

Gut microbiota composition and arterial hypertension improvement post-Roux-en-Y Gastric Bypass: GATEWAY Trial sub-analysis after 5 years

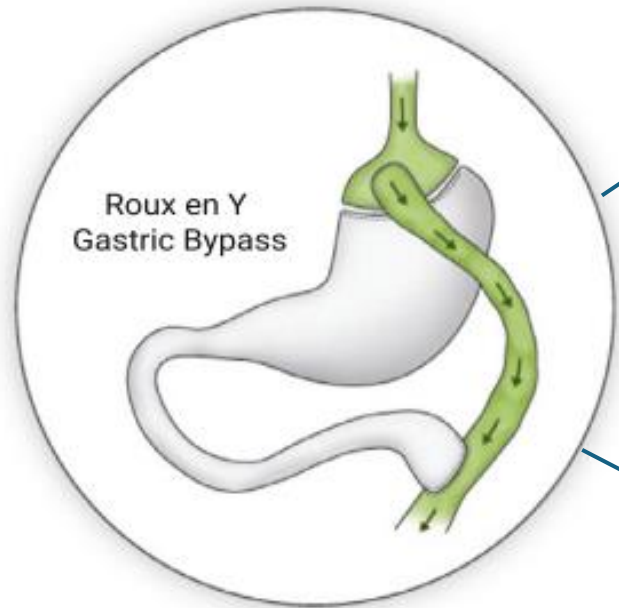
Danielle C. Fonseca, **Carlos Schiavon**, Aline Marcadenti, Rachel H.V. Machado, Lucas Tramuja, Ângela C. Bersch-Ferreira



Disclosures

- Research Grant:
 - Ethincon, Inc;
 - hcor, Associação Beneficente Síria;
 - **BiomeHub**

Bariatric and metabolic surgery – gut microbiota (GM)



↑ Proteobacteria
↑ Gemella
↑ Streptococcus

Frequently related to human diseases

↑ Veillonella
↑ Akkermansia

Frequently related to human health

↓ Blautia
↓ Bifidobacterium

Frequently related to human health

↓ SCFAs (Short-chain fatty acid)
↓ BCAAs (Branched chain amino acids)

Frequently related to human health

↑ BCFAs (Branched chain fatty acids)
↑ Tryptophan-derived metabolites
↑ TMAO (Trimethylamine N-oxide)

