



UNIVERSITÀ
DEGLI STUDI
DI TRIESTE

LIVER BACTERIAL COLONIZATION IN PATIENTS WITH OBESITY AND GUT DYSBIOSIS

*Lucia Paiano, Giuseppina Campisciano, Biagio Casagranda,
Manola Comar, Nicolò de Manzini, Silvia Palmisano*

**University of Trieste- Surgical Clinic Unit,
Cattinara Hospital, ASUGI**



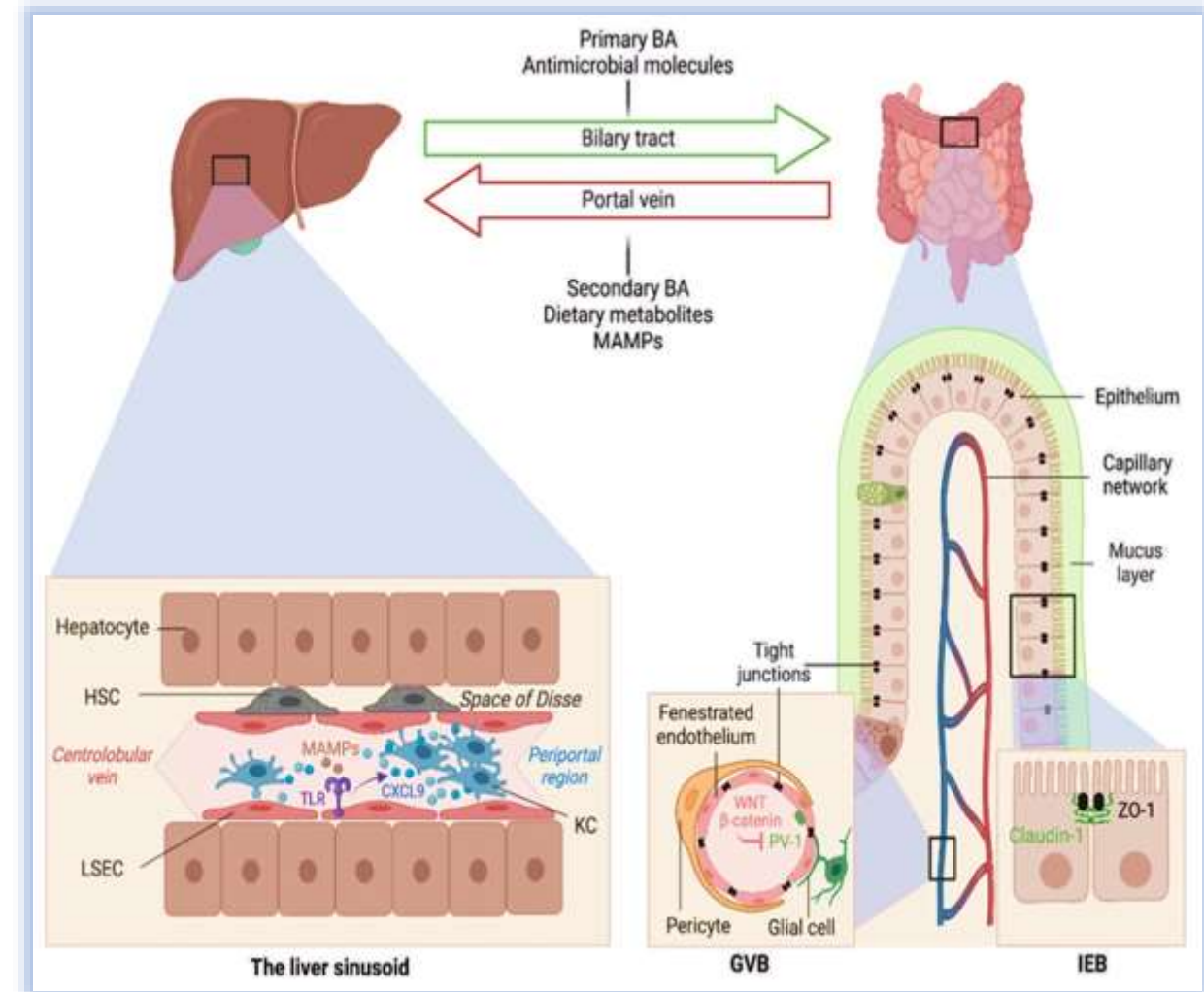
Background:

Link between

- Gut microbiota,
- Obesity
- Liver inflammation



FOCUS OF RESEARCH



Obesity → chronic dysbiosis gut microbiota

Background:

In the inflammatory status

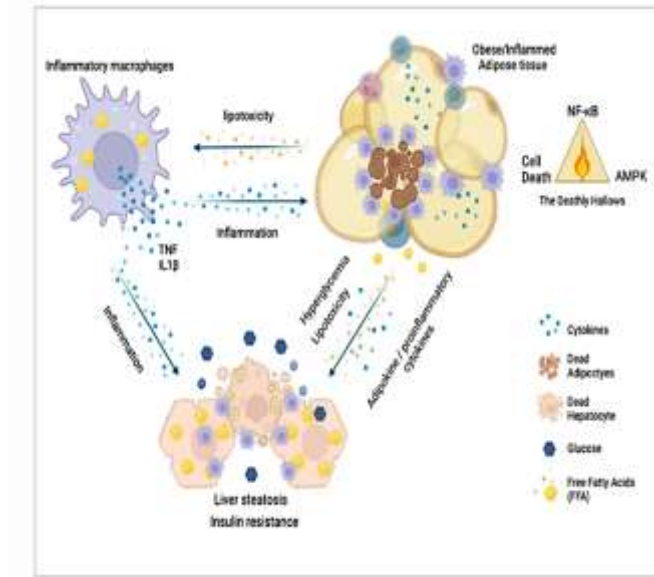
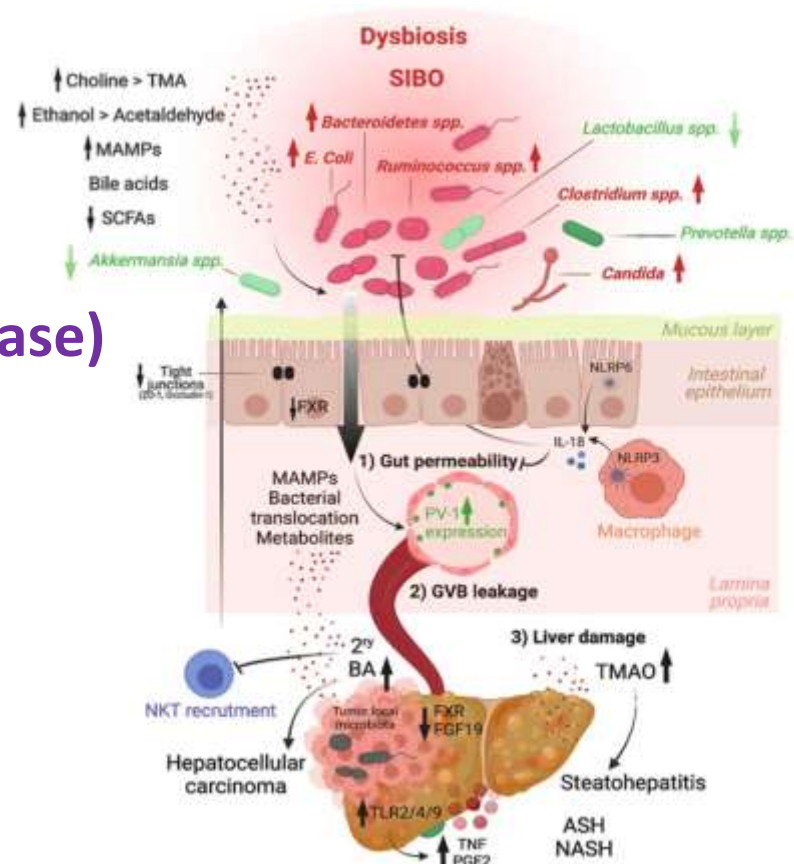
(Obesity → chronic inflammatory disease)



