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Exploiting the Gastrointestinal Microbiota as a Therapeutic Target for Type 1 Diabetes

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Introduction

Objective: To investigate a novel therapeutic for the treatment of Type 1 Diabetes (T1D).

The therapeutic would not kill, but just retard the growth of bacteria that are members of the genus *Bacteroides* while not effecting other bacterial species.

Why: T1D is a progressive autoimmune disorder characterized by the destruction of the insulin secreting Beta cells in the pancreas (Harrison *et al.* 2008).

While it has been previously acknowledged that there are genetic factors responsible for the onset of T1D, there is evidence to suggest environmental causes as well (Achenbach *et al.*, 2005; Gale, 2002).

Not everyone that has the genetic predisposition to developing the disease will have T1D.

Our interest lies in the observation that just before the onset of T1D in an individual, there is a bloom of bacteria from the Bacteroidetes phyla. (Giongo *et al.*, 2011).

How: We are investigating therapeutics that can inhibit the Starch Utilization System (SUS) specific to members of *Bacteroides*. Removing the SUS as a method of gaining biomass and energy for the cell would not kill the cell but slow its growth. The cell will have to find alternative and less effective methods of metabolism.

Current Work: We have investigated the effects of the drugs Acarbose and Miglitol on the growth of members from the *Firmicutes* and *Bacteroidetes* phyla.

Acarbose has been used previously in diabetes treatments because of its ability to retard the degradation of starch via inhibiting human α -amylases. α -amylases, similar to the ones observed in humans, can be found within the Sus.

