







**Results** 

**Behavior** 



## Microbiota influence on behavior: Integrative analysis of serotonin metabolism and behavioral profile in germ-free mice

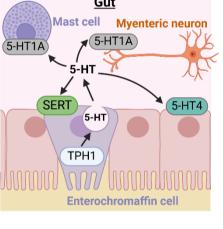
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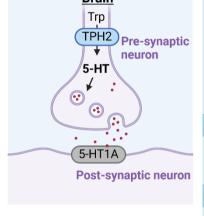
### Introduction

Previous studies on germ-free (GF) animals have described altered anxiety-like and social behaviors together with dysregulations in brain serotonin (5-HT) metabolism compared to conventional (CV) animals. In other studies, alterations in circulating 5-HT levels and gut 5-HT metabolism have also been reported in GF mice. We aimed to further this evidence through an integrative approach. We conducted an integrative analysis of various behaviors as well as markers of 5-HT metabolism in the brain and along the GI tract of GF male mice compared to CV ones.

# **Methods** Conventional (CV) **Germ-free (GF) Behavior** Locomotor activity Anxiety like behavior Social behavior Repetitive behaviors **Spatial memory**

#### Gut and brain 5-HT metabolism Gut **Brain**





Distance travelled % of distance in center p<0.0001

**Open-field test** 

→ Strongly **reduced** locomotor activity in GF

→ No difference in anxiety like-behavior in this test

#### Step-down test Latency to step-down

**CV** 

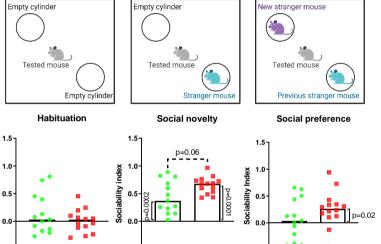
GF

Latency to 1st Grooming Total grooming time **ⓒ** 30

**Self-grooming test** 

→ Signs of increased anxiety in GF mice compared to CV in the step-down and selfgrooming tests

## Social behavior



social behavior in GF mice compared to CV

→Trend to an improved

No difference in repetitive behaviors and spatial memory were found.

for

## **Brain 5-HT metabolism**

→ We investigated gene expression of genes coding for TPH2 and 5-HT1A in the PFC, hippocampus, nucleus accumbens and cerebellum but found no differences between groups.

## **Conclusion**

- signs of increased anxiety-like behavior and a trend to improved sociability were observed in GF mice compared with CV mice. More in-depth analysis of locomotion and motor function in GF mice could be interesting.
- microbiota is involved in regulation of behavior, although heterogeneity among studies suggests a strong impact of genetic and environmental factors on this microbiotamediated regulation.
- its action on the immune system and the vagus nerve.

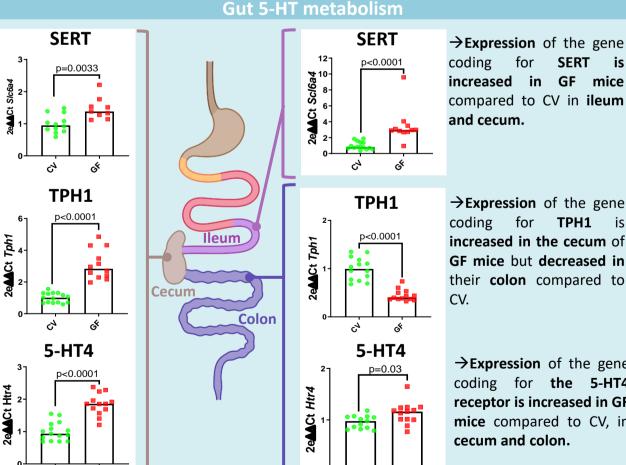
A strong decrease in locomotor activity, accompanied by

- These results add to previous evidence that the
- Brain gene expression analysis showed no differences in expression of genes coding for 5-HT1A and TPH2. It is possible that measuring other markers or studying other brain regions would have yielded different results.
- However, substantial differences in gut 5-HT metabolism were noted, with tissue-dependent results indicating a varying role of microbiota along the GI tract. The behavioral alterations we observed could be in part linked to those changes in gut 5-HT metabolism through
- As the gut is at the interface between intestinal microbiota and the central nervous system, it should be considered more systematically in future studies evaluating the influence of the gut microbiota on brain function and behavioral responses.

🗓 SCAN ME

**This work was** recently published:

**FASEB** Journal



→ Expression of the gene coding for TPH1 increased in the cecum of GF mice but decreased in their colon compared to

**SERT** 

GF

**→Expression** of the gene coding for the 5-HT4 receptor is increased in GF mice compared to CV, in cecum and colon.

**→Expression** of the gene coding for the 5-HT1A receptor is decreased in the colon of GF mice compared to CV.

**5-HT1A** 

e▲ACt Htr1a