Gut permeability



Translocation of MAMPs and DAMPs
(microbial and danger associated molecular pattern)
from the gut to the liver



Cell death and inflammation during obesity: "Know my methods, WAT(son)"
[Ximena Hildebrandt et al Nature Sep 2022](#)



Host-Microbiota Interactions in Liver Inflammation and Cancer
[Julie Giraud et al Cancer 2021](#)



NO studies that demonstrate the presence of the bacterial genome in obese patients liver biopsy.

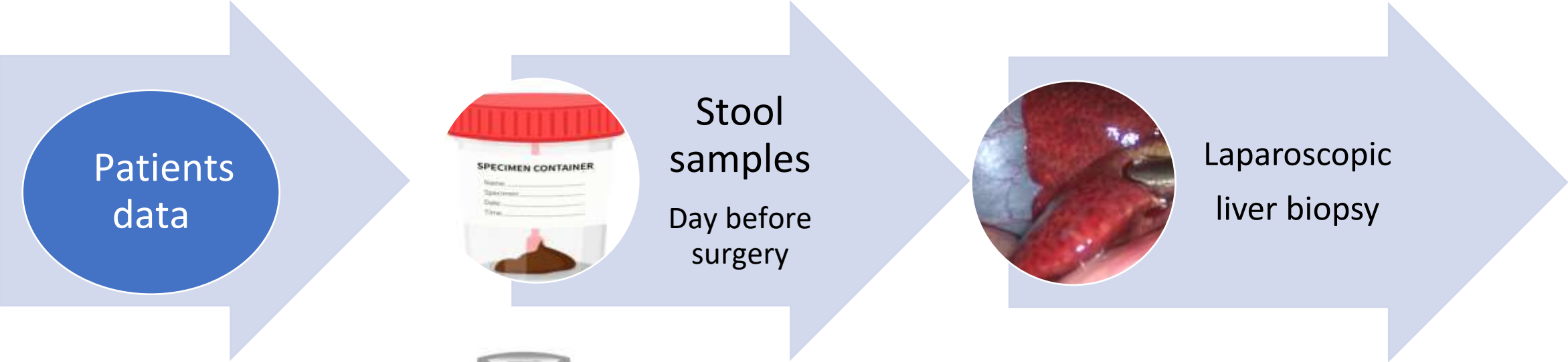
AIMs OF THE STUDY:

- ❑ Search for BACTERIAL GENOMA IN LIVER BIOPSY (neither MAMPs nor DAMPs)
- ❑ Compare the bacterial genome founded in the liver with gut microbiota composition in fecal samples (gut-liver axis)



Methods:

Prospective study → Enrollement patients undergoing bariatric surgery consequently .
Signature of informed consent, collection anamnestic/anthropometric information.



V3-16S rRNA region sequencing
Ion Torrent new generation sequencing platform

Results:

n= 23 patients
Stools samples and liver biopsy



V3-16S rRNA region sequencing



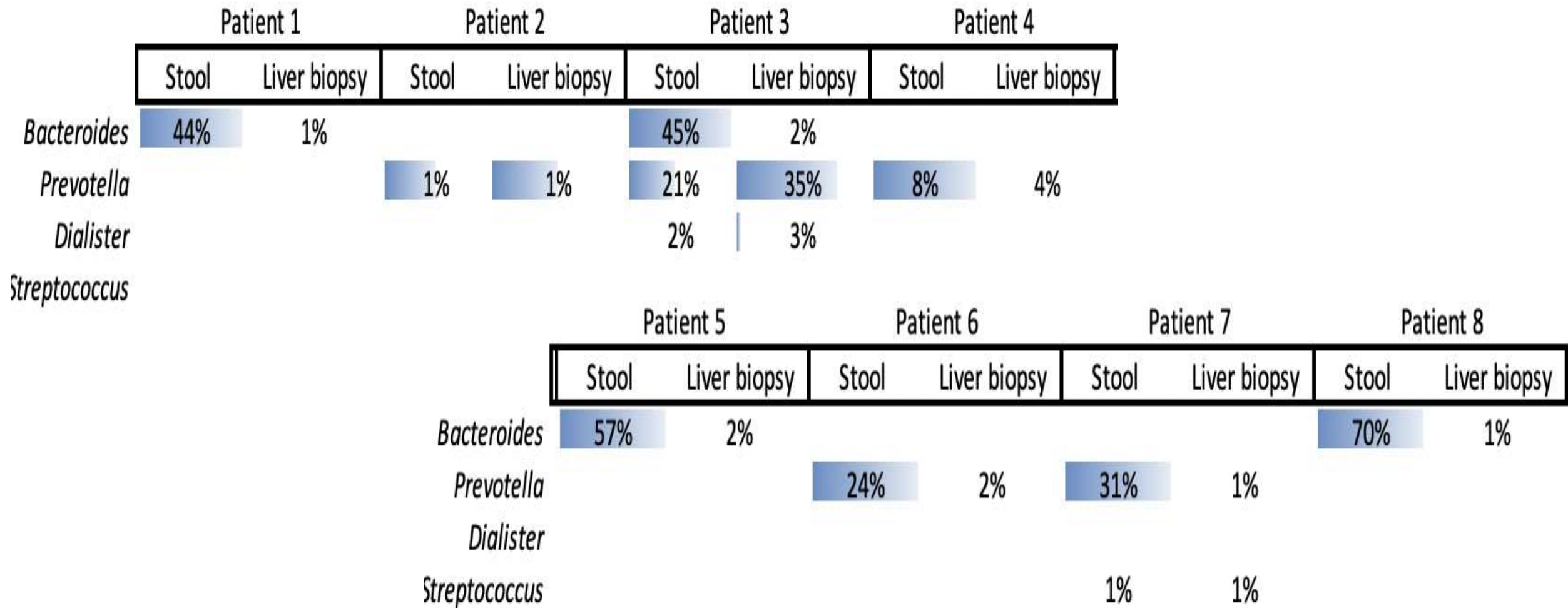
n=8 patients (34,7%)
The same bacterial genoma in stools samples and in liver biopsy

n=15 patients (65,3%)
Baterial genoma only in stools samples



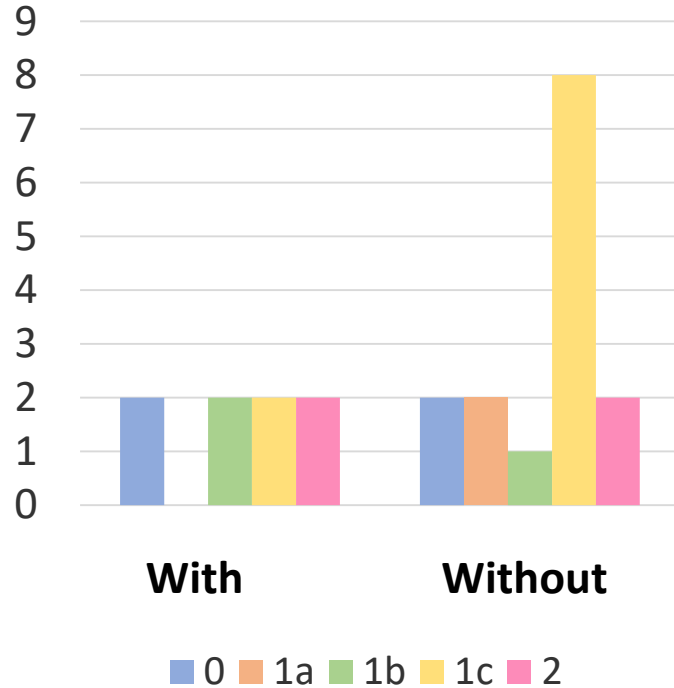
Prevotella (62,5%), Bacterioides (50%), Dalister (12,5%), Streptococcus (12,5%)

