

# Gut Biome and Graft-versus-Host-Disease: Do Organisms in the Gut Contribute to GVHD?

**Celebrating a Second Chance at Life  
Survivorship Symposium**

April 27 – May 3, 2024



**Robert Jenq MD**

Director of MD Anderson Microbiome Core  
Facility and Associate Professor in the  
Department of Genomic Medicine,  
MD Anderson Cancer Center

# Gut Biome and GVHD: Do organisms in the gut contribute to GVHD?

Robert Jenq, MD

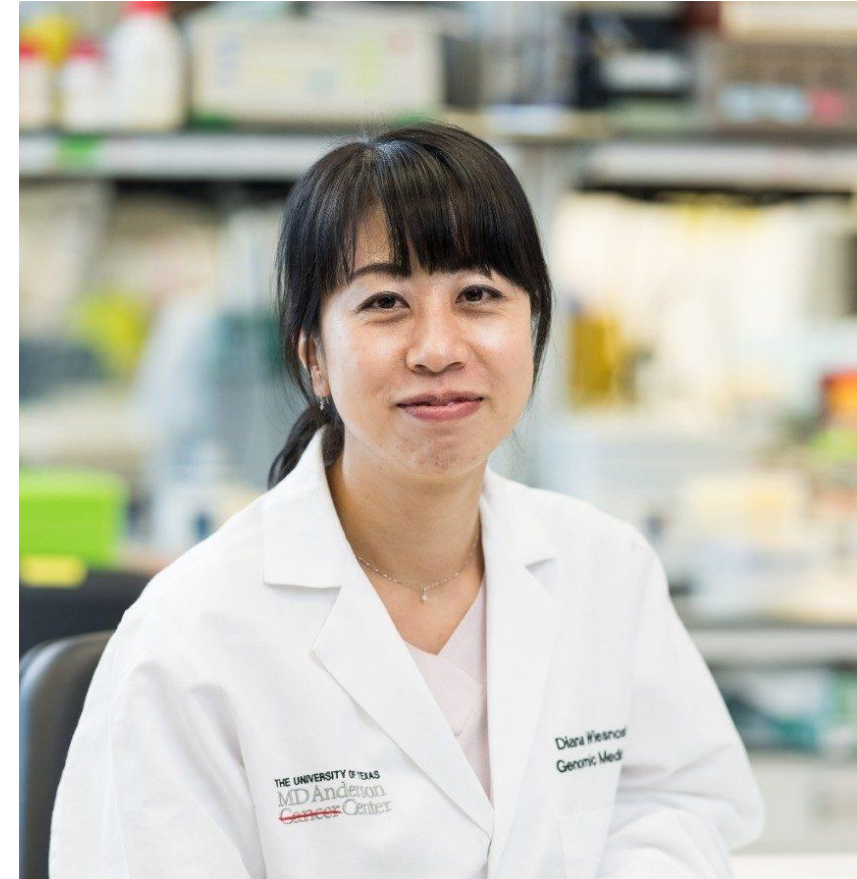
May 2, 2024

[rrjenq@mdanderson.org](mailto:rrjenq@mdanderson.org)

# Disclosures

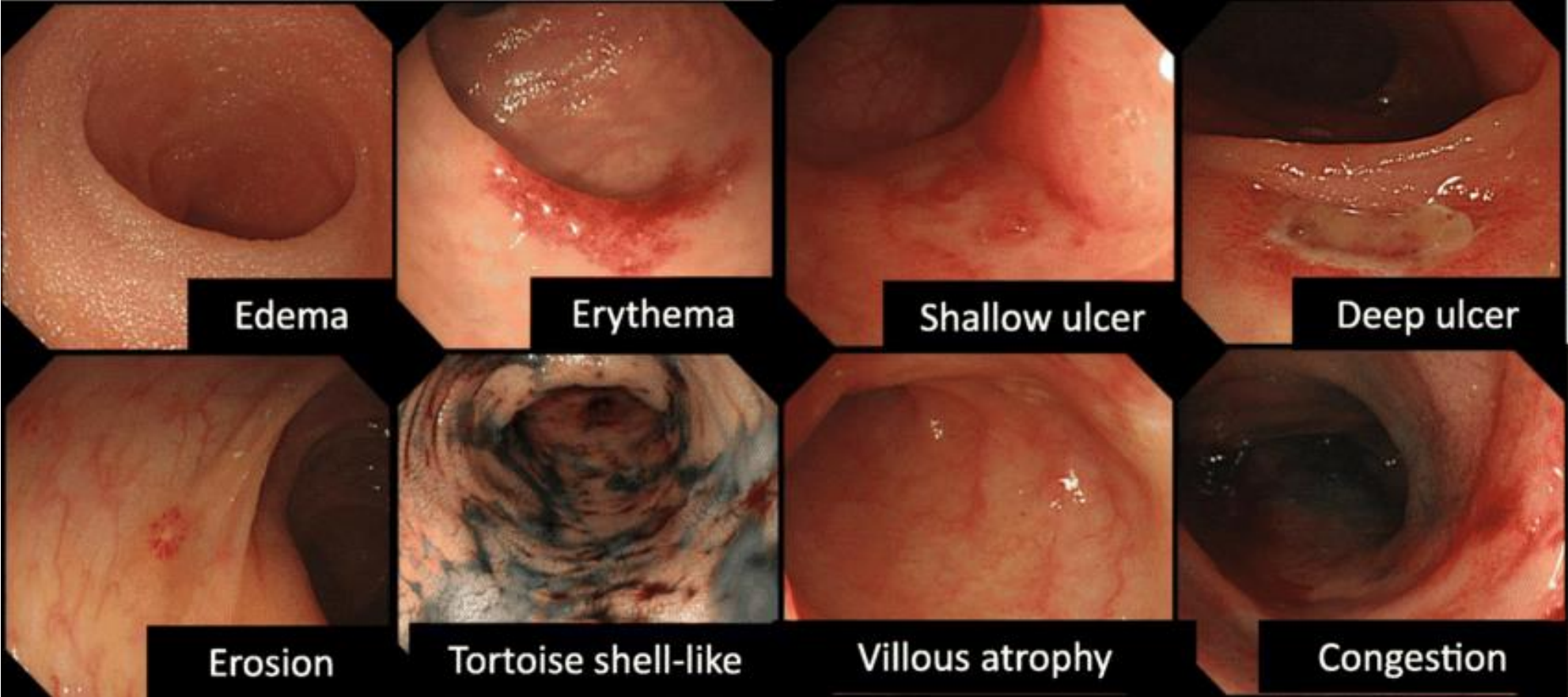
- Advisory board for LIScure Biosciences, Maat Pharma, Prolacta Biosciences, and Seres Therapeutics
- Royalties from Seres Therapeutics