Frequently related to human diseases

↑ Secondary bile acids **Frequently related to human health**

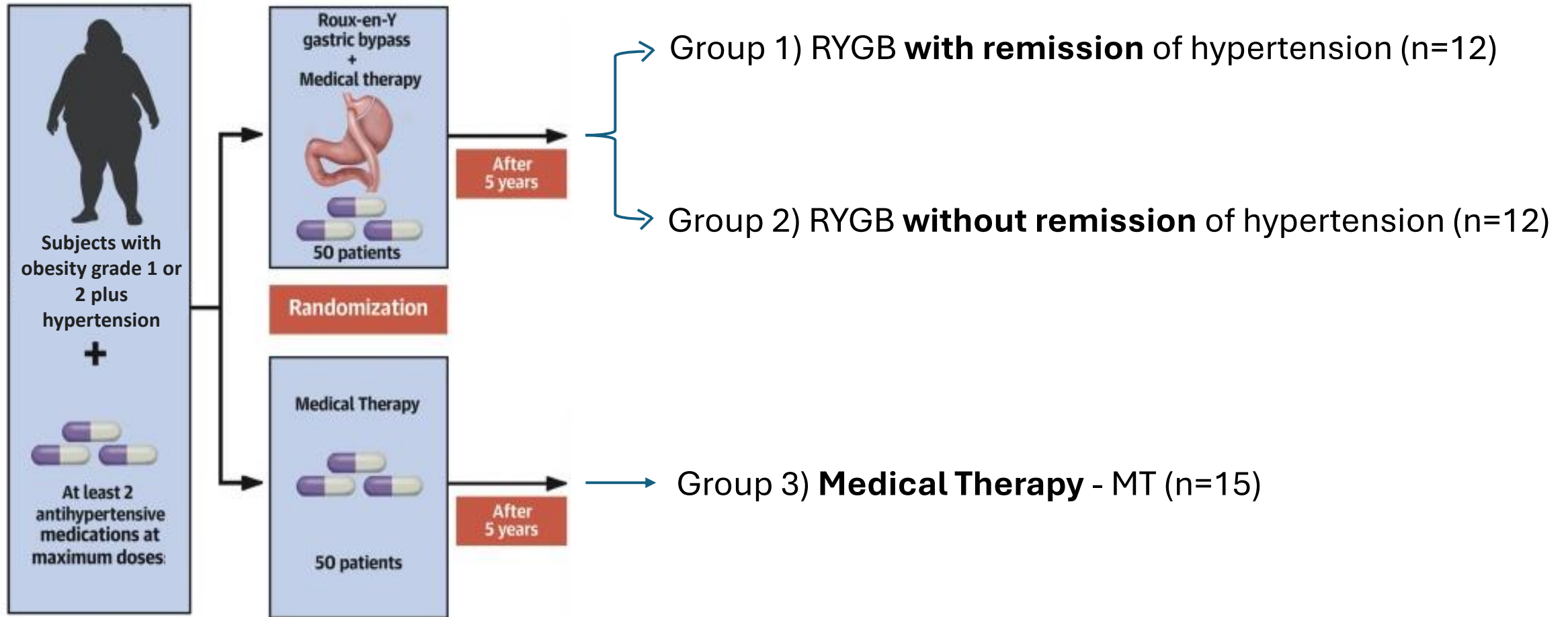
Aim



Our aim was to describe and evaluate the GM composition according to the type of response to RYGB compared to Medical Therapy after five years or more years of follow-up at the GATEWAY Trial.