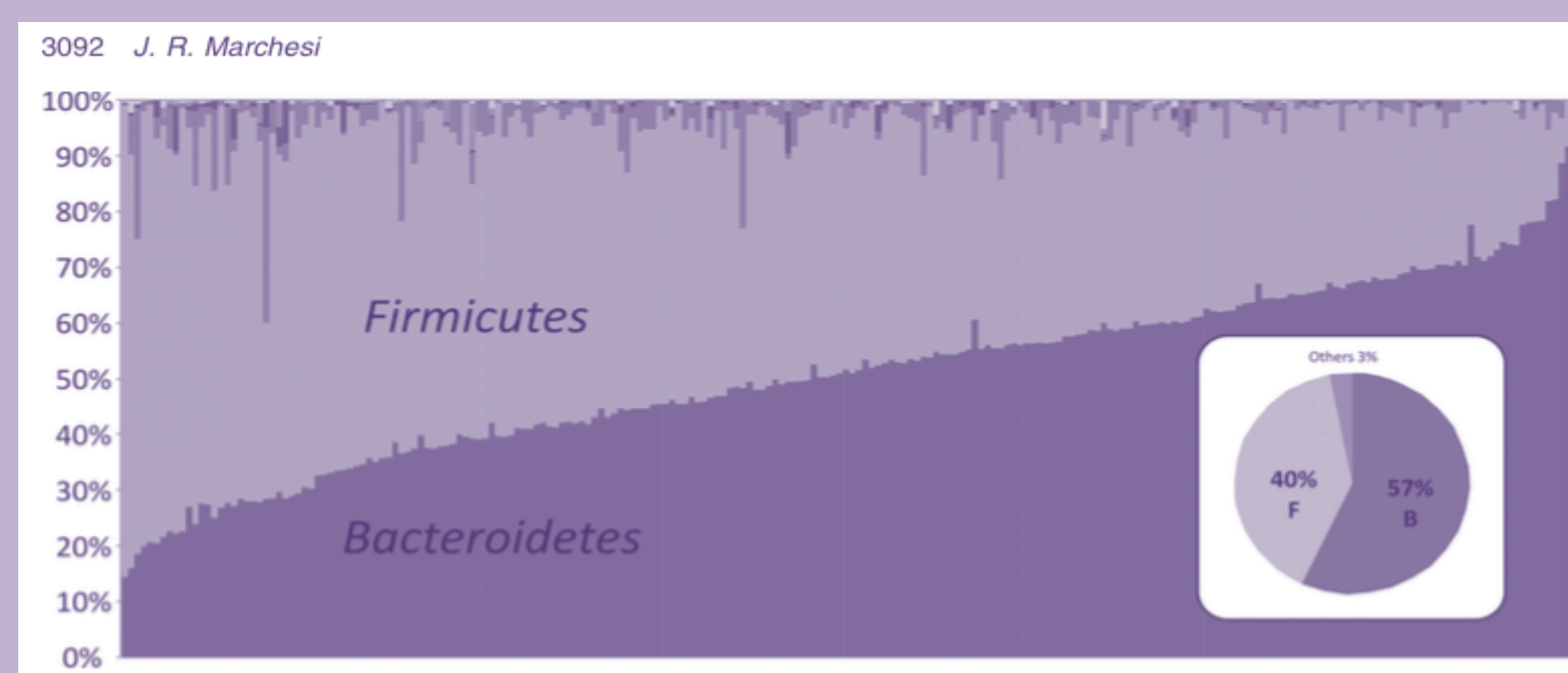


Figure 1. Variation of ratios of the two major phyla of bacteria, Firmicutes and Bacteroidetes, found in 386 fecal samples

(Figure from: Marchesi, J.R. 2011. Human distal gut microbiome. *Environmental Microbiology* 13: 3088-3102)