# Acknowledgements



*Cell, 2022 Hayase, et al  
Unpublished work, in revision*

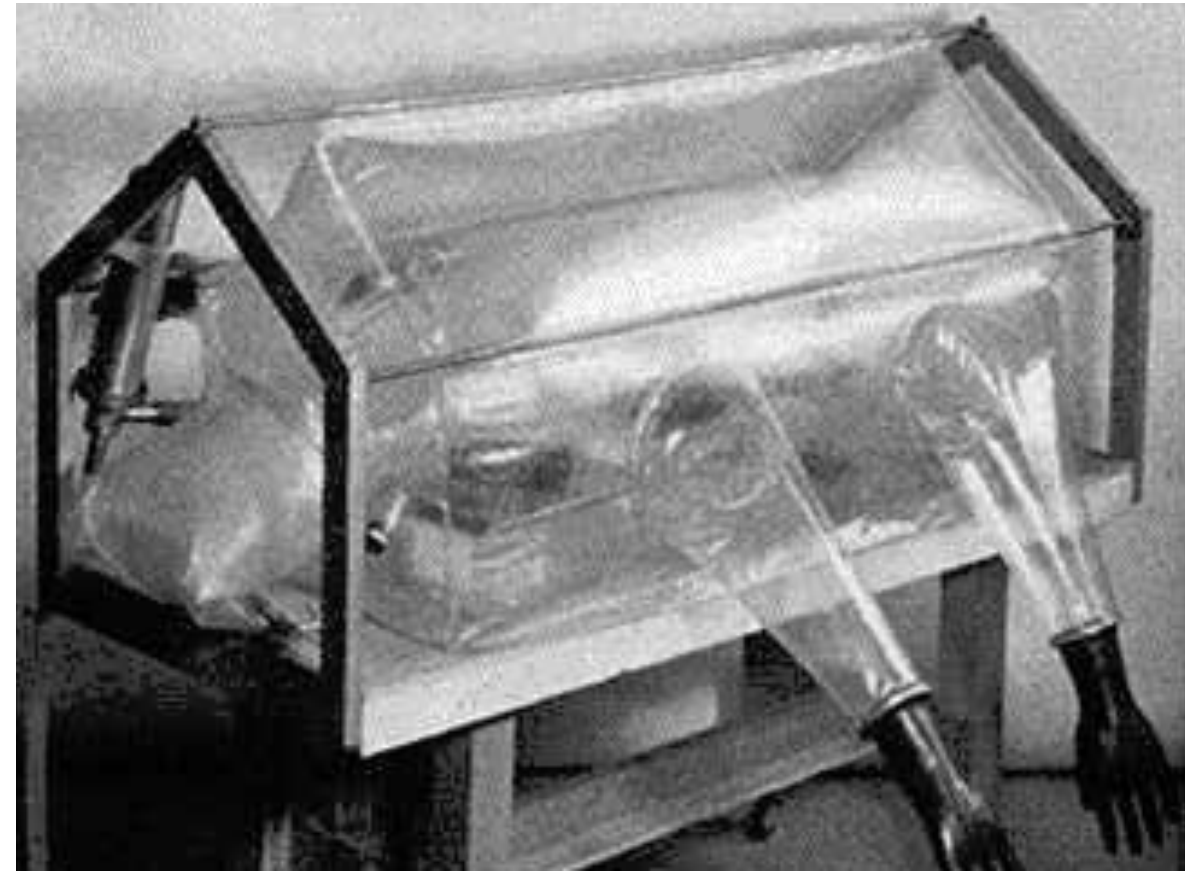
# Colonoscopy Findings in Acute GI GVHD



Sugihara, et al, BMC Gastroenterology 2018



Steam sterilizable metal isolator  
(Notre Dame University, 1928)



1962 Trexler isolator



2021 Class Biologically Clean (CBC) flexible film isolators

# Bacteria contribute to the development of GVHD

**Mortality and Gross Pathology of Secondary Disease in  
Germfree Mouse Radiation Chimeras<sup>1</sup>**

1971

J. MIRIAM JONES<sup>2</sup>, RAPHAEL WILSON, AND PATRICIA M. BEALMEAR



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**Mitigation of Secondary Disease of Allogeneic Mouse Radiation Chimeras by Modification of the Intestinal Microflora<sup>1</sup>**

1974

D. W. van Bekkum, J. Roodenburg, P. J. Heidt, and D. van der Waaij<sup>2</sup>



# Fred Hutch transplantation clinic in the 1980s

# Controversies in Intestinal Bacterial Decontamination in Allo-HCT

- Clinical studies initially suggested a benefit from near-total bacterial decontamination (Storb et al., 1983)
- Later studies, however, showed no clear benefit (Peterson et al., 1987; Passweg et al., 1998; Russell et al., 2000).

ORIGINAL ARTICLE

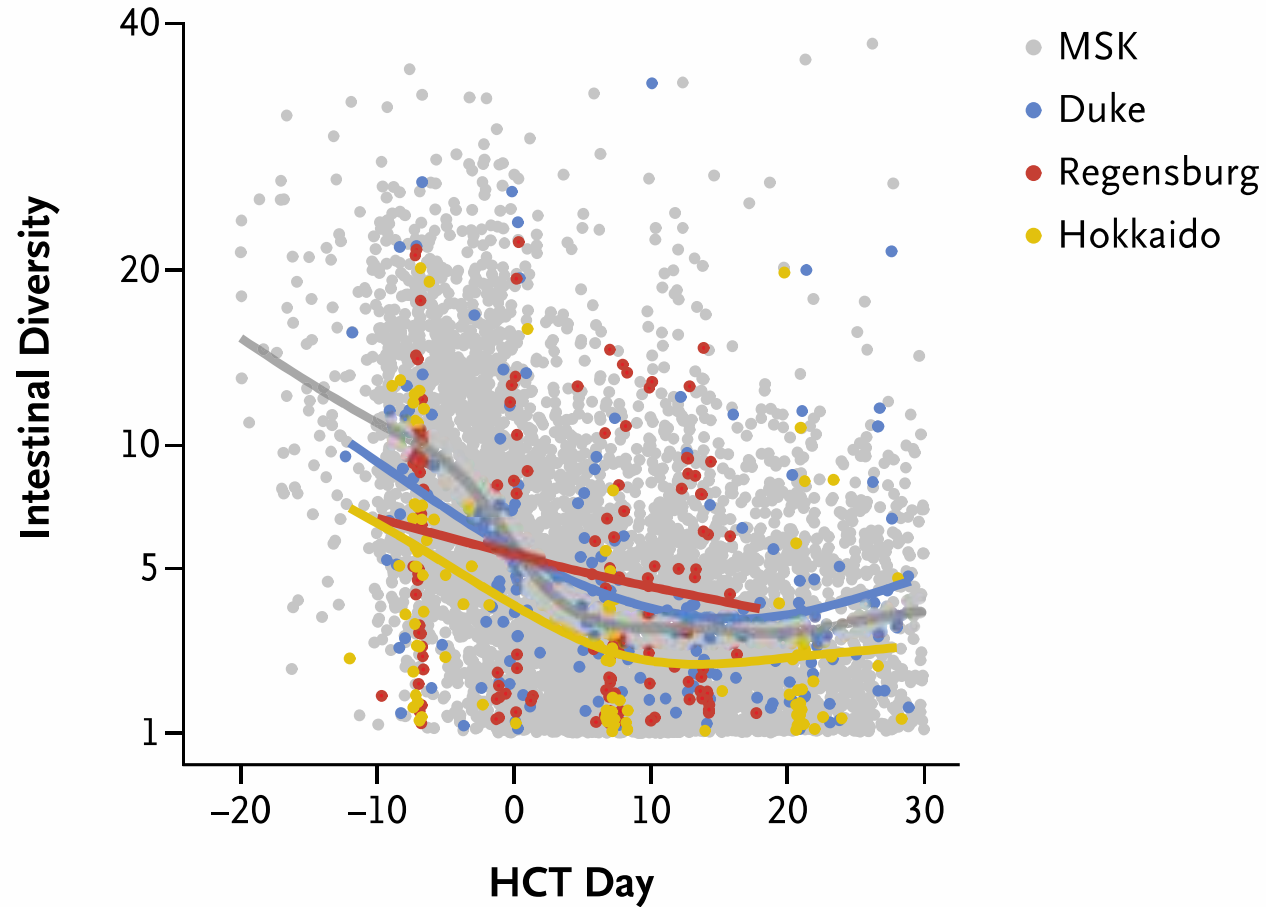
# Microbiota as Predictor of Mortality in Allogeneic Hematopoietic-Cell Transplantation

J.U. Peled, A.L.C. Gomes, S.M. Devlin, E.R. Littmann, Y. Taur, A.D. Sung, D. Weber, D. Hashimoto, A.E. Slingerland, J.B. Slingerland, M. Maloy, A.G. Clurman, C.K. Stein-Thoeringer, K.A. Markey, M.D. Docampo, M. Burgos da Silva, N. Khan, A. Gessner, J.A. Messina, K. Romero, M.V. Lew, A. Bush, L. Bohannon, D.G. Brereton, E. Fontana, L.A. Amoretti, R.J. Wright, G.K. Armijo, Y. Shono, M. Sanchez-Escamilla, N. Castillo Flores, A. Alarcon Tomas, R.J. Lin, L. Yáñez San Segundo, G.L. Shah, C. Cho, M. Scordo, I. Politikos, K. Hayasaka, Y. Hasegawa, B. Gyurkocza, D.M. Ponce, J.N. Barker, M.-A. Perales, S.A. Giralt, R.R. Jenq, T. Teshima, N.J. Chao, E. Holler, J.B. Xavier, E.G. Pamer, and M.R.M. van den Brink

