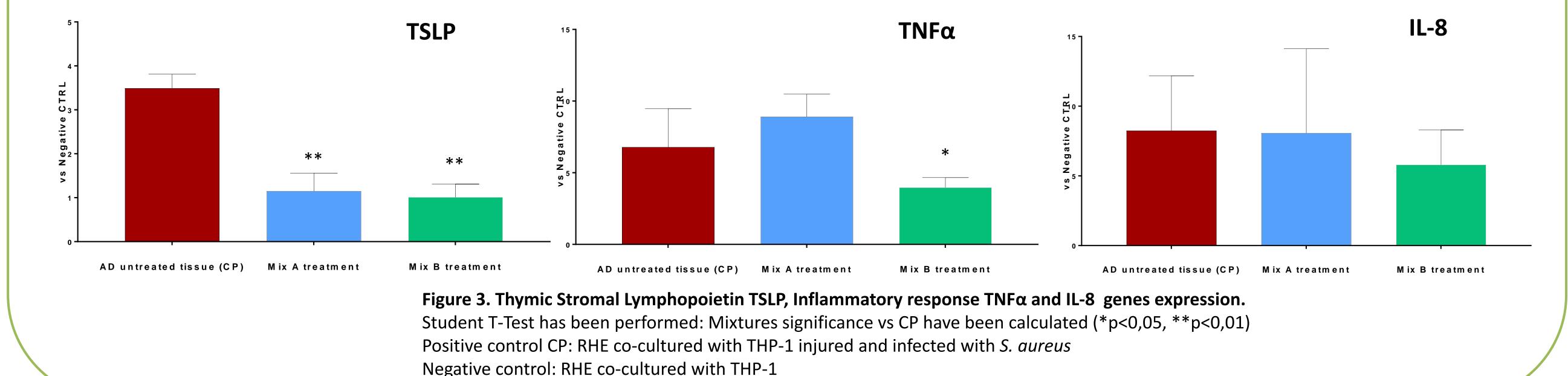


Figure 2. Histomorphological results

Gene espression

Results, expressed as fold with respect to the negative control and reported into figure 3, show that TSLP is highly expressed in the atopic skin model [4] and both mixtures reduced in significantly manner the TSLP overexpression suggesting a role in restoring/enhancing immunity response. Up regulation of IL-8 and TNF α confirms the inflammatory response [4] and mixture B was the most effective in reducing the inflammatory response reducing the IL-8 and TNF α mediated inflammation.



Conclusions

The combined use of prebiotics to boost the growth of the protective friendly skin bacteria, lipid molecules that are effective in improving barrier properties, and a bioactive compound that exhibits potential anti-inflammatory activity, could explain the reduction of inflammatory response and ameliorative effects observed on tissue by mixture B, which can be favourable used in cosmetic formulations for baby care to support atopic dermatitis treatment [5].

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