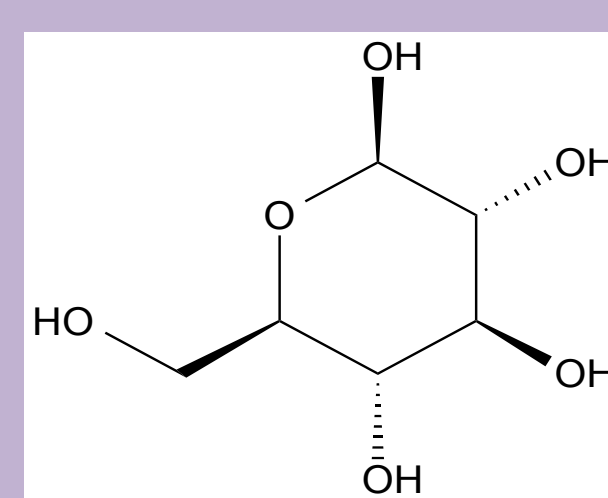


Figure 2. Molecular structure of D-glucose

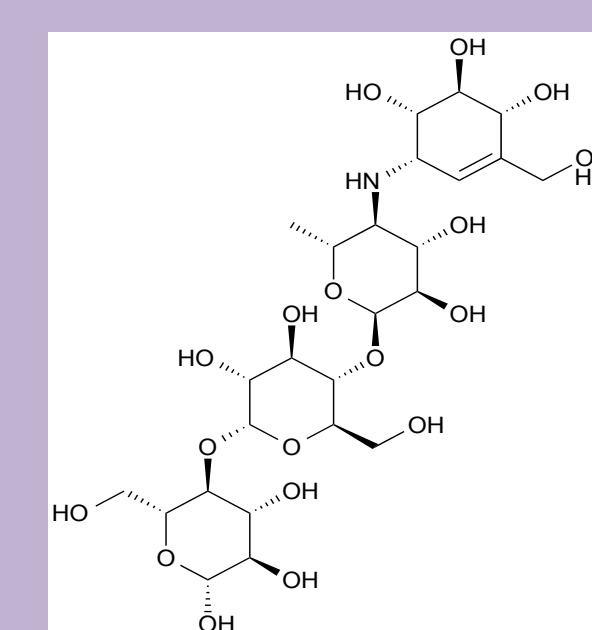


Figure 3. Molecular structure of Acarbose

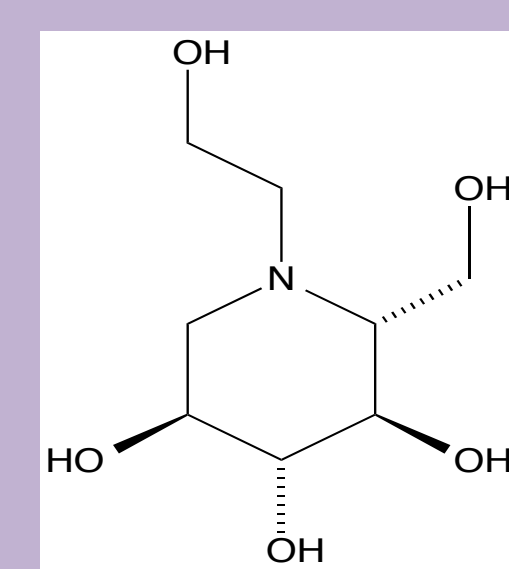


Figure 4. Molecular structure of Miglitol

Methods and Materials

- Bacteroides thetaiotaomicron* was grown on tryptone yeast glucose (TYG) agar anaerobically.
- Colonies from plates were used to prepare overnight cultures of *B. thetaiotaomicron* TYG broth (grown anaerobically).
- Test tubes were filled with minimal media containing either 0.5% glucose, maltose, or pullulan as a carbon source, cysteine, *Bacteroides* culture, and varying amounts of acarbose.
- An anaerobic environment was created by inserting a cotton ball in the tube and lighting it on fire.
- Upon completion of burning, sodium bicarbonate and pyrogallol were added.
- The tube was stoppered, sealed, and incubated at 37°C for 24 hours.
- Inhibition was indicated the following day by measuring the optical density at a wavelength of 600nm.
- Lactobacillus reuteri* strains were grown on de Man, Rogosa and Sharpe (MRS) agar aerobically.
- Colonies from plates were used to prepare overnight cultures of *L. reuteri* MRS broth (grown aerobically).
- Test tubes were filled with MRS broth containing either 0.5% glucose, maltose, or pullulan as a carbon source, cysteine, *Lactobacillus* culture, and varying amounts of acarbose.
- The test tubes were incubated at 37°C for 24 hours.
- Inhibition was indicated the following day by measuring the optical density at a wavelength of 600nm.

Preliminary Results

Bacteroidetes representative: *B. ovatus*, *B. theta*, *B. fragilis*

Firmicutes representatives: *L. reuteri* ATCC 55730 and *L. reuteri* PTA 6475

Monosaccharide representative: Glucose

Disaccharide representative: Maltose

Polysaccharide/Glycan representative: Pullulan

Table 1. Inhibition of Bacterial Growth by Acarbose

Species	Glucose	Maltose	Pullulan
<i>B. theta</i>	None	At 200 μ M	At 50 μ M
<i>B. fragilis</i>	None	Not tested	At 10 μ M
<i>L. reuteri</i> ATCC 55730	None	None	None
<i>L. reuteri</i> PTA 6475	None	None	None