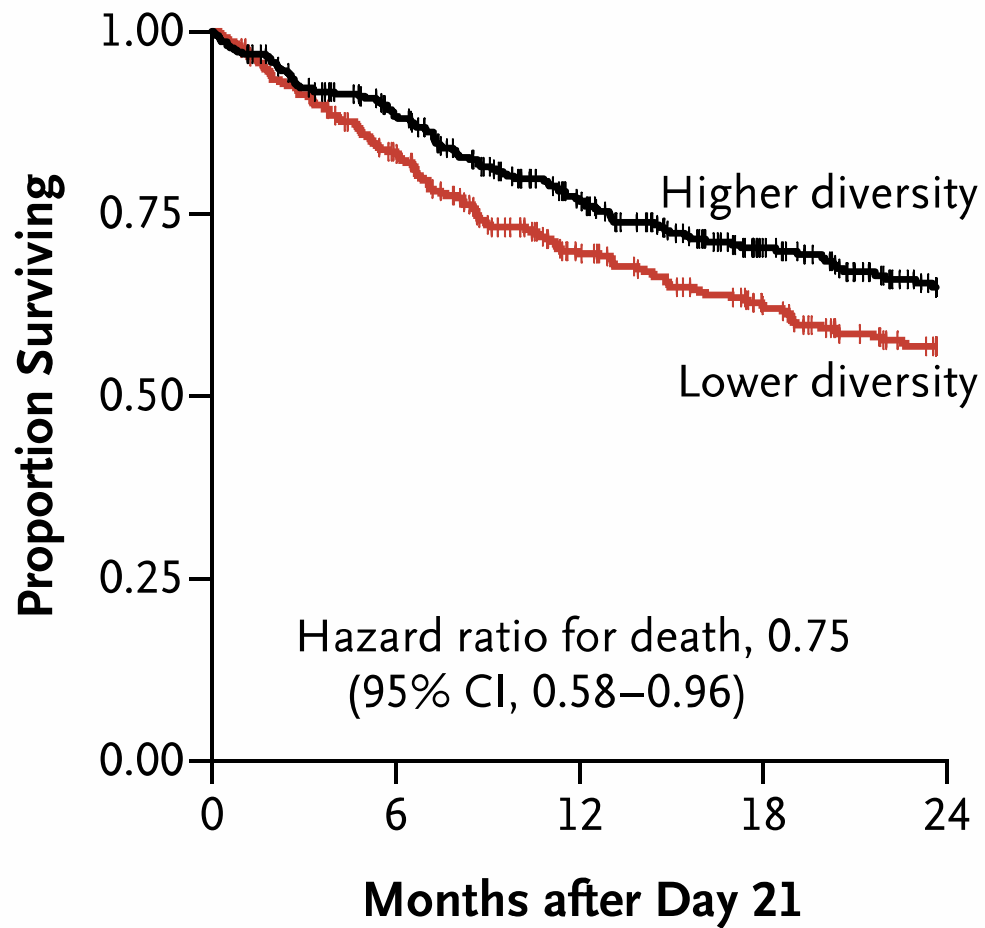
## RESULTS

We profiled 8767 fecal samples obtained from 1362 patients undergoing allogeneic hematopoietic-cell transplantation at the four centers

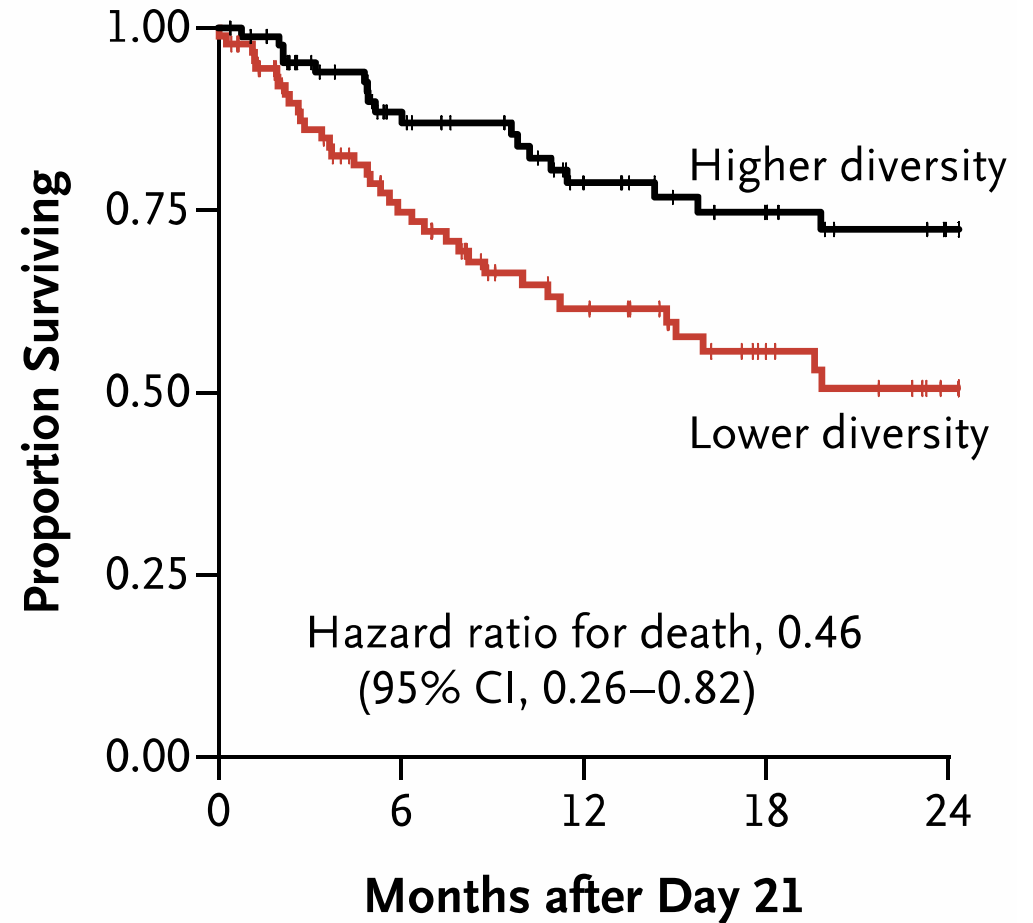
Change in Diversity of Intestinal Microbiota during HCT Period



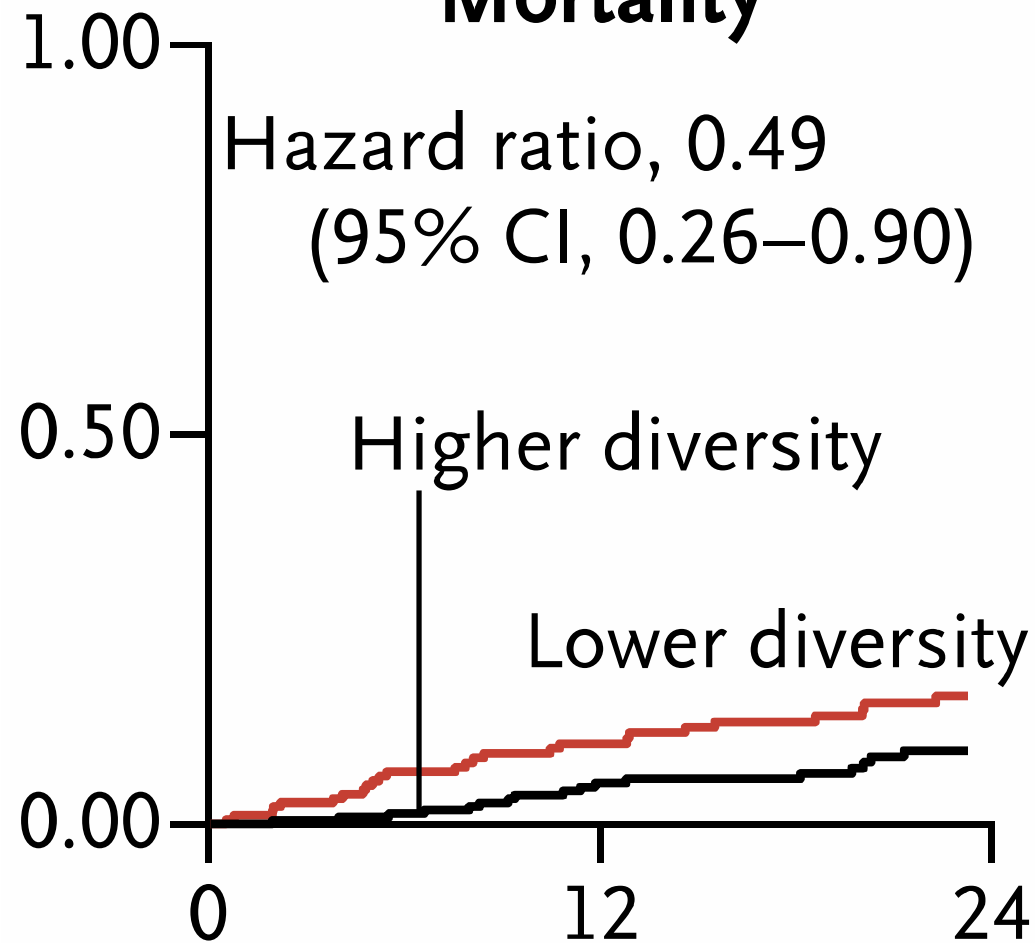
### Overall Survival — Cohort 1



### Overall Survival — Cohort 2



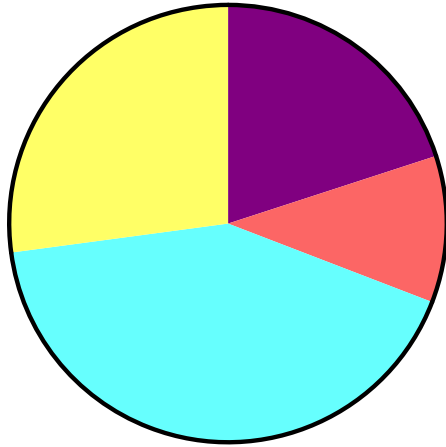
# GVHD-Related Mortality







	MSKCC Estimate ± S.E.	Regensburg Estimate ± S.E.	Duke Estimate ± S.E.	Hokkaido Estimate ± S.E.
Δt	<b>-0.33 ± .02</b>	-0.10 ± 0.17	<b>-0.15 ± 0.06</b>	-0.01 ± 0.06
cefepime	0.07 ± .07	-	-0.13 ± 0.22	0.18 ± 0.19
doripenem	-	-	-	0.11 ± 0.23
meropenem	0.08 ± 0.21	-0.02 ± 0.24	-0.51 ± 0.33	<b>-0.36 ± 0.15</b>
piperacillin-tazobactam	<b>-0.13 ± 0.05</b>	-0.24 ± se 0.29	-0.35 ± 0.28	-
teicoplanin	-	-	-	-0.15 ± 0.12



