Gut dysbiosis and its Association with ASD

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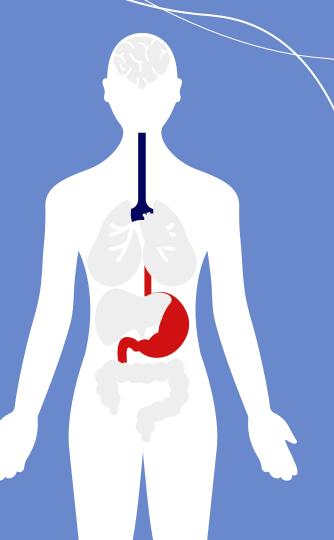
Outline

Introduction

Specific Aims

Research Plan

Results and Conclusion



Introduction

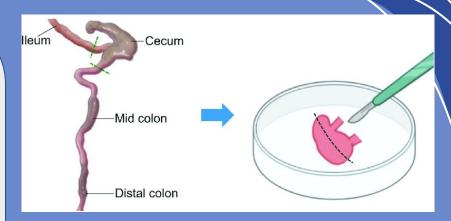
Gut Microbiota

- A system of trillions of microorganisms
- Function in the body
- Diet and microbiota
- Cecum



Cecum

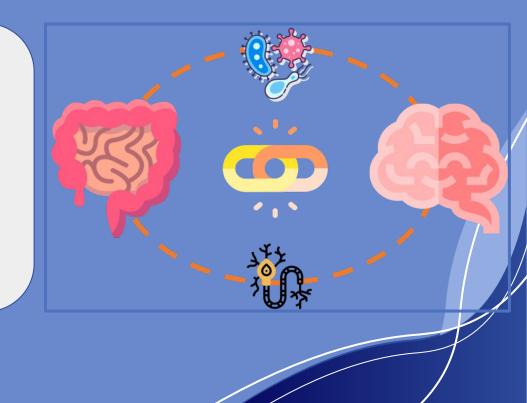
- Located in the lower left side of the abdominal cavity
- **Cecum in mice is enlarged**
- Contains bacteria that aid in digestion of plant matter
- Facilitates nutrient absorption





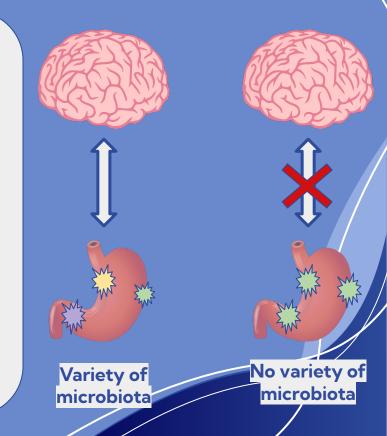
Gut-Brain Axis

Two-way biochemical
signaling that takes place
between the GI tract and
the CNS



Gut Dysbiosis and ASD

- Imbalance of gut microbiota
- Lifestyle choices can have an impact
- Correlation between severity of gut issues and ASD
- Affects certain signaling pathways
- Disrupts Blood Brain Barrier
- Probiotics role



Probiotics

Healthy bacteria

(eg. Bifidobacterium & Lactobacillus)

Improve gastrointestinal diseases

Probiotics

Boosts immunity

Modulates gut microbiota

Bifidobacterium Bifidum

- Live in the GI tract
- Digests food, absorb nutrients and prevents infection
- **Game Bifidobacterium** species generate GABA

GABA (Gamma-Aminobutyric Acid) :

- Inhibitory neurotransmitter
- Decreasing nerve cell hyperactivity (fear, stress and anxiety)

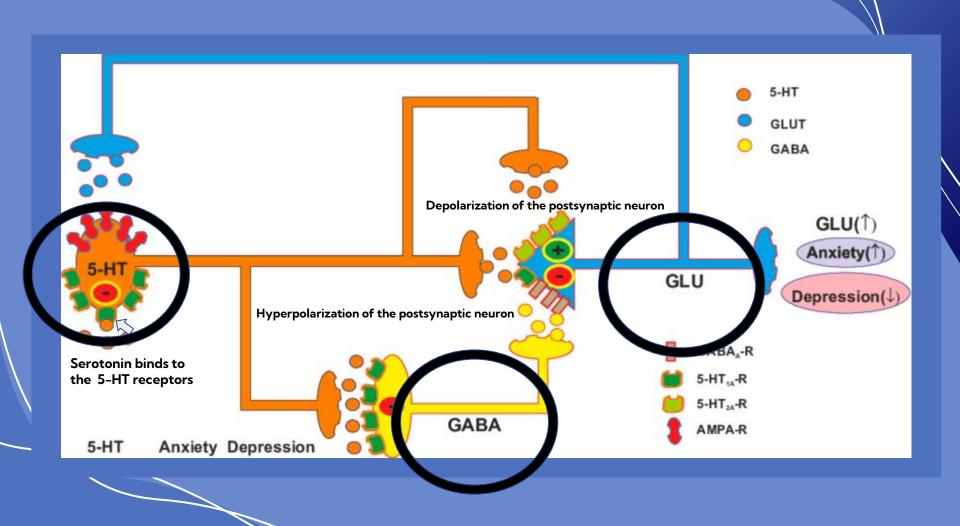
Children with ASD

- Lower levels of GABA \rightarrow high levels of anxiety

Lactobacillus

- Rod-shaped bacteria
- Produces lactic acid
- Production of fermented dairy products