Results:

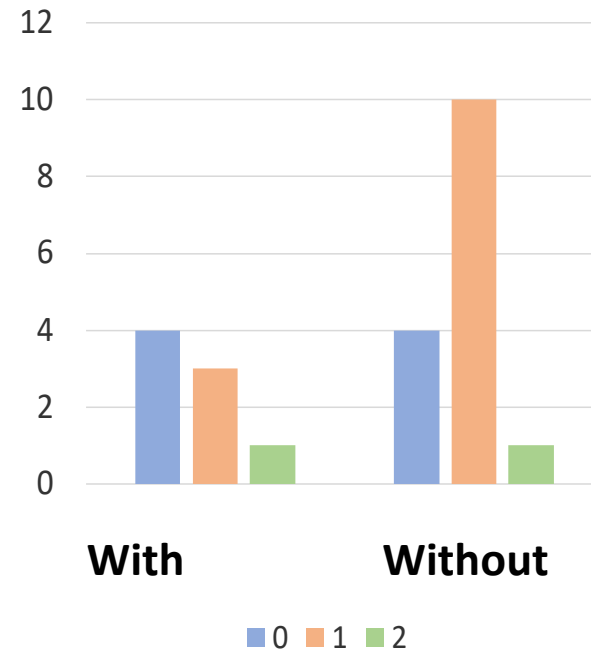


Results:

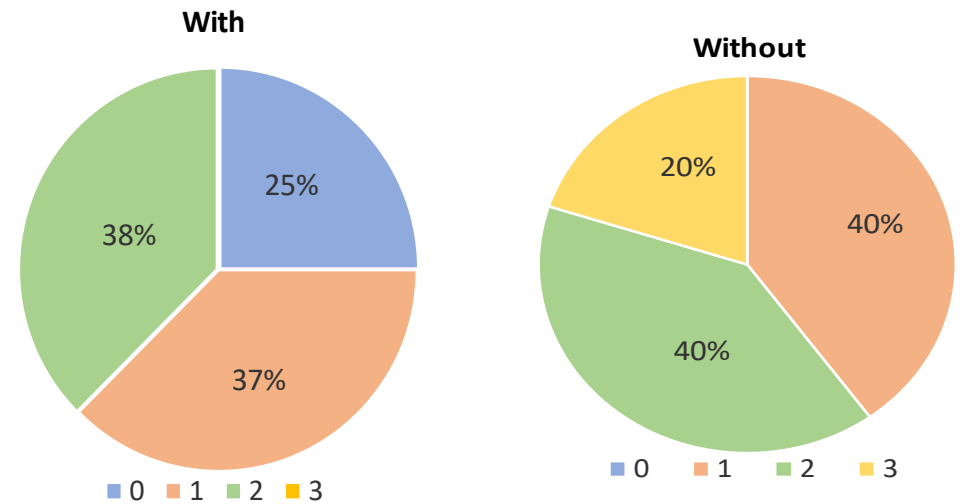
Kleiner fibrosis



Kleiner lobular inflammation



Brunt steatoepatitis



$p = 0.2$

Results:

Complication rate
according to
Dindo Clavien
90 days

	Obese w liver colonization	Obese w/o liver colonization	
0	12 (80%)	6 (75%)	<i>p</i> = 0.9
1	1 (6.7%)	1 (12%)	
2	2 (13%)	1 (12%)	