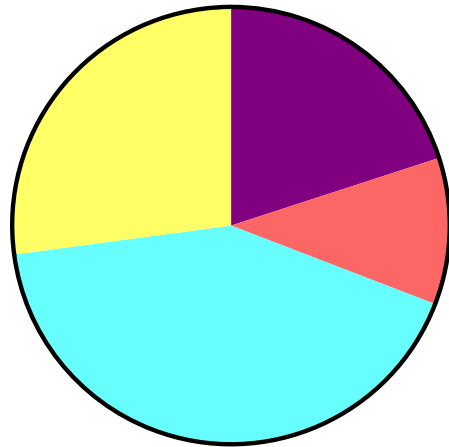
# Cohort of 295 patients with AML or MDS who underwent allogeneic HCT at MD Anderson



Total=295

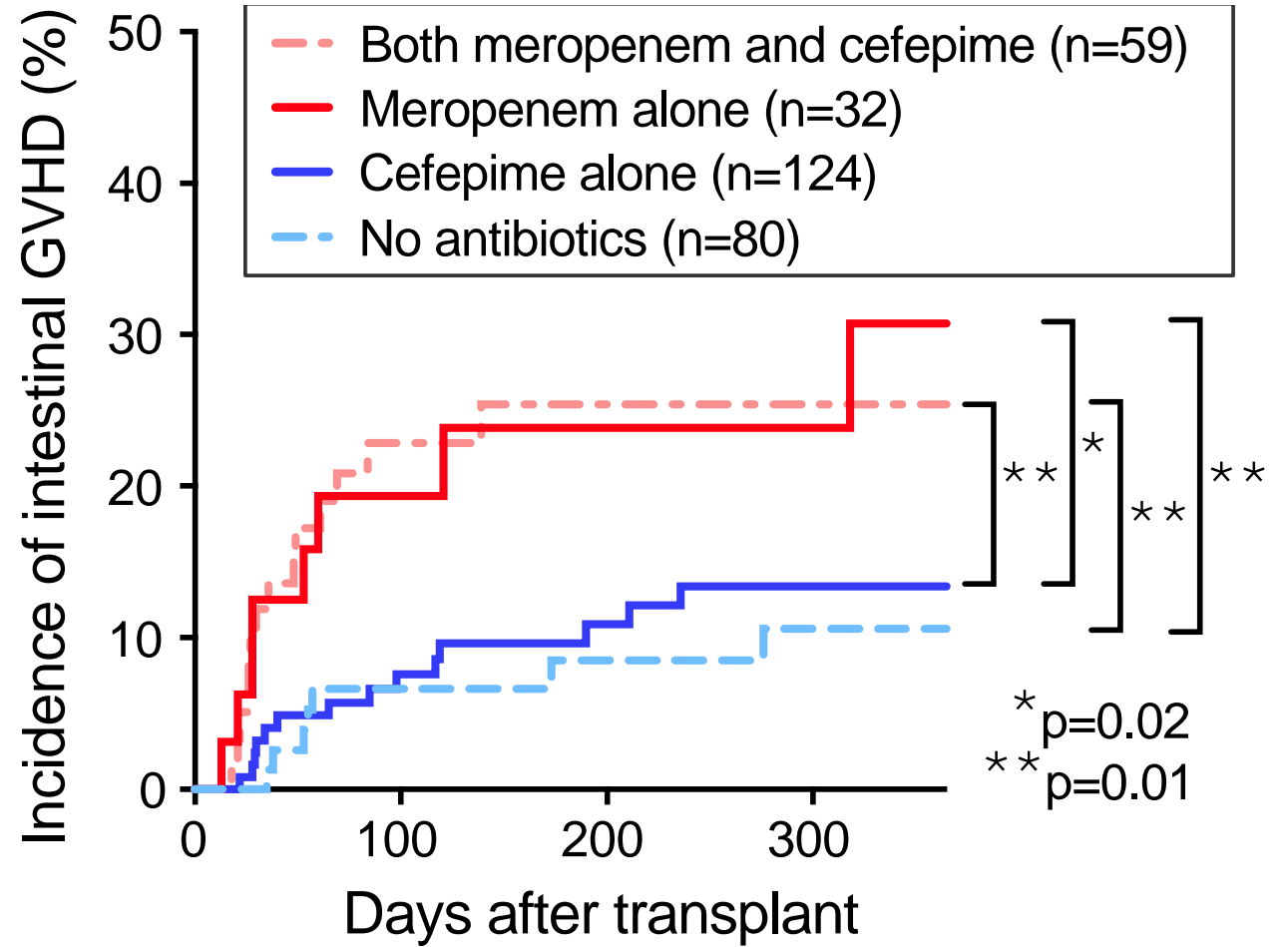
-  No antibiotics
-  Cefepime
-  Meropenem
-  Both meropenem and cefepime

# Cohort of 295 patients with AML or MDS who underwent allogeneic HCT at MD Anderson

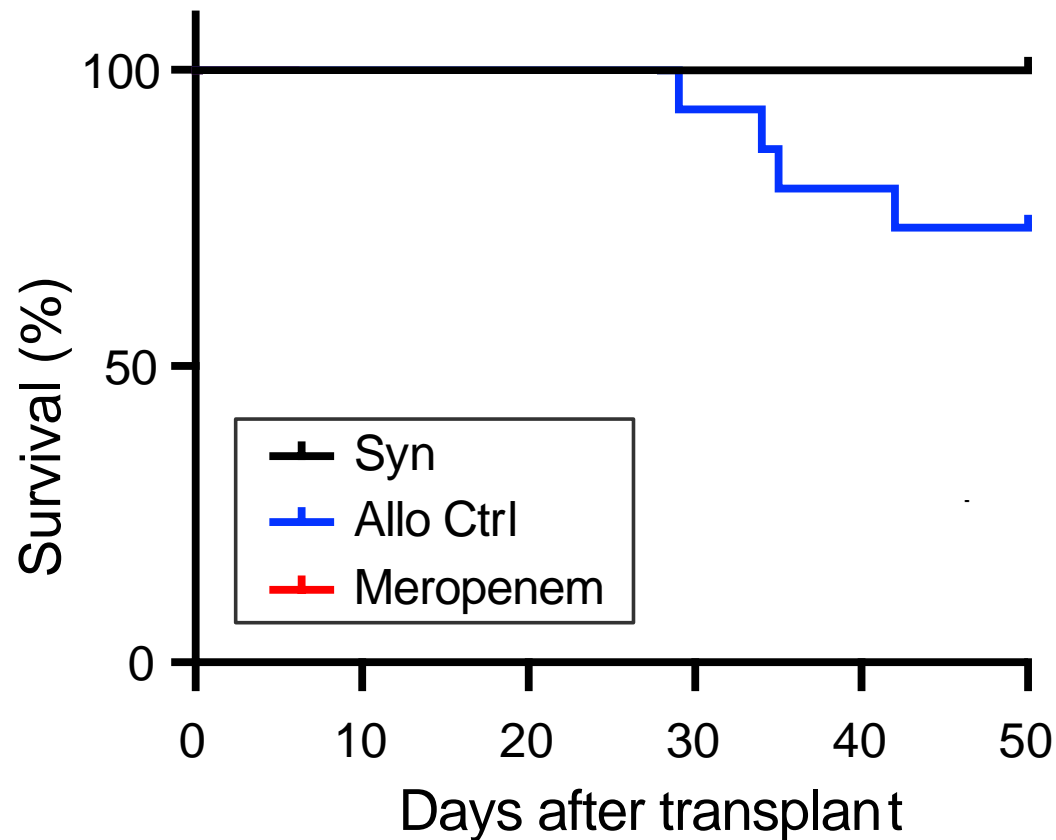
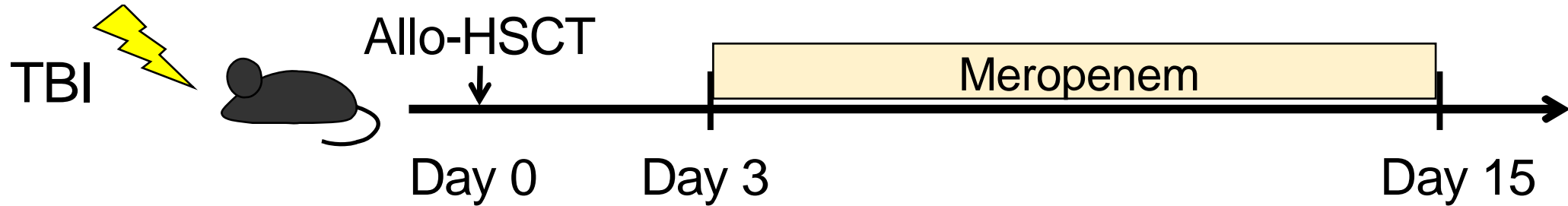


