



# Restoration of gut microbiota diversity with oral pooled fecal microbiotherapy in acute myeloid leukemia patients after intensive chemotherapy: the phase 1b CIMON trial

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## INTRODUCTION

- Acute myeloid leukemia (AML) treatment combines intensive chemotherapy (IC) with broad-spectrum antibiotics (ATB) that induces a strong gut microbiota dysbiosis, promoting pathological conditions and increasing incidence of complications.
- Restoration of the full gut microbiota ecosystem is a promising therapeutic tool to improve clinical outcomes in patients with acute myeloid leukemia (AML) receiving IC and ATB.
- Here we report the tolerability, safety and efficacy of MaaT033, an oral, delayed-release capsule containing lyophilized pooled full ecosystem fecal microbiota, in restoring the gut microbiota in 21 AML patients having undergone IC and ATB: results from the Phase Ib CIMON trial (NCT04150393).

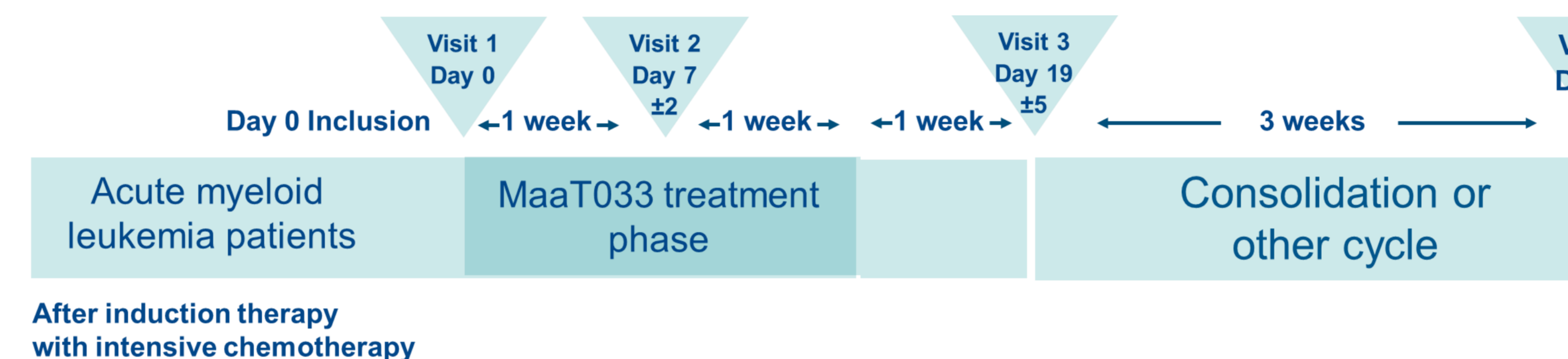
## Main inclusion criteria

- Age  $\geq$  18 years
- Patients diagnosed with **AML or high-risk myelodysplastic syndrome**, receiving intensive chemotherapy
- Patients healthy enough to likely receive **their consolidation or second cycle of chemotherapy after induction chemotherapy**
- Patients healthy enough to likely receive HSCT
- Patient recovered from neutropenia

## Main exclusion criteria

- Active uncontrolled infection** according to the attending physician
- Any gastro-intestinal **bleeding in the past 3 months**
- [...]

## STUDY FLOW CHART



## PATIENTS' CHARACTERISTICS

|  |                                   | Treated patients (N=21), n(%)      |          |
|--|-----------------------------------|------------------------------------|----------|
| Gender, n (%)  | Male                              | 16 (76%)                           |          |
|  | Female                            | 5 (24%)                            |          |
| Age at inclusion (years)   | Median [range]                    | 62 [22-77]                         |          |
| Leukemia risk category (ELN 2022), n (%)                                   | Favorable                         | 7 (33%)                            |          |
|  | Intermediate                      | 6 (29%)                            |          |
|  | Adverse                           | 8 (38%)                            |          |
| Induction chemotherapy, n (%)  | Cytarabine + anthracycline        | 20 (95%)                           |          |
|  | Azacitidine                       | 1 (5%)                             |          |
|  | Consolidation chemotherapy, n (%) | Cytarabine alone or in combination | 17 (81%) |
|  |                                   | Daunorubicin                       | 1 (5%)   |
|  |                                   | Azacitidine + Venetoclax           | 1 (5%)   |
| None   | 1 (5%)                            |                                    |          |
| Unknown  | 1 (5%)                            |                                    |          |
| Patients who received antibiotics during induction chemotherapy, n (%)     | Prophylactic antibiotics          | 10 (48%)                           |          |
|  | Therapeutic antibiotics           | 19 (90%)                           |          |
|  | No systemic antibiotics           | 2 (5%)                             |          |
| Patients who received antibiotics during consolidation chemotherapy, n (%) | Prophylactic antibiotics          | 3 (14%)                            |          |
|  | Therapeutic antibiotics           | 10 (48%)                           |          |
|  | No systemic antibiotics           | 7 (33%)                            |          |
|  | Unknown                           | 1 (5%)                             |          |