% Body Weight Lost at 1 year
(Exces poids)

83,6 %

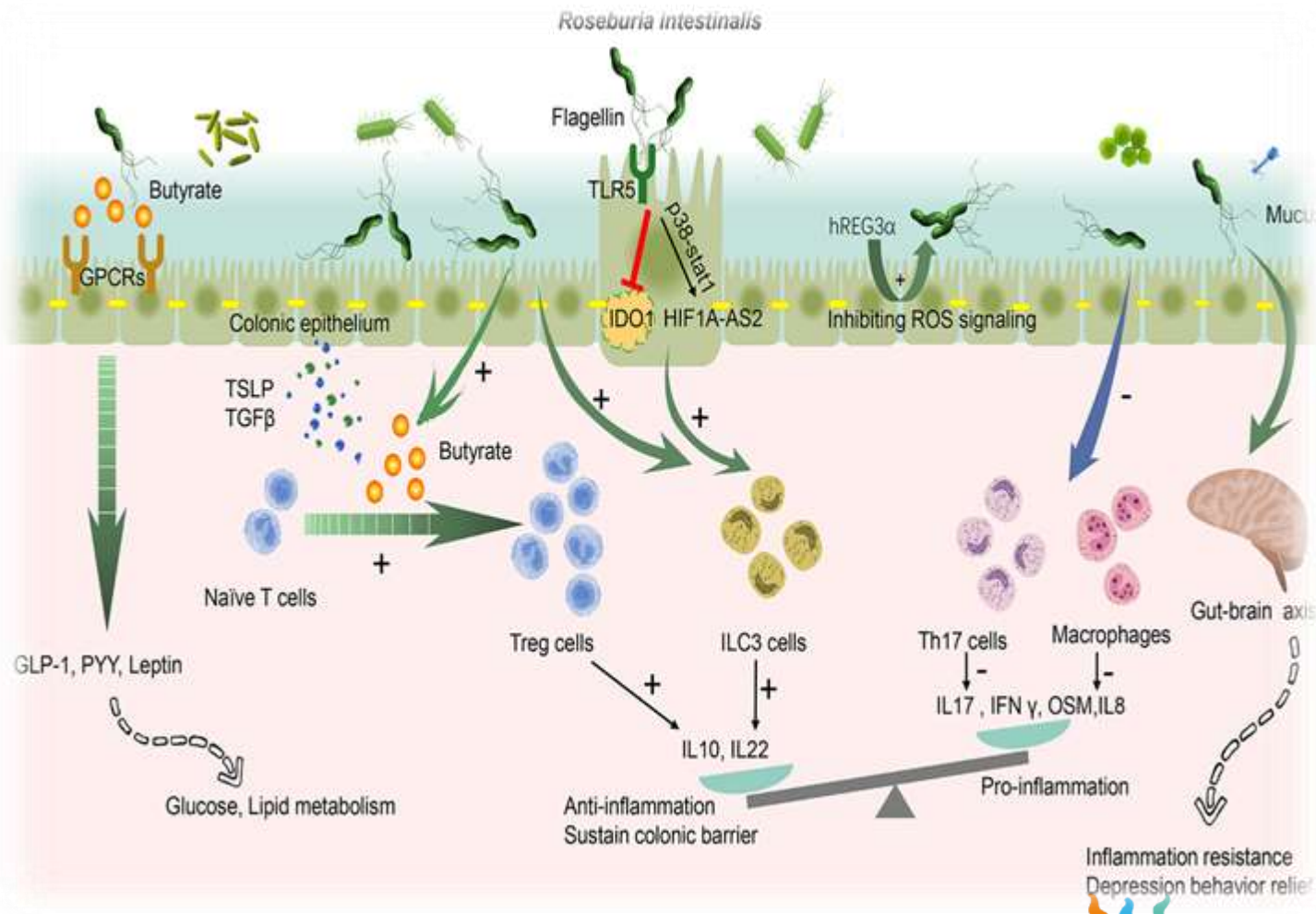
74,4 %

***p* = 0.4**

Microbiota analysis (median) :