Total=295

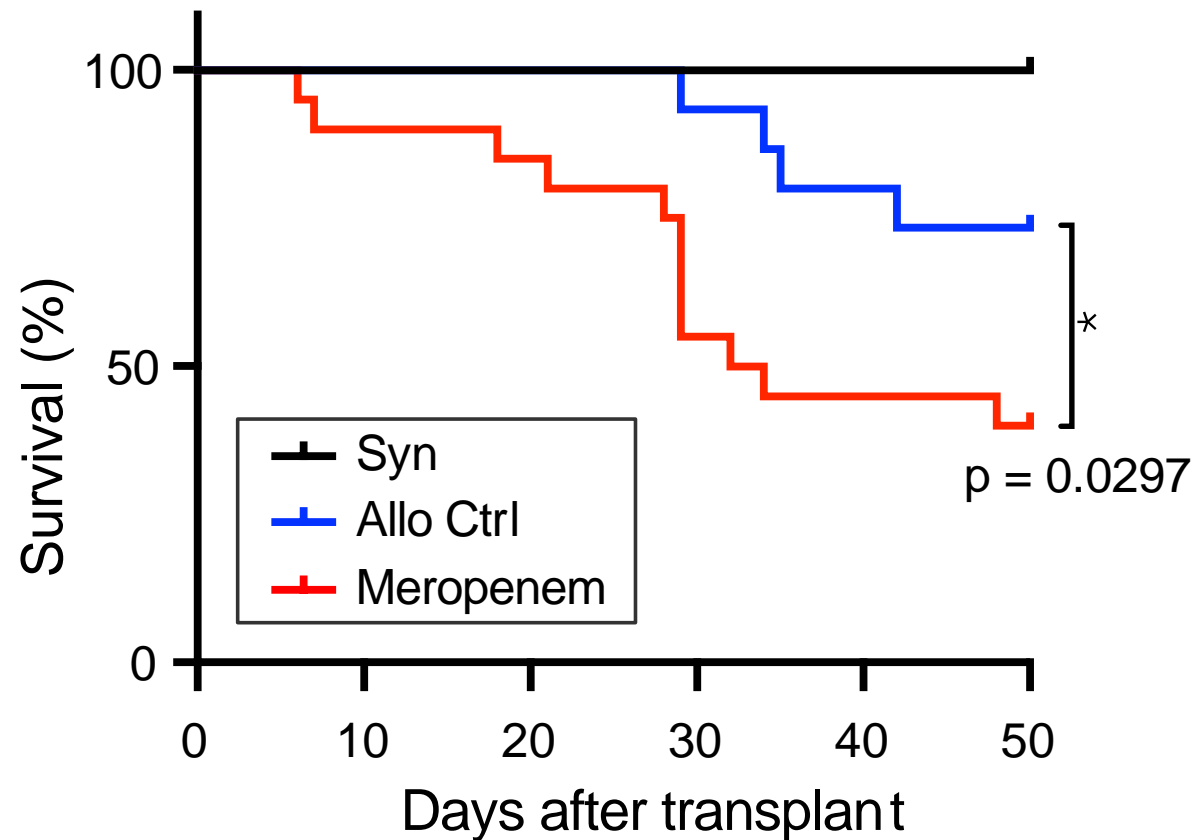
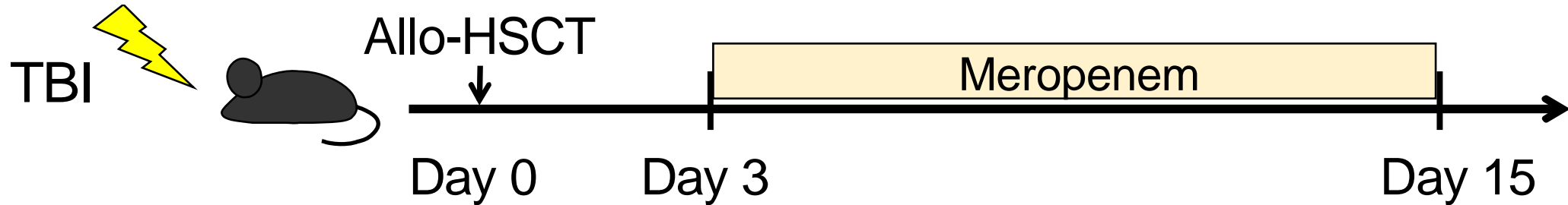
- No antibiotics
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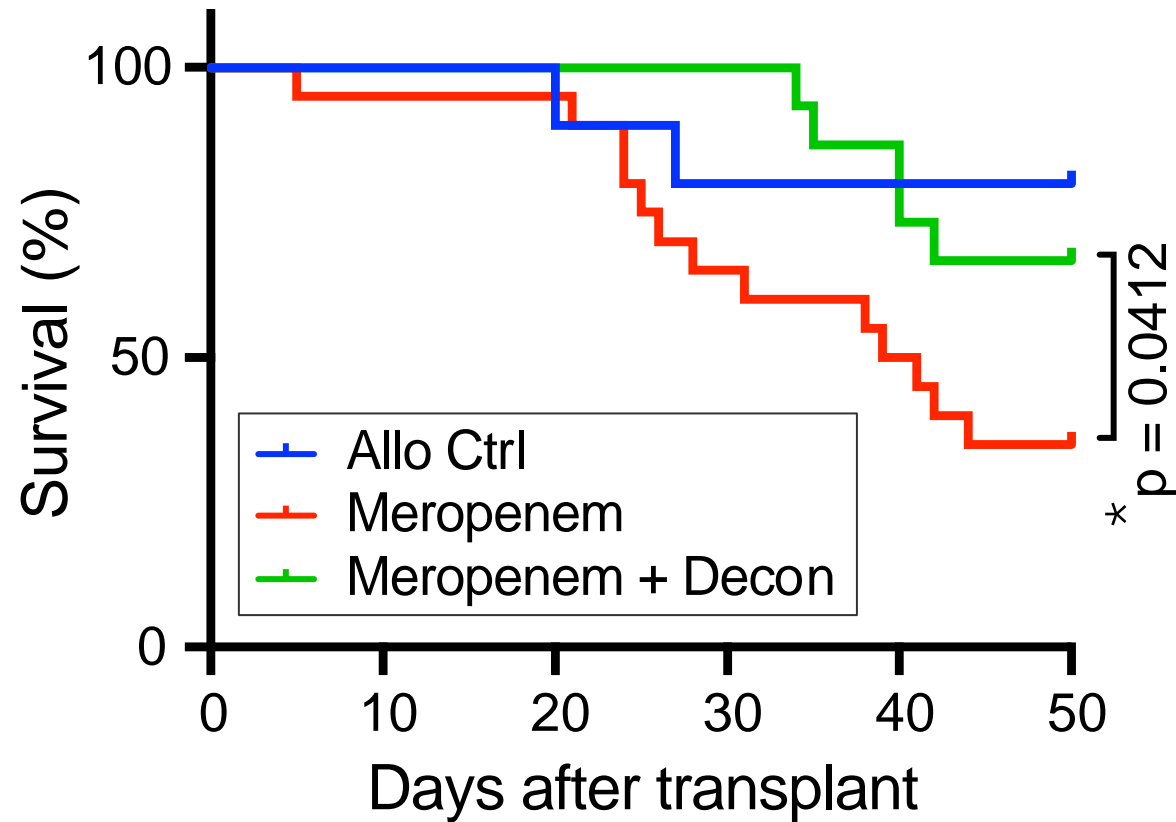
# Meropenem treatment is associated with aggravated intestinal GVHD