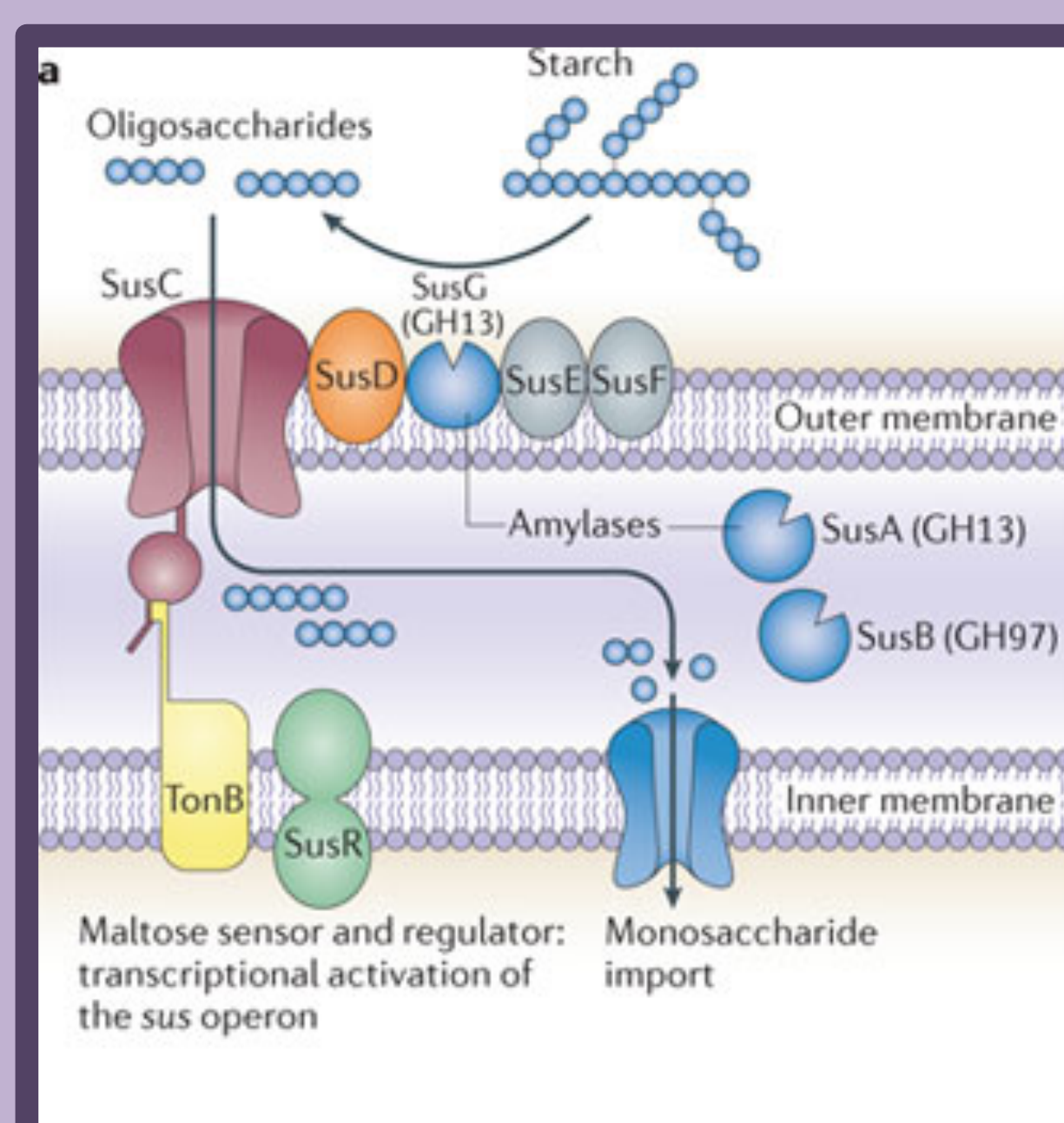


Figure 5. Starch Utilization System (Sus) for *B. thetaiotaomicron*. This is the pathway responsible for recognition, degradation, and importation of complex carbohydrates. (Figure from: Koropatkin, N.M., E.A. Cameron, E.C. Martens. 2012. *Nature*, 10: 328.)