Species	Obese w/o liver colonization	Obese w liver colonization	P value
<i>Bacteroides coprocola</i>	2%	0%	0.810
<i>Bacteroides dorei</i>	1%	0%	0.070
<i>Bacteroides faecis</i>	1%	1%	0.441
<i>Bacteroides massiliensis</i>	2%	2%	0.246
<i>Bacteroides thetaiotaomicron</i>	1%	0%	0.515
<i>Bacteroides uniformis</i>	3%	6%	0.582
<i>Bacteroides vulgatus</i>	6%	4%	0.496
<i>Barnesiella intestinihominis</i>	0%	1%	0.718
<i>Blautia wexlerae</i>	1%	1%	0.794
<i>Clostridium bartlettii</i>	1%	1%	0.347
<i>Clostridium glycyrrhizinilyticum</i>	1%	0%	0.818
<i>Clostridium symbiosum</i>	2%	1%	0.219
<i>Faecalibacterium prausnitzii</i>	4%	2%	0.496
<i>Fusicatenibacter saccharivorans</i>	1%	2%	0.496
<i>Gemmiger formicilis</i>	0%	1%	0.258
<i>Parabacteroides distasonis</i>	1%	0%	0.631
<i>Parabacteroides merdae</i>	1%	1%	0.794
<i>Roseburia intestinalis</i>	3%	1%	0.034
<i>Suttereilla waadsworthensis</i>	1%	0%	0.896

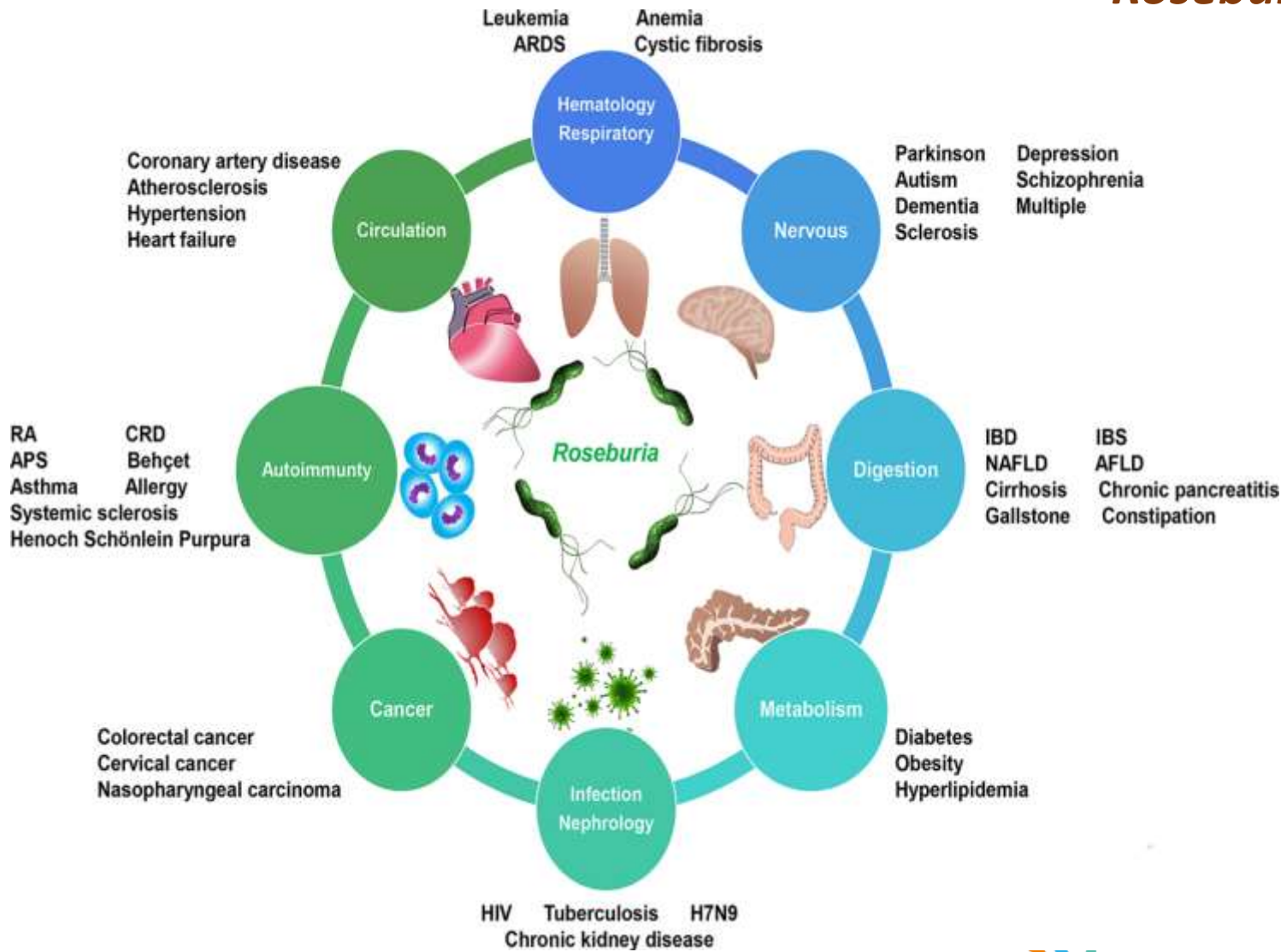
Roseburia intestinalis modulation in the gut tract