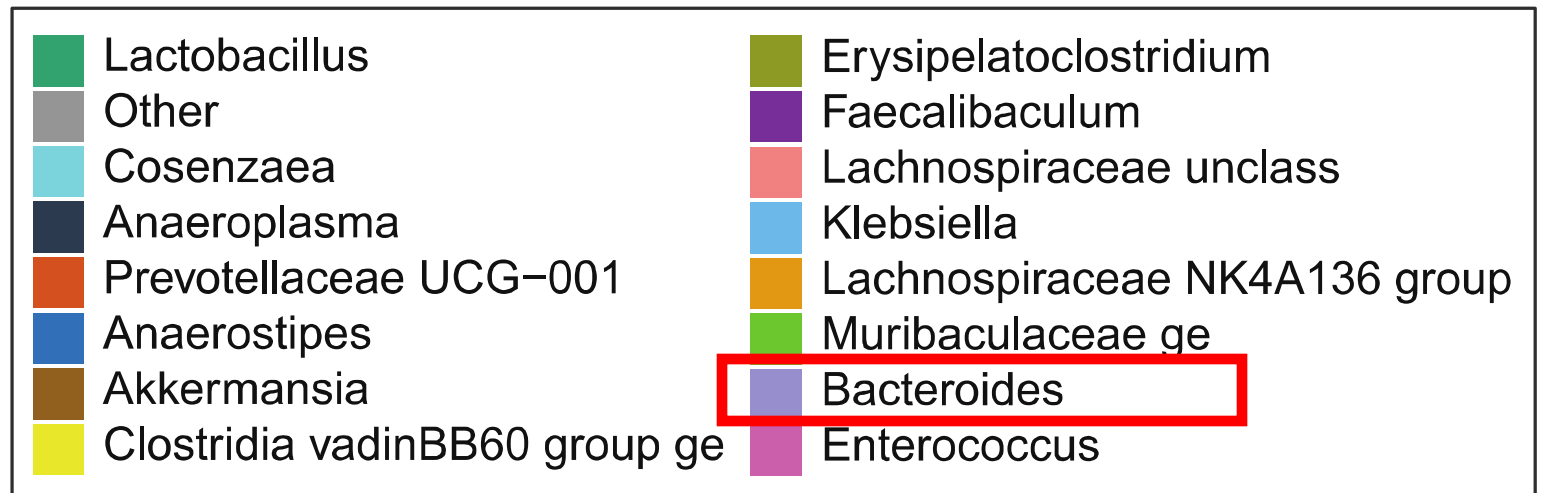
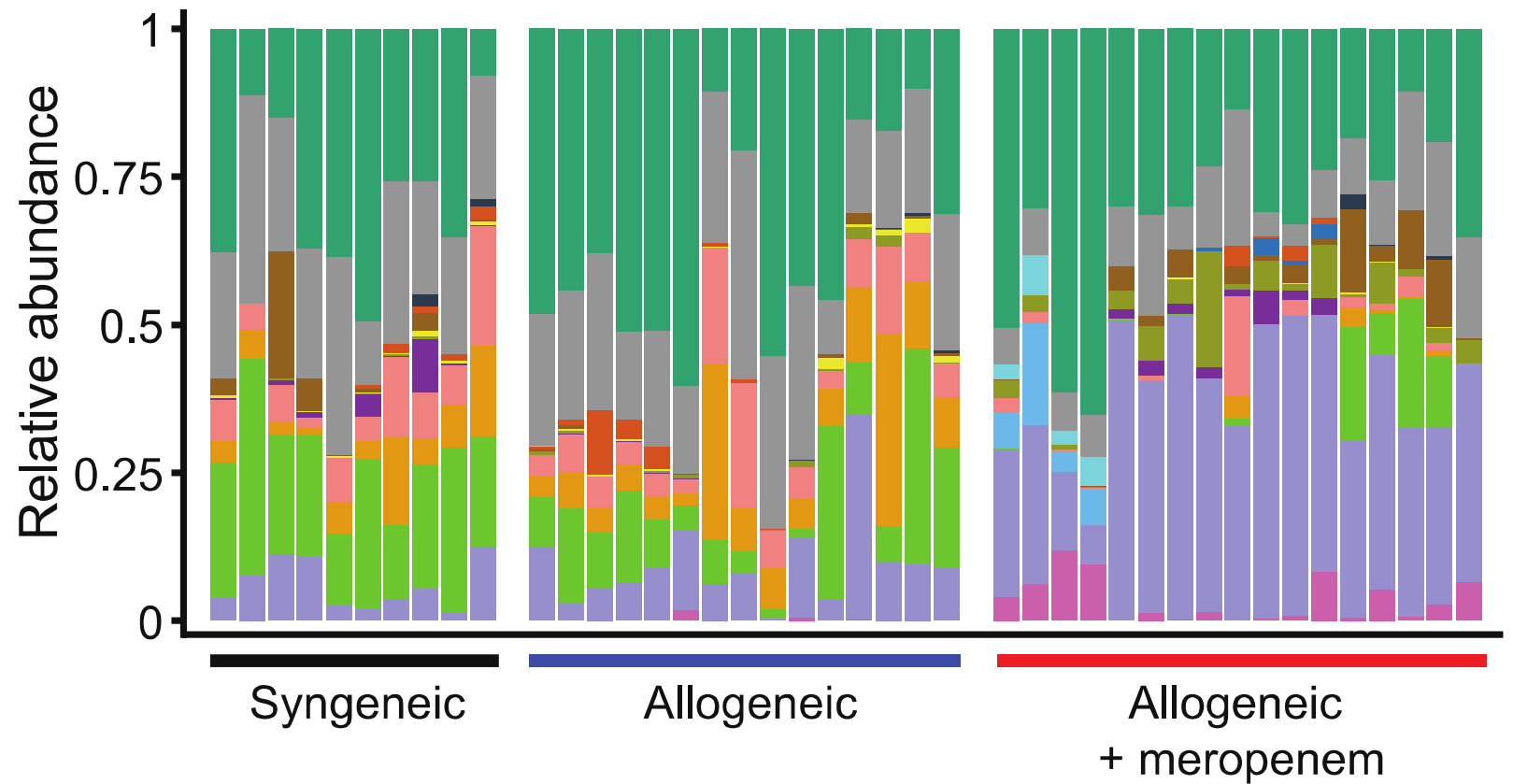
# Meropenem treatment is associated with aggravated intestinal GVHD



# Decontamination Can Rescue Meropenem-Treated GVHD mice

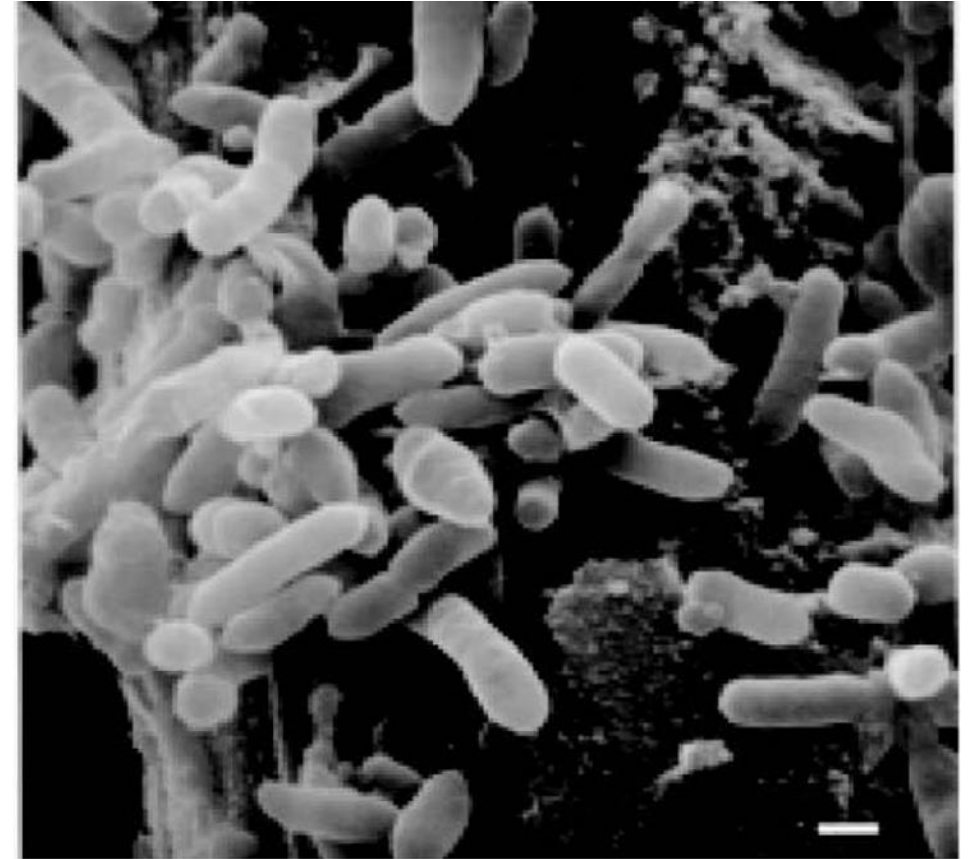


*Bacteroides* is increased in meropenem-treated GVHD mice



# *Bacteroides thetaiotaomicron*

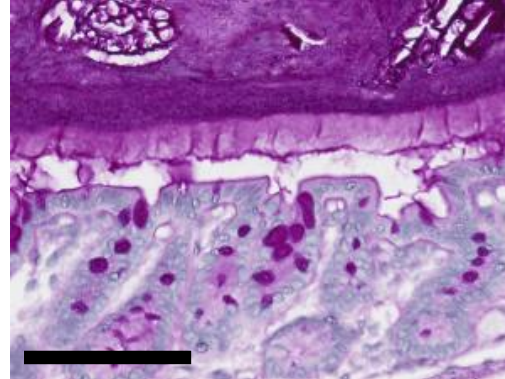
- An intestinal bacterial found in both mice and humans
- Has a broad ability to digest dietary fibers and intestinal mucus



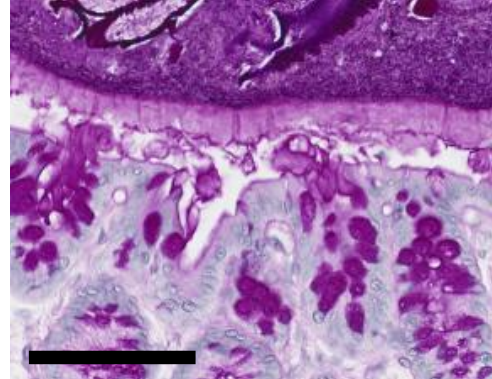
*TerAvest MA et al; Biotechnol Bioeng, 2014*