	Lactobacillus Helveticus		Lactobacillus Paracasei	
•	Regulates the 5HT that balances the excitatory and inhibitory neurotransmission in the PNS and CNS	•	Increases glutamate and GABA in ASD patients.	



Gut Dysbiosis and ASD

Knowns

- The human gut is home to a variety of microbes
- ASD is associated with an unbalanced gut microbiota
- Children with ASD are at a greater risk of Gl concerns
- Probiotics exert beneficial effects in both the gut and the brain

Questions

- Is there a cause- effect relationship between ASD and gut microbiota?
- Is gut microbiota in ASD patients different from someone who does not have ASD ?
- Does adding probiotics alter the gut microbiota in ASD patients or impact their behaviors ?

Hypothesis

16s rRNA gene can help us identify gut microbiota and manipulating microbiota by administering probiotics can be used as a treatment for ASD by decreasing autism-related behaviors.

Specific Aims

Specific Aims

Specific Aim 1

Identification of gut microbiota through 16S rRNA gene in BTBR and C57 mice.

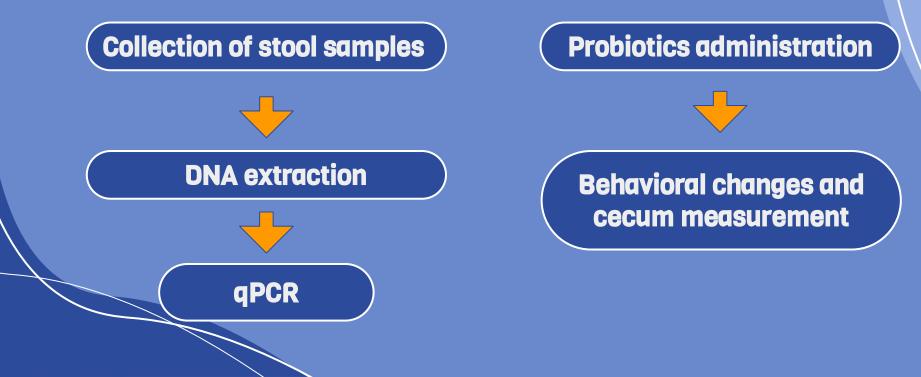


Evaluation of autism-related behavioral phenotypes and cecum measurement in BTBR and C57 mice after administering probiotics.

Research Plan

Experimental approach

1. Identifying Microbiota



2. Modification of diet

Animal models

3 BTBR mice

Absence of corpus callosum
Smaller Hippocampal Commissure
Exhibits autism-like behaviors

3 C57BL/6 mice
Most widely used strain in
biomedical research
Exhibits wild type behaviors





Identifying Microbiota

1. Collection of stool samples

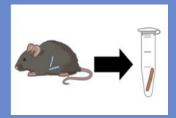
Place mouse in cage Disinfect cage with 70% ethanol Wait for mice to defecate

Up to 40 minutes of waiting.



Collect sample and place in tube

Later on used for DNA extraction

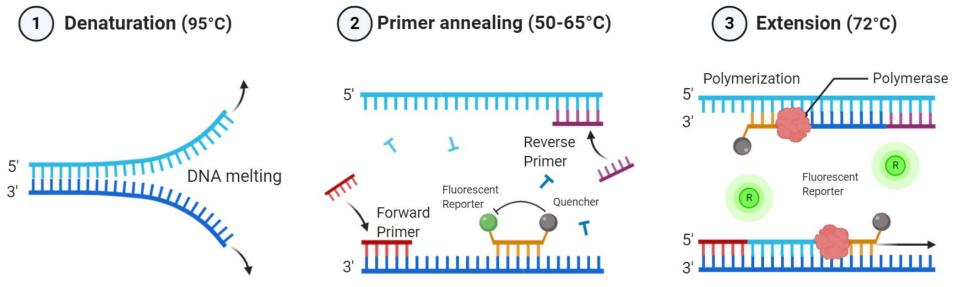


Identifying Microbiota

2. DNA extraction

- 1. Disruption of cells to release DNA for collection
- 2. Removal of Contaminating Biomolecules with buffers and washers