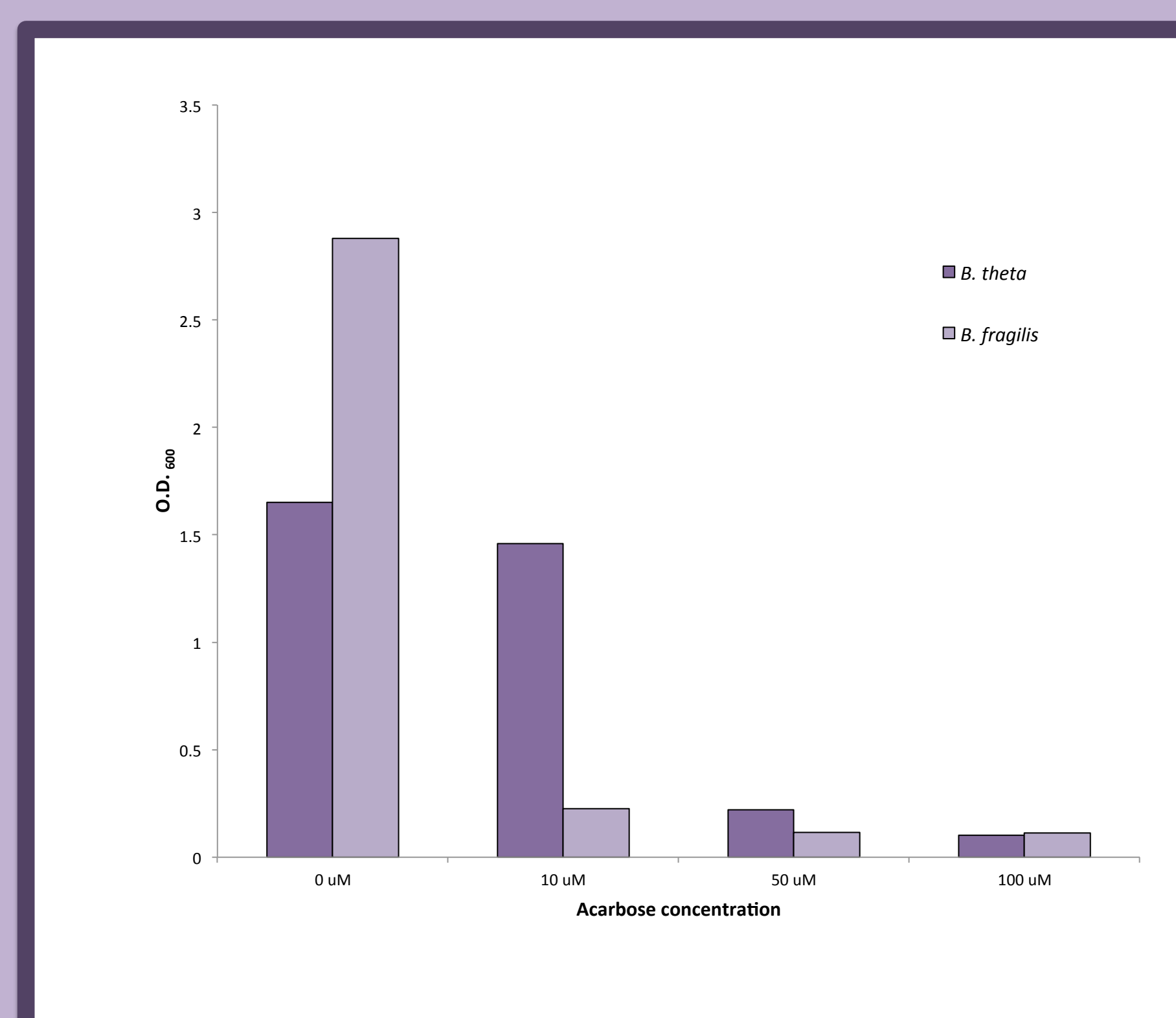


Figure 6. Representative growth assay demonstrating acarbose inhibition of *B. theta* and *B. fragilis* grown with pullulan.