# Meropenem Arm Showed Thinned Mucus Layer Decontamination Suppressed Mucus Degradation

Syn



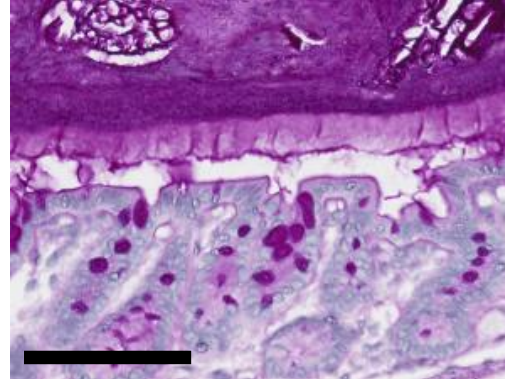
Allo Ctrl



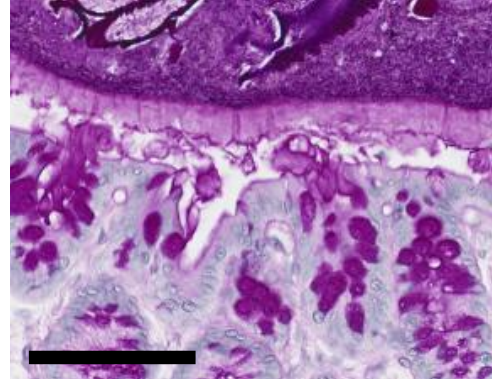


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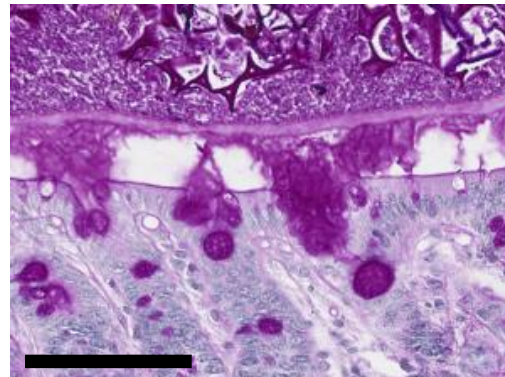
Syn



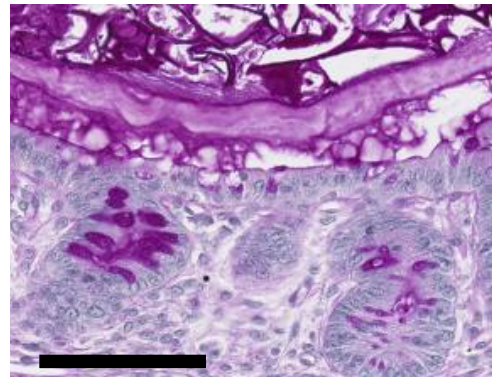
Allo Ctrl



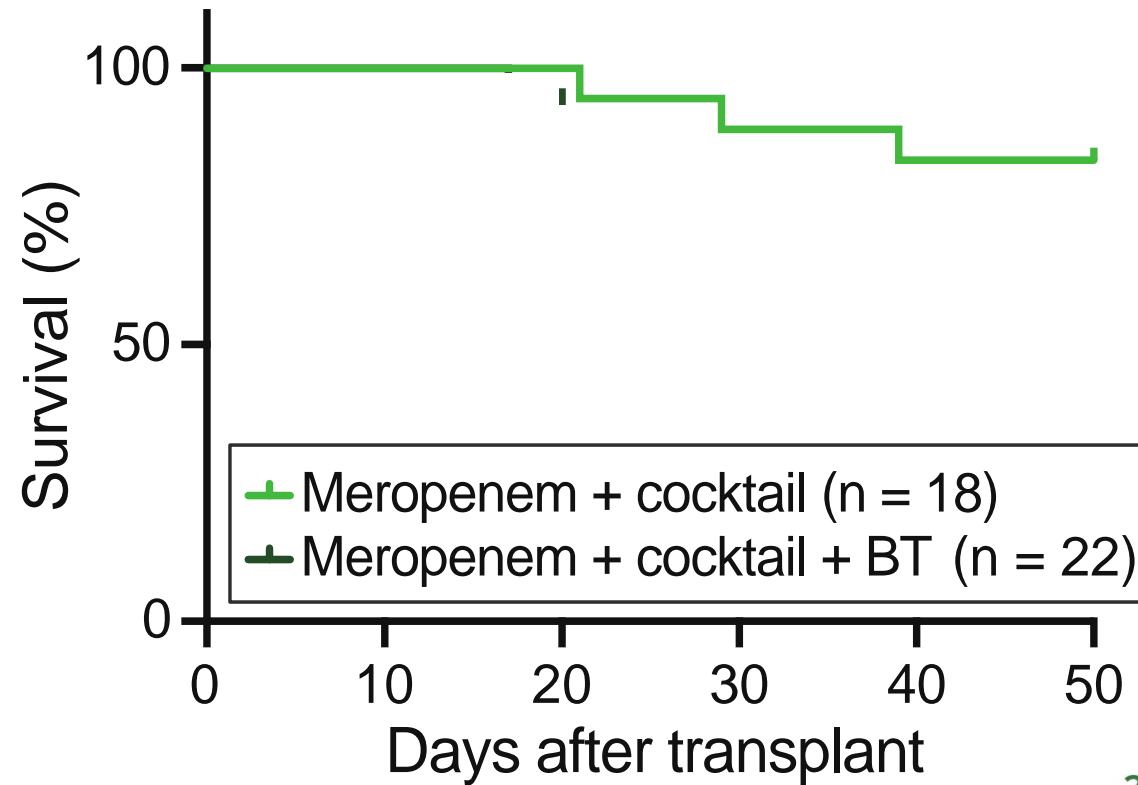
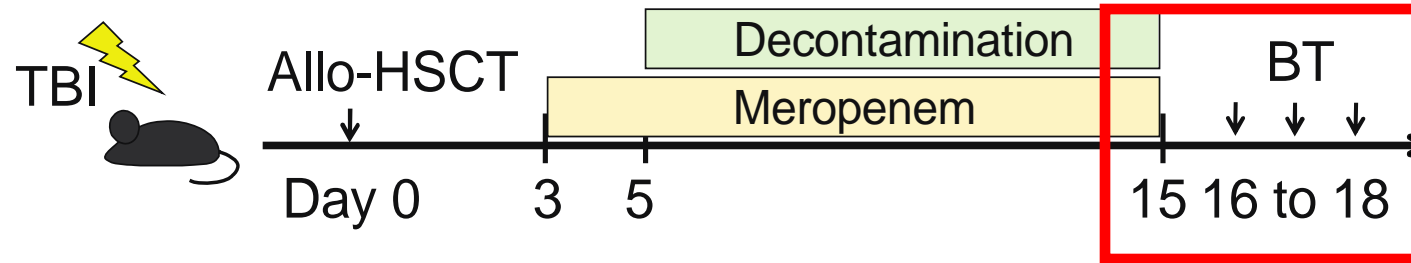
Meropenem



