

# Exploration of synthetic communities as a tool to modulate the vaginal microbiome

Commonly used acronym: SynCom

Created on: 09-07-2025 - Last modified on: 16-07-2025

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# **Organisation**

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**Country** Belgium

Geographical Area Flemish Region

# **SCOPE OF THE METHOD**

The Method relates to	Human health
The Method is situated in	Translational - Applied Research
Type of method	In vitro - Ex vivo

## **DESCRIPTION**

**Method keywords** 

vaginal microbiome
synthetic communities
live biotherapeutic products
consortia
lactobacilli
vaginal infections

## Scientific area keywords

vaginal health

probiotics

microbiology

bioreactor technology

female health

Biopharmaceuticals

biotechnology

## **Method description**

This method/PhD project explores an innovative approach using synthetic communities (SynComs) to develop vaginal therapeutics. First, a top-down approach will be used to screen and select consortia of vaginal microbiota, emphasizing synergistic interactions and eliminating antagonistic effects. A bottom-up approach will validate the top-down findings, providing defined communities suitable for automation and scaling up. Simultaneously, these defined SynComs will undergo testing for their probiotic potential, comparing them with single strains through *in vitro* anti-pathobiont assays, community fitness analyses, and implementation into cutting-edge vaginal cell models. This research project will offer ground-breaking insights into a novel approach to establishing robust, safe, and effective microbiome-targeted therapies.

# Lab equipment

BSL2 lab, hypoxic chamber, biosafety cabinet, bioreactor equipment (pumps, tubing, magnetic stirring plates), qPCR, sequencing techniques (e.g., ONT minION), cell culture room, vaginal cells, etc.

#### **Method status**

Still in development

# PROS, CONS & FUTURE POTENTIAL

## **Advantages**

- Allows high-throughtput screening in mini bioreactors,
- Scale-up possible up till larger volumes (5L and 7L),
- Consortia of synergistic bacteria compared to individual strains may offer more benefits in application as LBP, such as easier niche engraftment and long-term effects due to robustness,
- Easier implemented than vaginal microbiome transplants (safety, cost, regulation-wise).

## Challenges

- Still in optimization stage,
- Culture bias may not always lead to target lactobacilli,
- Whole genomes required for safety testing and unveiling modes of action.

#### **Modifications**

Additional metabolic modeling to screen for consortia of genome-complementary lactobacilli strains (e.g., screen for potential of cross-feeding) in strain selection of defined bottom-up SynComs and *in vitro* testing

## **Future & Other applications**

This can be applied to gut microbiome research for the same purpose, namely modulating the gut microbiome in a positive way. Additionally, this can be used in environmental research, modulating the microbiome of plants, crops, other ecosystems to enhance resistance and resilience against different

types of disease.

# REFERENCES, ASSOCIATED DOCUMENTS AND OTHER INFORMATION

## References

Manuscript preprint on the use of SynComs: Vander Donck, Leonore and Victor, Maline and Van Beeck, Wannes and Van Rillaer, Tim and Dillen, Jelle and Ahannach, Sarah and Wittouck, Stijn and Allonsius, Camille Nina and Lebeer, Sarah and Administrator, Sneak Peek, Host-Independent Synergism between Lactobacillus crispatus and Other Vaginal Lactobacilli. Available at SSRN: https://ssrn.com/abstract=5243601 or http://dx.doi.org/10.2139/ssrn.5243601

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