## METHODS AND OBJECTIVES



### CIMON study design

Phase I, open-label, single-arm  
6 investigational sites in France



### DOSE ESCALATION DESIGN

5 cohorts : dose escalation of MaaT033

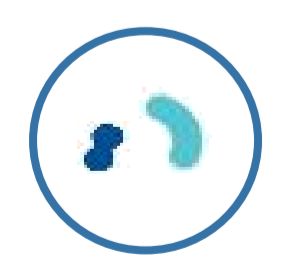
- Cohort 5 (n=6) Not performed, due to sufficient data from Cohort 1-4
- Cohort 4 (n=6) 1 capsule / day, 2 weeks
- Cohort 3 (n=6) 1 capsule / day, 1 week
- Cohort 2 (n=6) 1 capsule / day, 1 week
- Cohort 1 (n=3) 2 capsules / week, twice

Dose escalation



### STUDY OBJECTIVES

- ✓ MaaT033 tolerability
- ✓ Dose regimen evaluation (based on safety and MaaT033 activity / engraftment)
- ✓ Patient compliance



### GUT MICROBIOTA CHARACTERIZATION

- ✓ 16S sequencing



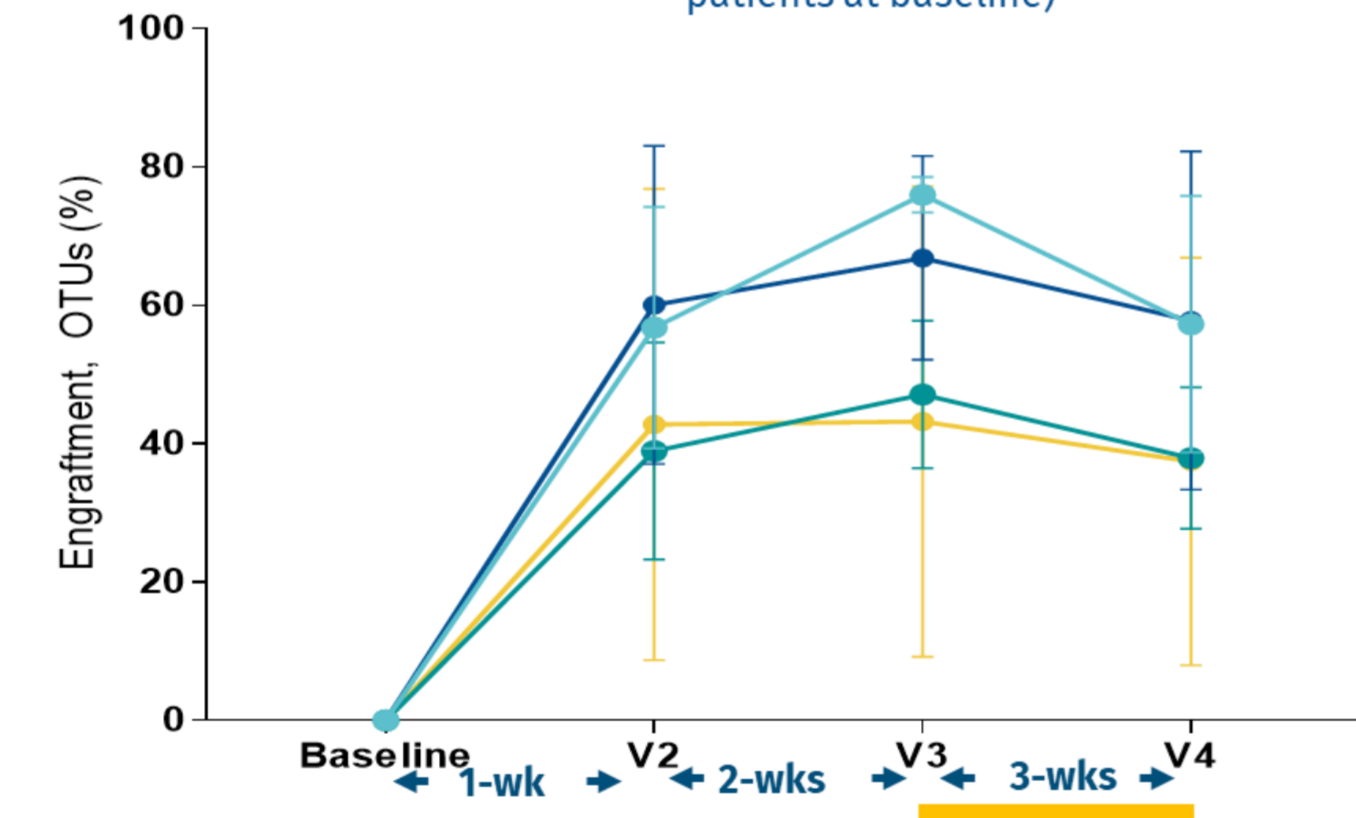
### MaaT033 CHARACTERISTICS

Oral pooled fecal microbiota

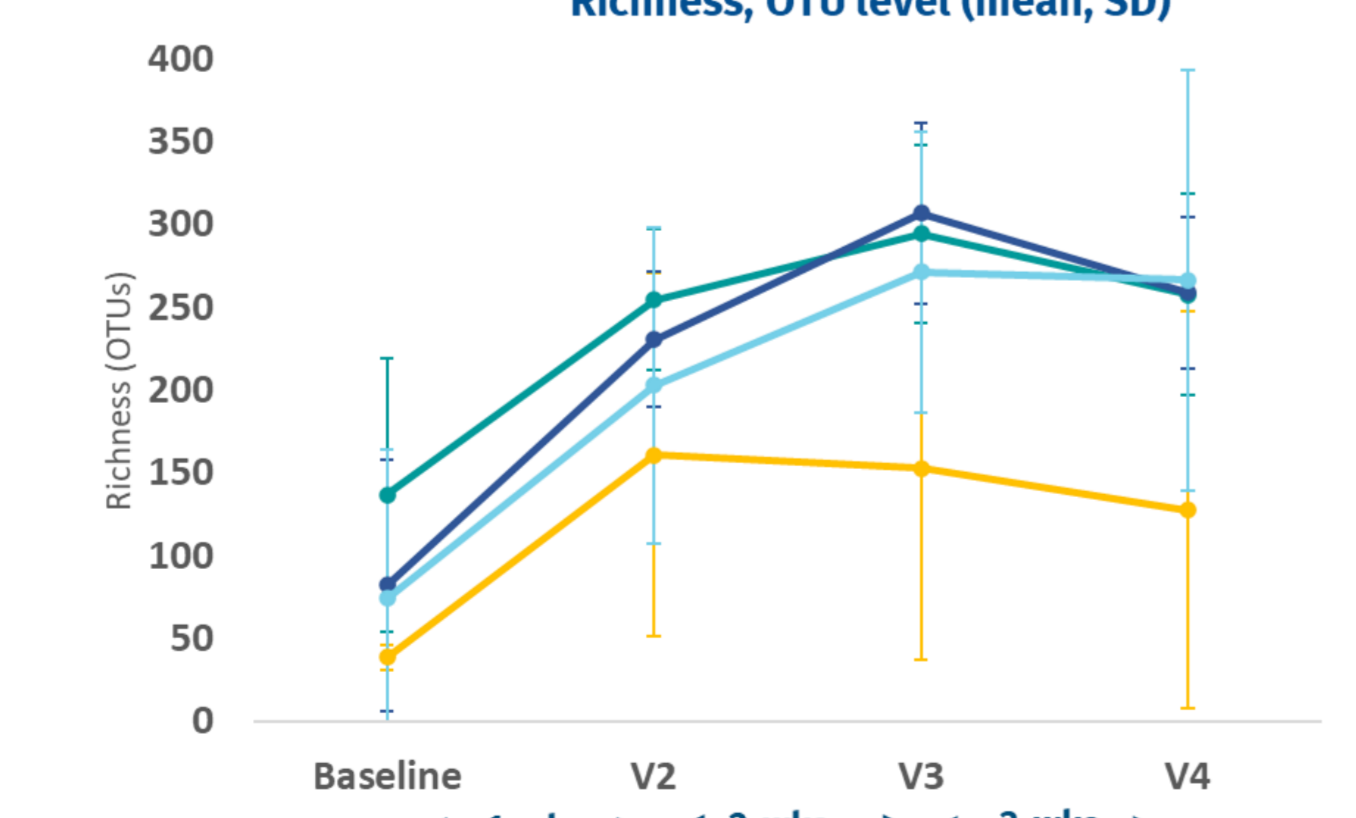
- ✓ Strict screening process for healthy donors (> 40 tested parameters)
- ✓ High-richness
- ✓ High-diversity
- ✓ Full ecosystem
- ✓ Ileo-caecal delivery
- ✓ > 10<sup>9</sup> viable bacteria/capsule

