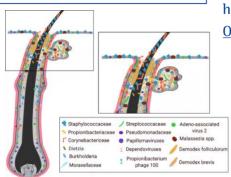
# Investigating changes in the human scalp hair follicle microbiome ex vivo



#### Claim substantiation:

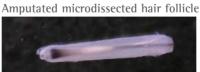
- Healthy skin with a balanced microbiome
- anti-dandruff
- anti-acne



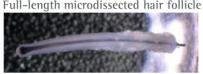
The human hair follicle is colonized by unique and complex microbiota and serves as a microbial habitat, providing a maximal surface area for microbe-host interactions regulating scalp health. Hair follicle microbiome dysbiosis is involved in hair follicle disorders, such as acne vulgaris, alopecia areata, androgenic alopecia, dandruff.

Therefore, targeting the hair follicle microbiome leads to the development of novel cosmetic strategy to manage hair disorders and maintaining scalp health.

Our models:



Full-length microdissected hair follicle



We can also investigate human skin microbiome ex vivo!

We have access to affected skin samples: Alopecia Areata, Androgenetic Alopecia, Acne Inversa,...

## Case study: Compound X restores a healthy microbiome in the hair follicle

#### 1. Compound X stimulates the production of DCD in the hair follicle

Healthy HFs manage and efficiently control their microbiome by production of antimicrobial peptides (AMPs) with different microbial specificities, and which are inducible by bacterial products/metabolites. We analyze how manipulation of the microbiome changes AMP expression and how treatment with selected compounds may help in boostering certain antimicrobial peptides.

#### Compound X



Dermcidin (DCD)

Representative images from n=6 healthy donors, qualitative observation by in situ hybridization using a DCD specific probe



# bacteria and fungi

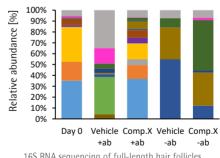
We analyze how treatment with cosmetic compounds changes the hair follicle microbiome and whether this treatment could be a suitable adjunct therapy in hair disorders characterized by dysbiosis, in order to have a more equilibrated microbiome.

2. Compound X prevents hair follicle contamination by

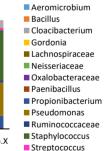




Representative images of full length hair follicle treated with vehicle or compound X in absence of antibiotics.



16S RNA sequencing of full-length hair follicles



#### **Additional Read-Out Parameters:**

Representative images of DCD positive cells (immunostaining)

Intrafollicular DCD mRNA expression

Data from n=5-6 healthy donors, analysis 6h

after treatment, mean±SEM

Vehicle w/o antibiotics

in the HF epithelium

In vitro: agar diffusion assay; MIC determination; 96 well liquid cultures and determination of viability Ex vivo: shotgun sequencing; alpha diversity and taxonomic evaluations; Pas/Lightgreen staining to quantify fungal colonies; in situ hybridization of bacteria strains; analysis of inflammatory markers, antimicrobial peptides, pigmentation, etc.

Compound X w/o antibiotics

#### Selection of our publications:

Edelkamp et al., 2023 SPP; Lousada et al., 2022 Bioessays; Bispo Lousada et al., 2021 Exp Dermatol: Lousada et al., 2021 Br J Dermatol

## Contact us for a customized study:

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## Reasons why you should choose Monasterium Laboratory:

- Cutting edge methodologies and techniques
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- A focus on novel targets and therapeutics for skin & hair disorders: identify-characterize-validate
- Delivering mechanistic action stories, biomarkers & predictors of response
- Claims support for cosmetic ingredients in skin or hair follicle models
- Clinical trials carried out with strategic partners for healthy skin and hair benefits
- Comprehensive project reports & manuscript drafting and submission

### Monasterium Laboratory

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