Conclusions

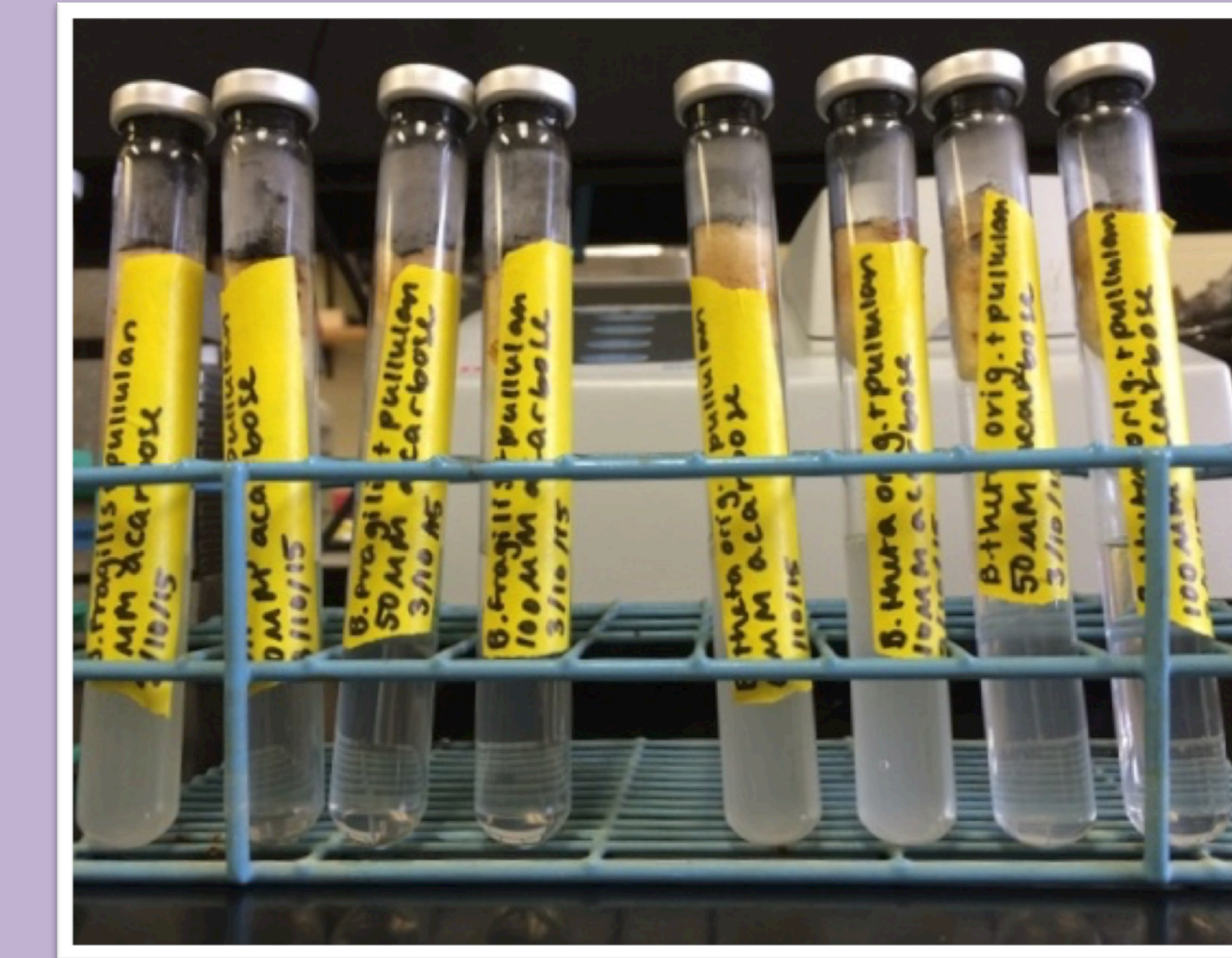


Figure 7. Anaerobic cultures of *B. theta* and *B. fragilis* with varying levels of acarbose.

What we have found:

- Acarbose inhibits the growth of *Bacteroides* spp. and is not lethal to the cells.
- Acarbose does not inhibit *Lactobacillus* growth.
- Acarbose has structural similarities to many glycan molecules and is most likely effecting the Sus and therefore prohibiting its ability to breakdown available glycan molecules.

These findings have the possibility of providing a novel T1D treatment by preventing the bloom of Bacteroidetes in the large intestine.

Our future directions include:

- Assay inhibition on other complex carbohydrates
- Investigate inhibitory effects with another *Bacteroides* sp. and a non-*Bacteroides* sp. within the Bacteroidetes phylum
- Establish exact mode of inhibition of Sus system in *B. thetaiotaomicron* (*i.e.* where does it bind to inhibit?)
- Determine if correlation is an indicator of causation by conducting *in vivo* experiments using the Non-obese diabetic (NOD) mouse model to examine the effectiveness of acarbose administration at delaying or preventing the onset of T1D.