## RESULTS: GUT MICROBIOTA AND HOST PARAMETERS ANALYSIS

MaaT033 engraftment (mean, SD)  
(specific to MaaT033 i.e., exclude shared OTUs between MaaT033 and patients at baseline)



Richness, OTU level (mean, SD)



OTU: operational taxonomic unit; V: visit; V1 (baseline): after induction chemotherapy and neutropenia recovery, before MaaT033 start; V2: 1 week after MaaT033 start; V3: before consolidation cycle; V4: at the end of consolidation cycle; C: Cohort; SCFAs: short chain fatty acids; wk: week

- MaaT033 induces an increased microbiota richness at OTUs level

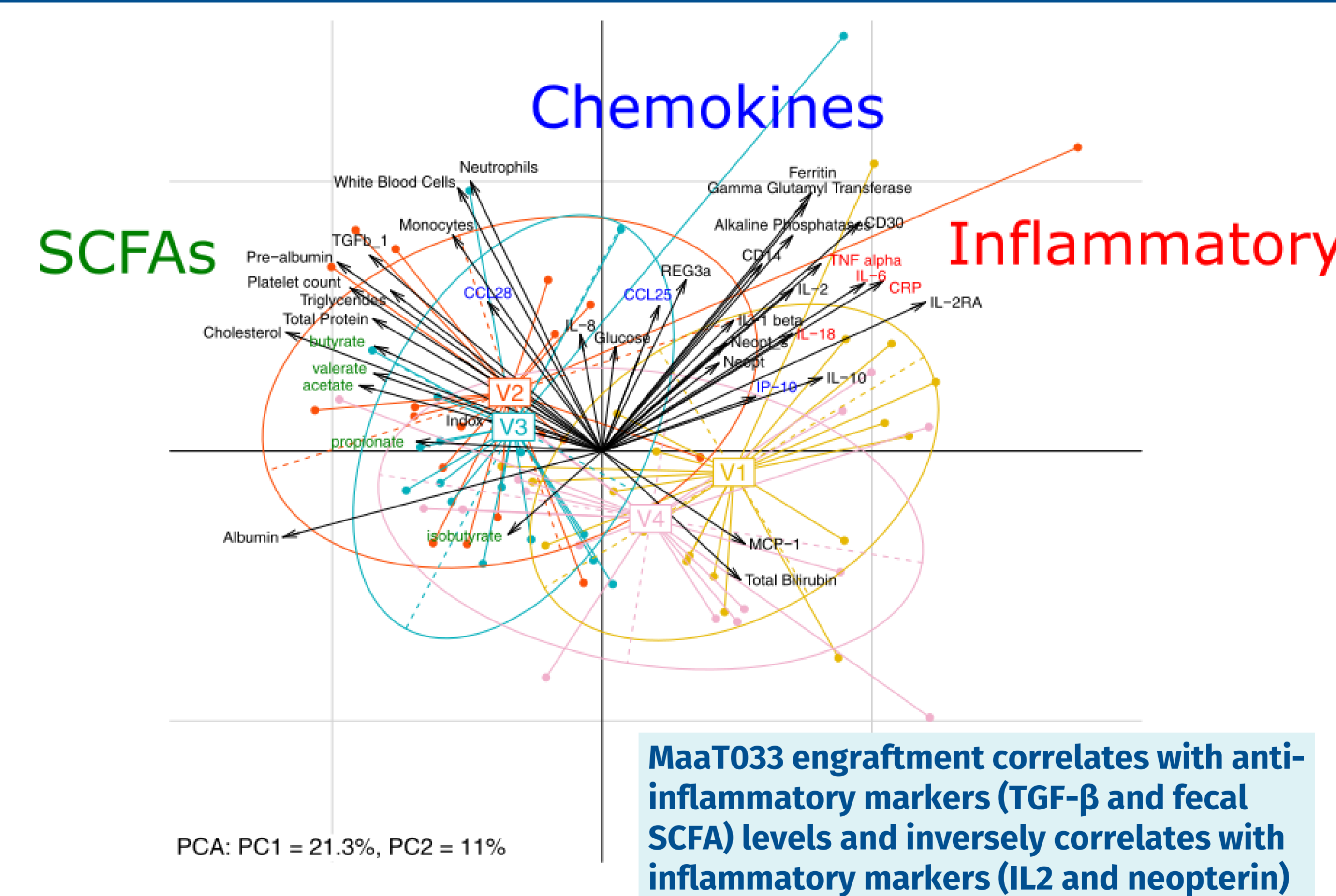
- MaaT033 displays a strong and persistent bacterial engraftment, higher in cohort 3 and cohort 4 (3 capsules per day) compared to cohorts 1 and 2 (1 capsule per week / 1 capsule per day).

