

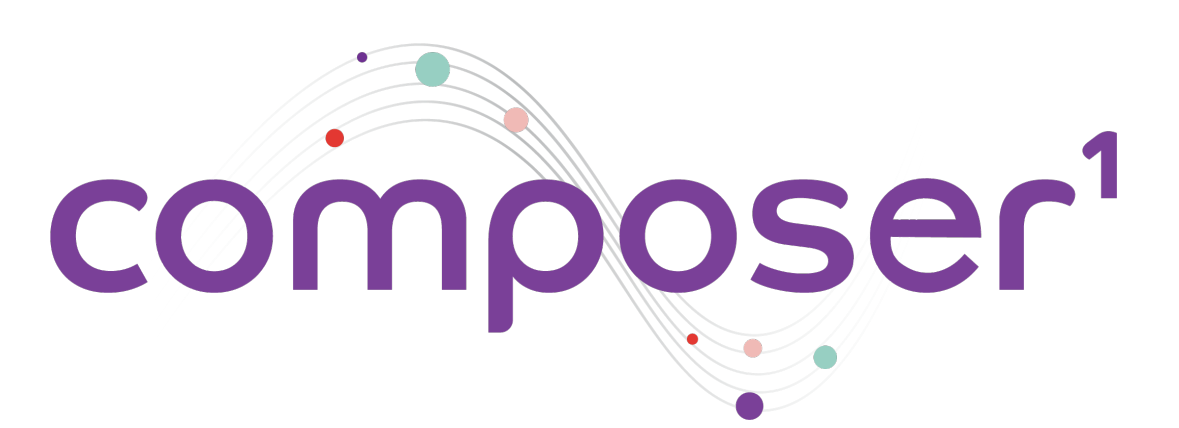
Gut commensal bacteria, associated with clinical remission in ulcerative colitis, protect and heal gut epithelial barrier through three novel pathways



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Introduction

- Precision microbiome profiling of a human fecal microbiota transplantation (FMT) study identified 8 bacteria associated with clinical response in mild-to-moderate Ulcerative Colitis (UC) patients. These 8 strains form **MB310**, a Live Biotherapeutic Product (LBP) for the treatment of UC
- In vitro primary human cell studies with MB310 species showed inflammatory response modulation and regulatory T cell (Treg) induction
- Aim of study: explore MB310 species impact on barrier function and mechanisms of action behind their protective effect**

MB310 strains improve barrier function and repair

Barrier Function

Condition	ST	FP	Sp.1	Sp.5	Sp.8
Normalised resistance	-0.8	0.8	0.2	0.7	0.1

Barrier Repair

Condition	ST	FP	Sp.1	Sp.5	Sp.8
Normalised resistance	-0.8	0.2	0.2	0.3	0.6

3 of the MB310 strains (species 1, 5 and 8) enhance epithelial barrier function and repair

Species 5 induces epithelial barrier repair by release of polyamine

Polyamine release

Condition	Control	Sp. 5
Polyamine (µM)	0.2	4.5

Inhibition of polyamine production in the barrier repair assay

Condition	ST	FP	Sp. 5	Sp. 5 + 10mM DFMO
Normalised resistance	-0.8	0.4	0.4	0.15

Species 5 produces high levels of polyamines. Blocking polyamine production by species 5 with Difluoromethylornithine (DFMO) inhibits the barrier repair

Species 8 induces epithelial barrier repair by release of nicotinic acid

Nicotinic acid release

Condition	Control	Sp. 8
Relative intensity (x1e7)	0.1	5.2

Inhibition of GPR109a in the barrier repair assay

Condition	ST	FP	Sp. 8	Sp. 8 + MPN
Normalised resistance	-0.8	0.4	0.2	0.0

Species 8 releases the metabolite nicotinic acid. The protective effect on barrier repair by species 8 is reversed by blocking the nicotinic acid receptor GPR109a with Mepenzolate (MPN)

Methods

To understand how MB310 drives clinical benefit we tested how the individual species impacted barrier function

- Barrier assays:** transepithelial electrical resistance on Caco-2 cells, untreated (Barrier Function) or damaged with LPS (Barrier Repair) to mimic inflammation. *Salmonella typhimurium* (ST) and *Fecaelibacterium prausnitzii* (FP) were used as negative and positive controls, respectively
- Mechanisms of action** were investigated through
 - Comparative genomics:** comparing strains of same species with positive and negative effect on barrier function
 - Cloning** of novel protein into *E. coli*
 - Tight junction protein expression:** measured by qPCR
 - Metabolomics:** determination of bacterial metabolites in bacterial culture supernatants
 - Blocking of bacterial metabolites:** determine the role of metabolites in barrier function by blocking receptors

Novel internalin-like ZO1 regulator protein (IZOR) promotes protective barrier function effects in Sp. 1

Screen of 12 strains of Species 1 on barrier function

Strain	Sp.1	Sp.1a	Sp.1b
AUC (hrs)	18	18	18

Strain	Sp.1c	Sp.1d	Sp.1e	Sp.1f	Sp.1g	Sp.1h	Sp.1i	Sp.1j	Sp.1k	Sp.1l
Normalised Resistance	-0.1	-0.1	-0.2	-0.2	-0.1	0.3	-0.1	0.1	0.1	0.1

Comparative genomics on Species 1 strains

IZOR gene

Regulation of barrier function by IZOR-expressing *E. coli*

Condition	ST	FP	Control <i>E. coli</i>	IZOR <i>E. coli</i>
Normalised resistance	-1.2	0.2	-0.2	0.3

Regulation of tight junction expression by IZOR

Condition	Control <i>E. coli</i>	IZOR <i>E. coli</i>	Sp. 1
Fold change	1.0	1.6	1.6

12 strains of Species 1 were tested in the barrier repair assay in 2 batches. 4 beneficial (**healer**) strains and 10 non-beneficial (**neutral**) strains were identified

When comparing the genomes of **healer** strains and **neutral** strains only 1 gene was differentially present, internalin-like ZO1 regulator protein (IZOR). IZOR is a novel protein only present in a subset of species 1

Transduction of *E. coli* with IZOR gene confers beneficial outcome in barrier function assay indicating it is sufficient for protective effect

IZOR-expressing *E. coli* and species 1 regulate tight junction proteins in Caco-2 cells. Both strongly induce ZO1, which is key for barrier integrity

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Conclusions

- The 8 MB310 species, identified as being associated with clinical benefit, interact with the host in different ways to oppose key UC disease pathologies
- 3 of the MB310 species improve gut epithelial barrier integrity, particularly promoting recovery of epithelium post inflammatory damage.
- These species act through 3 distinct mechanisms of action:
 - Species 1 expresses a novel protein, IZOR, which regulates the expression of tight junction proteins and mediates the barrier protective effects of species 1.
 - Species 5 and species 8 promote recovery of epithelium post pro-inflammatory damage by secretion of bacterial metabolites; polyamines and nicotinic acid, respectively
- A combination of anti-inflammatory and barrier repair address key UC disease pathologies, offering therapeutic potential for a broad spectrum of UC patients
- COMPOSER-1 is an ongoing Phase 1b trial testing MB310 in UC patients