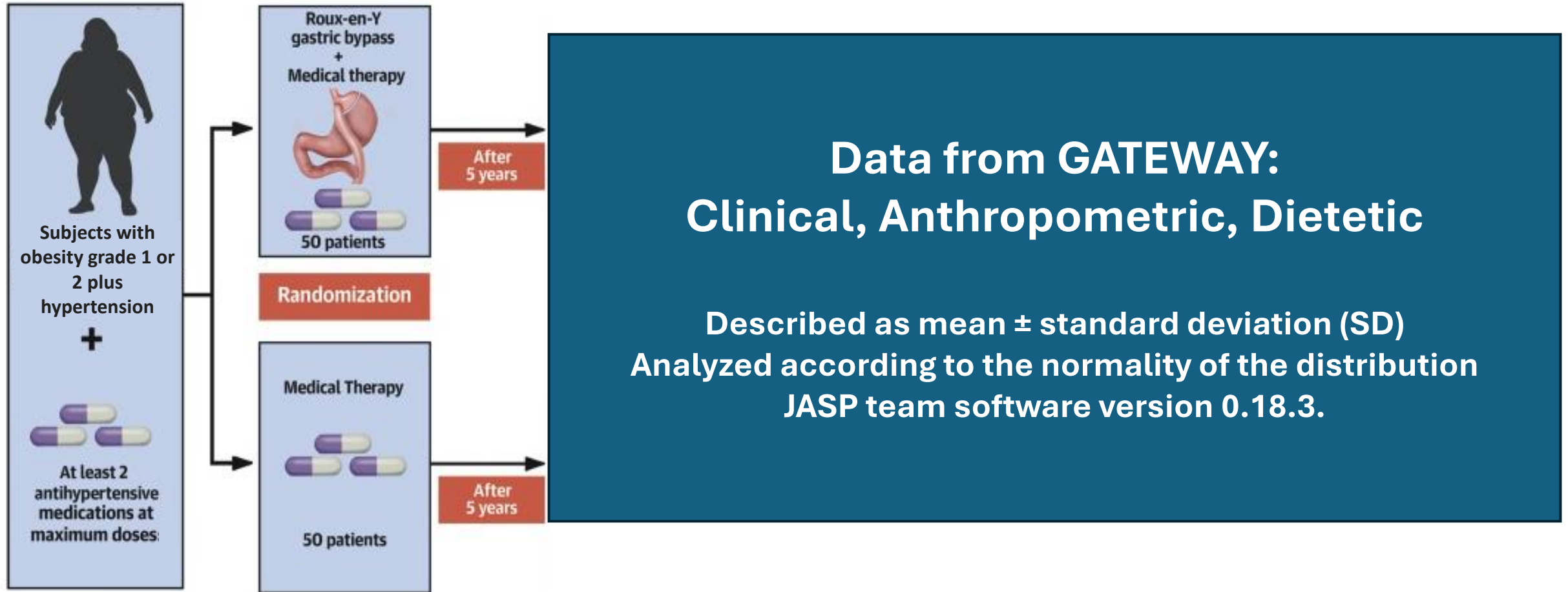
Methods

This is a sub-analysis of the GATEWAY RCT (NCT01784848) that included individuals with obesity and hypertension after at least five years follow-up.



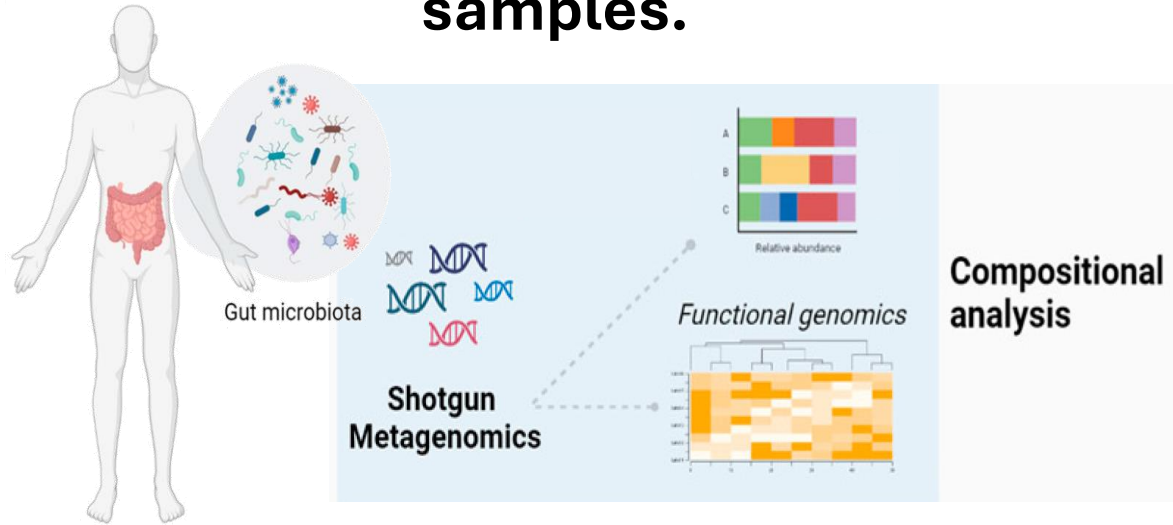
Methods

This is a sub-analysis of the GATEWAY RCT (NCT01784848) that included individuals with obesity and hypertension after at least five years follow-up (cross-sectional analysis).



Methods

GM was assessed by shotgun metagenomic sequencing of fecal samples.



Descriptives

- Relative abundance (GM profile):
Percentual of Phyla and Genera

Descriptives:

- **Richness:** the number of species present in a system
- **Alfa-diversity (Simpson and Shannon):** the ecological diversity of a single sample, taking into account the number of different bacteria and their relative abundances (distribution)



Considered measures of gut microbiota health

Analyses were performed:

- LinDA in MicrobiomeStat R package version 1.1
- Correlation among GM and clinical data in JASP team software version 0.18.3.

Results

Population Characteristics After 5-years follow-up:

Variable	Group1 (n=12)	Group2 (n=12)	Group3 (n=15)	P-value
Age (years)	47.7 ± 8.3	55.8 ± 9.4	55.3 ± 8.2	0.044
BMI (kg/m ²)	27.6 ± 2.0	28.7 ± 5.2	37.4 ± 4.2	p<0.001
SBP (mmHg)	119.4 ± 14.3	130.7 ± 12.4	127.7 ± 17.3	0.544
DBP (mmHg)	78.6 ± 12.0	83.4 ± 11.6	80.9 ± 8.3	0.173

Anova

Results

GM profile:

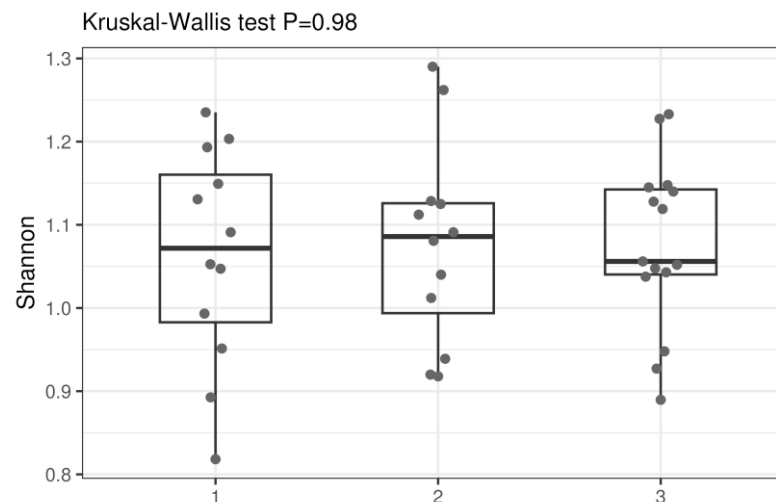
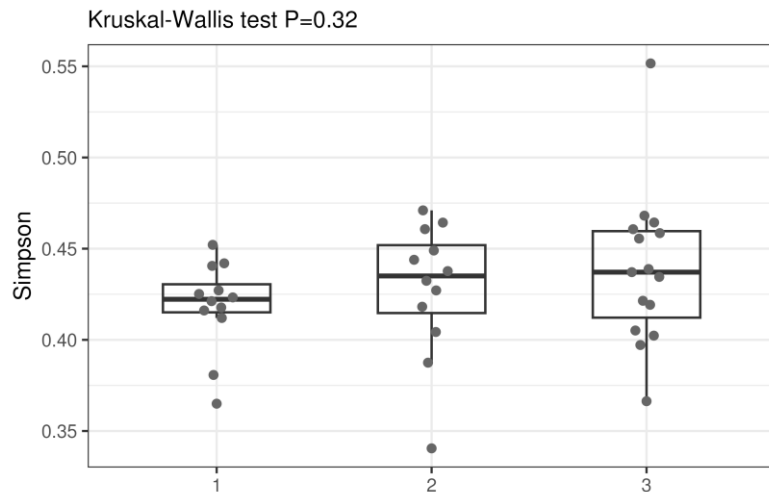
- **Phyla (p>0.05): Similar among groups**

Firmicutes, Bacteroidetes, Proteobacteria and Actinobacteria → The dominant gut microbial of human

- **Top five genera (p>0.05): Similar among groups**

Prevotella (24.5%); Faecalibacterium (6.2%); Blautia (4.3%); Escherichia (3.8%); Phocaeicola (3.8%)