- MaaT033 bacterial engraftment is inversely correlated with patients' baseline microbiota richness

(up) Microbiota richness, defined as number of OTUs found in patients' gut microbiota, described for each cohort

(down) Engraftment of MaaT033 in each cohort

Engraftment = ratio of OTUs that were not present in patient at baseline but were present in MaaT033 and were found in patient after treatment. Redundant OTUs found in patient and MaaT033 are not taken into account.



Principal Component Analysis performed with the host parameters dataset.

Individual samples are represented by points, colored and linked according to the corresponding visit. The arrows depict the variables, here the host parameters. The longer they are the more they drive the samples projections, the closer they are, the more they vary together. Specific parameter groups are highlighted in color (SCFAs, Chemokine, and inflammatory parameters). For a better visualization, variables with negligible impact on sample projections (with small arrows) were removed from the figure after computation.

MaaT033 engraftment correlates with anti-inflammatory markers (TGF- $\beta$  and fecal SCFA) levels and inversely correlates with inflammatory markers (IL2 and neopterin)

4 serious adverse events (SAEs) reported in 4 patients. Only one (in bold) was considered as possibly related by the investigator.

| Event                       | Cohort 1 (N=3) n (%) | Cohort 2 (N=6) n (%) | Cohort 3 (N=6) n (%) | Cohort 4 (N=6) n (%) | Total (N=21) n (%) |
|-----------------------------|----------------------|----------------------|----------------------|----------------------|--------------------|
| Febrile neutropenia         | 0                    | 1 (17%)              | 0                    | 0                    | 1 (5%)             |
| Neutropenic colitis         | 0                    | 0                    | 1 (17%)              | 0                    | 1 (5%)             |
| <b>Diarrhoea infectious</b> | 0                    | 0                    | 0                    | 1 (17%)              | 1 (5%)             |
| Hyperkalaemia               | 0                    | 1 (17%)              | 0                    | 0                    | 1 (5%)             |

The SAE infectious diarrhea with enteropathogenic *E. coli* (EPEC) started 3 days after MaaT033 treatment initiation.

- ✓ EPEC strain not found in MaaT033 batch (genome sequencing) nor in patient's feces before MaaT033 administration
- ✓ Absence of pillbox contamination verified by PCR
- => Relationship to MaaT033 was unlikely, although it could not be formally excluded.

Six infectious events reported :

- 1 during MaaT033 treatment phase (SAE described above)
- 5 after MaaT033 treatment and during the consolidation cycle : cellulitis, Escherichia sepsis, paronychia, Pseudomonas infection, skin infection, none of them being reported as serious nor potentially related to MaaT033

MaaT033 treatment is safe :

After IC in immunocompromised AML patients : good tolerance, minimal toxicity after treatments and post-neutropenia

## CONCLUSIONS

- MaaT033 appears to be **safe and effective for gut microbiota restoration** in AML patients receiving IC and ATB.
- 3 MaaT033 capsules per day for 1 week** induce an **increase in microbiota richness** and an **effective and persistent MaaT033 bacterial engraftment** in AML patients
- A Phase IIb trial is planned to start in 2023 to evaluate MaaT033 as an adjunctive and maintenance treatment in patients with hematological malignancies receiving allogeneic hematopoietic stem cell transplantation.