- Production of butyrate
- Stimulation excretion cytokines
- Promotion differentiation and regulation T cells
- Activation ILC3
- Suppression TH17 and macrophages
 - Its flagellin displays an anti-inflammation effect through TLR5.

Roseburia intestinalis: A Beneficial Gut Organism From the Discoveries in Genus and Species Kai Nie et al

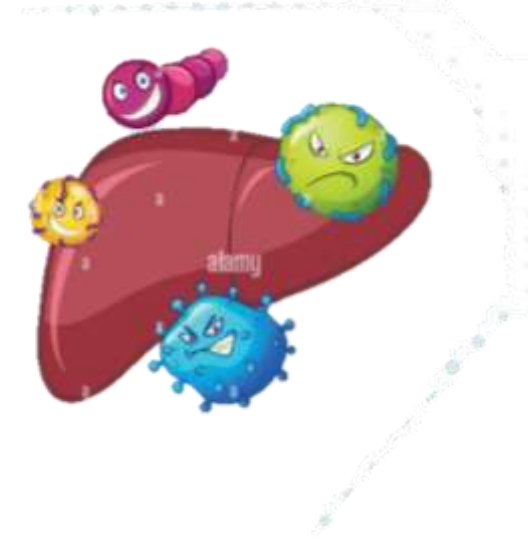
Roseburia dysbiosis-associated diseases exist in different systems



IBD (inflammatory bowel diseases)
 IBS (irritable bowel syndrome)
 NAFLD
 AFLD,
 CRD (chronic rheumatoid disease)
 APS(antiphospholipid syndrom)
 etc...

Conclusions:

- **First** study reporting the presence of bacterial genome in a liver biopsy on bariatric patients.
- In 8 patients, we found ***same bacterial genoma in stools samples and in liver biopsy*** → *bacterial traslocation from gut or vascular barrier to the liver*
- In our population, the ***Roseburia intestinalis* dysbiosis** was associated with the presence of bacterial genome in the liver, probably related with a greater permeability of the gut and vascular barriers.



In accordance with «EACCME criteria for the Accreditation of Live Educational Events»:

I have no potential conflict of interest to report





Thank you for your attention!!!