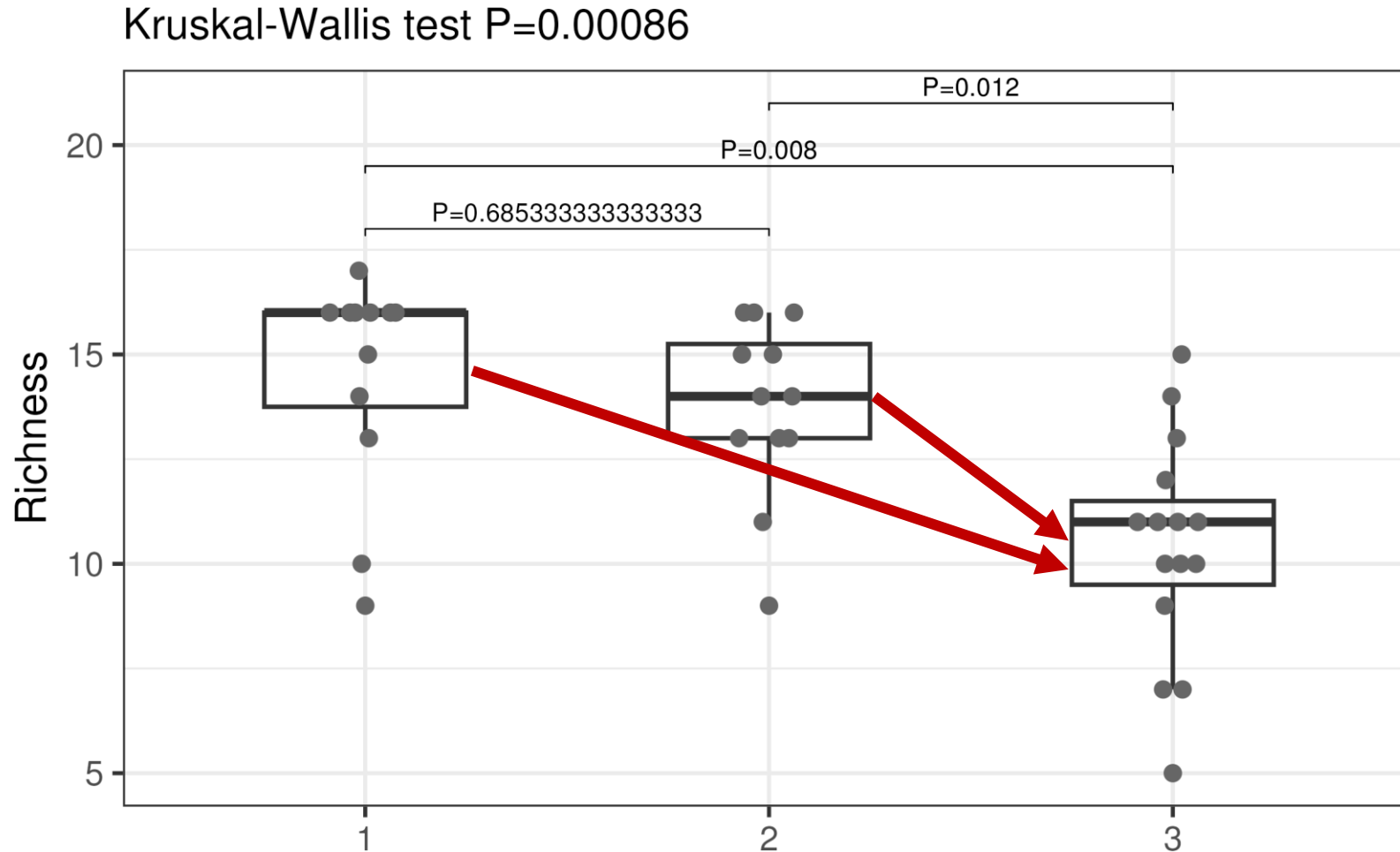
- **Alfa-diversity measures (p>0.05): Similar among groups**



Regarding the
compositional aspect,
no differences were
identified between the
GM of the groups

Results

Microbial richness was notably different between BS groups (1 and 2) comparing MT



Literature suggests that: low richness of GM is characterized by more adiposity, insulin resistance and a more pronounced inflammatory phenotype when compared with high GM richness individuals.

Le Chatelier E, Nielsen T, Qin J, et al. *Nature*. 2013 Aug 29;500(7464):541-6.
Balmant BD, Fonseca DC, Rocha IM, et al. *Nutrients*. 2023 Oct 5;15(19):4261.

Pairwise comparisons used Wilcoxon rank sum tests with Holm correction for multiple comparisons.

Conclusion

- This comprehensive analysis unveiled no disparities in the taxonomic composition of the GM when comparing groups.
- Nevertheless, it did identify higher richness in the GM of participants undergoing RYGB compared to those who remained on drug treatment throughout the study's follow-up period.
- The increased GM richness has previously been observed after short-term bariatric surgery. Studies with follow-up of more than five years are scarce, especially in patients with hypertension. **The findings of this subanalysis provide opportunities to explore GM characteristics in patients with and without hypertension remission after RYGB.**

A hand is shown holding a pink, anatomical model of a brain. The background is a light blue field filled with various blue, textured, 3D models of brain cells or neurons, some of which are out of focus. The overall scene is a scientific or medical illustration.

Thank you for your attention!

carlos.schiavon@bp.org.br