References

- Achenbach, P., E. Bonifacio, K. Koczwara, A. Ziegler. 2005. Natural history of type 1 diabetes. *Diabetes* 54:S25-S31.
- Gale, E.A.M. 2002. The rise of childhood type 1 diabetes in the 20th century. *Diabetes* 51:3353-3361.
- Caicedo, R.A., N. Li, C. Des Robert, P.O. Scumpia, C.P. Hubsher, C.H. Wasserfall, D.A. Schatz, M.A. Atkinson, J. Neu. 2008. Neonatal formula feeding leads to immunological alterations in an animal model of type 1 diabetes. *Pediatric Research* 63: 303-307.
- Visser, J., S. Brugman, F. Klatte, L. Vis, H. Groen, J. Strubbe, J. Rozing. 2003. Short-term dietary adjustment with a hydrolyzed casein-based diet postpones diabetes development in the diabetes-prone BB rat. *Metabolism-Clinical and Experimental* 52: 333-337.
- Giongo, A., K.A. Gano, D.B. Crabb, N. Mukherjee, L.L. Novelo, G. Casella, J.C. Drew, J. Ilonen, M. Knip, H. Hyoty, R. Veijola, T. Simell, O. Simell, J. Neu, C.H. Wasserfall, D. Schatz, M. Atkinson, E.W. Triplett. 2011. Toward defining the autoimmune microbiome for type 1 diabetes. *ISME Journal* 5: 82-91.
- Greiner, T. and F. Backhed. 2011. Effects of the gut microbiota on obesity and glucose Homeostasis. *Trends in Endocrinology and Metabolism* 22:117-123.
- Musso, G., R. Gambino, M. Cassader. 2011. Interactions Between Gut Microbiota and Host Metabolism Predisposing to Obesity and Diabetes. *Annual Review of Medicine*, 62:361-380
- Wen, L., R.E. Ley, P.Y. Volchkov, P.B. Stranges, L. Avanesyan A.C. Stonebraker, C.Y. Hu, F.S. Wong, G.L. Szot, J.A. Bluestone, J.I. Gordon, A.V. Chervonsky. 2008. Innate immunity and intestinal microbiota in the development of Type 1 diabetes. *Nature* 455:1109-U1110.
- Brown, C.T., A.G. Davis-Richardson, A. Giongo, K.A. Gano, D.B. Crabb, N. Mukherjee, G. Casella, J. C. Drew, J. Ilonen, M. Knip, H. Hyoty, R. Veijola, T. Simell, O. Simell, J. Neu, C.H. Wasserfall, D. Schatz, M.A. Atkinson, E.W. Triplett. October 2011. Gut Microbiome Metagenomics Analysis Suggests a Functional Model for the Development of Autoimmunity for Type 1 Diabetes. *PLoS ONE* 6:1-9. [Online.] <http://www.plosone.org>.
- Koropatkin, N.M., T. J. Smith. 2010. SusG: A Unique Cell-Membrane-Associated α -Amylase from a Prominent Human Gut Symbiont Targets Complex Starch Molecules. *Structure*, 18: 200-215
- Marchesi, J.R. 2011. Human distal gut microbiome. *Environmental Microbiology* 13: 3088-3102
- Frank, D.N., A.L. St Amand, R.A. Feldman, E.C. Boedeker, N. Harpaz, N.R. Pace. 2007. Molecular-phylogenetic characterization of microbial community imbalance in human inflammatory bowel diseases. *Proc Natl Acad Sci USA* 104: 13780-13785.
- Harrison, L.C., M.C. Honeyman, G. Morahan, J.M. Wentworth, S. Elkassaby, P.G. Colman, S. Fourlanos. 2008. Type 1 diabetes: lessons for other autoimmune diseases? *J Autoimmun* 31: 306-310.

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