Identifying Microbiota

3. qPCR

Modification of diet

1. Probiotics administration

10⁶ CFU of probiotic for a total of 3 days Including Lactobacillus and Bifidobacterium









Modification of diet

2. Behavioral data collection process 10 minute home cage recordings Excessive Grooming Rearing Behaviors Nibbling "Spidermouse" Cage location





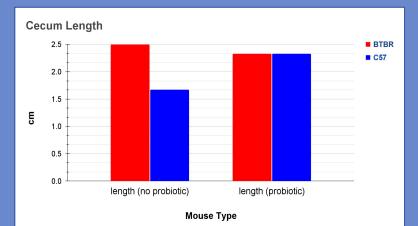
qPCR Microbiota Identification

Qualitative - <u>qPCR</u> <u>Identification</u> shows us whether or not a microbiota is present

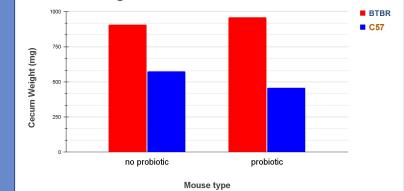
Next steps - <u>qPCR</u> <u>Profiling</u> which would give us *quantitative results* of how much of a specific microbiota is expressed

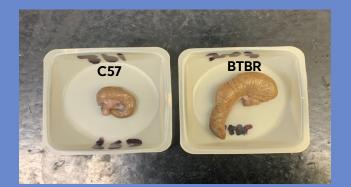
	C57	C57	C57	BTBR	BTBR	BTBR
Bacteroides fragilis	+	+	+	+	+	+
Escherichia coli	+	+	+	+	+	+
Candida albicans	+	+	+	+	+	+
Streptococcus agalactiae	+	+	+	+	+	+
Bifidobacterium longum	+	+	+	+	+	+
Enterococcus faecalis	+	+	+	+	+	+
Helicobacter pylori	+	+	+	+	+	+
Streptococcus mitis	+	+	+	+	+	+
Citrobacter freundii	+	+	+	+	+	+
Clostridium perfringens	+	+	+	+	+	+

Cecum measurements



Mice cecal weight



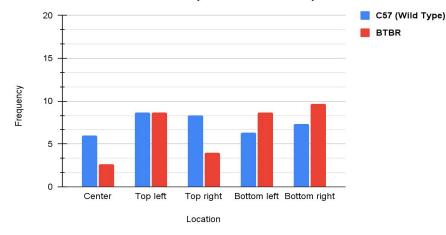


Behavioral observations

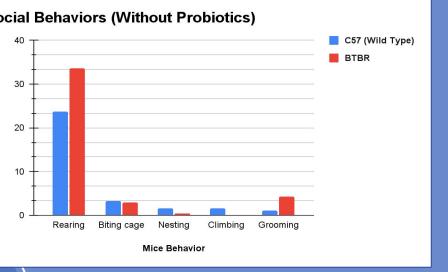
Mice Movement Patterns

Mice Movement Patterns (Without Probiotics)

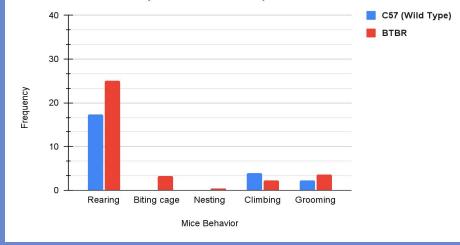
Mice Movement Patterns (With Probiotics)



Social Behaviors



Social Behaviors (With Probiotics)



Conclusion

qPCR Identification

The data showed us that gut microbiota associated with ASD is present in both mice

<u>Behavioral observation</u>

Repetitive behaviors associated with anxiety decreased while interactive and explorative behaviors increased

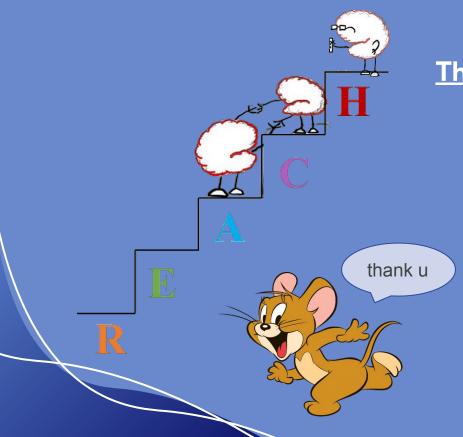
<u>Cecum measurements</u>

- Probiotics caused a slight increase BTBR cecal size probably due to an influx of microbiota
- □ C57 cecal size minor fluctuation due to already normal function

Limitations/Future Steps

- Mice might not eat the probiotics
- □ Time (conducting the procedure within a specific time frame)
- Explore Different types of behavioral analysis experiments
- We will experiment on different types of probiotic strains

Acknowledgements



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Thank you! :)

Questions?