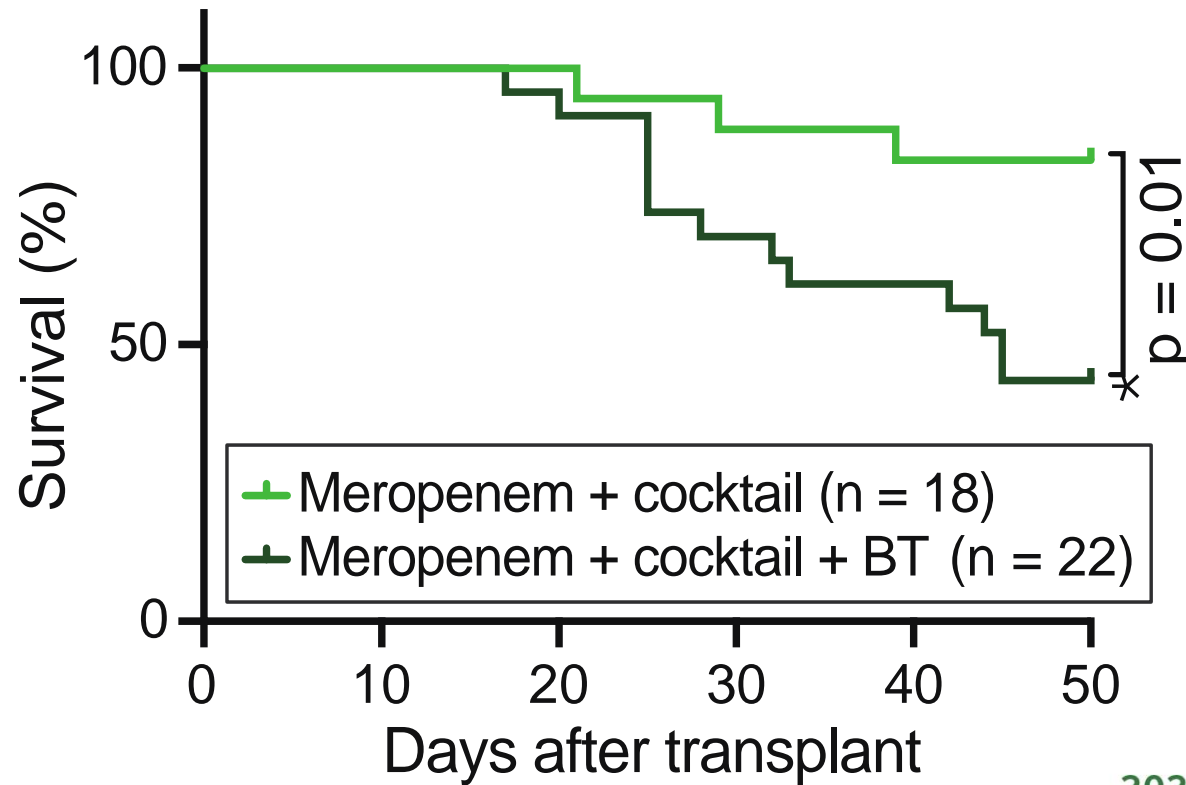
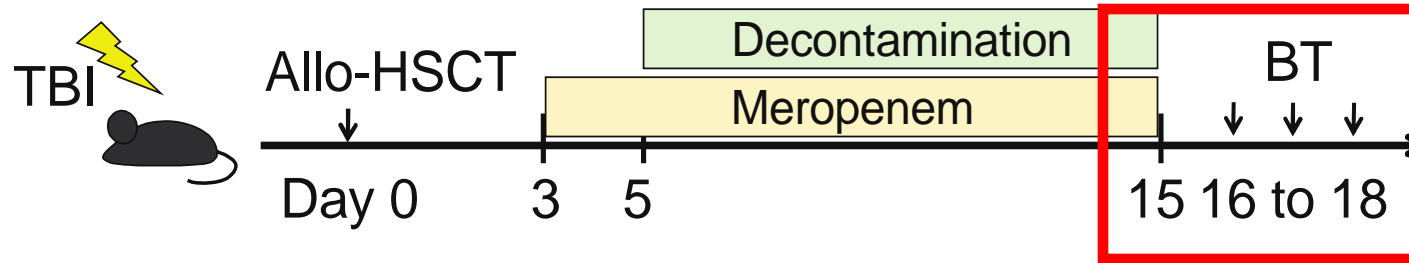
Meropenem + Decon



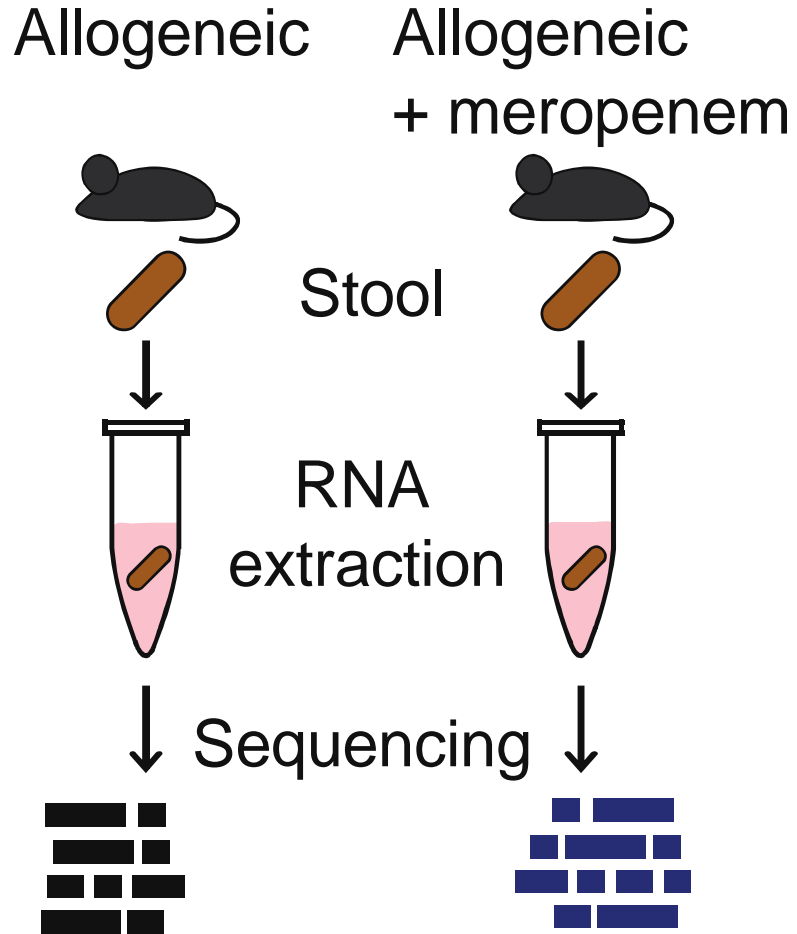
# *B. thetaiotaomicron* Reintroduction Aggravates GVHD in Decontaminated Mice



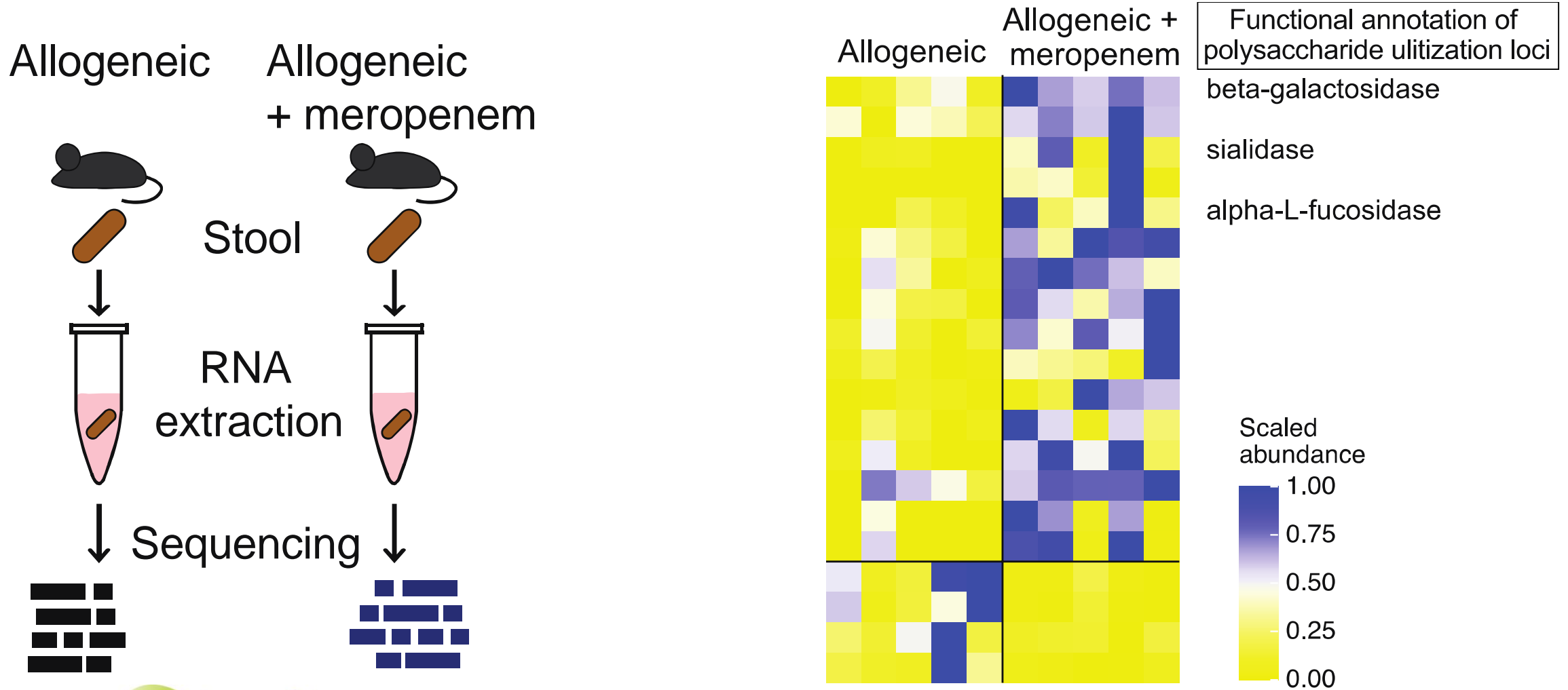
# *B. thetaiotaomicron* Reintroduction Aggravates GVHD in Decontaminated Mice



# Meropenem Upregulated the Expression of Mucus-Degrading Enzyme in *B. Thetaiotaomicron*



# Meropenem Upregulated the Expression of Mucus-Degrading Enzyme in *B. Thetaiotaomicron*



Next question:

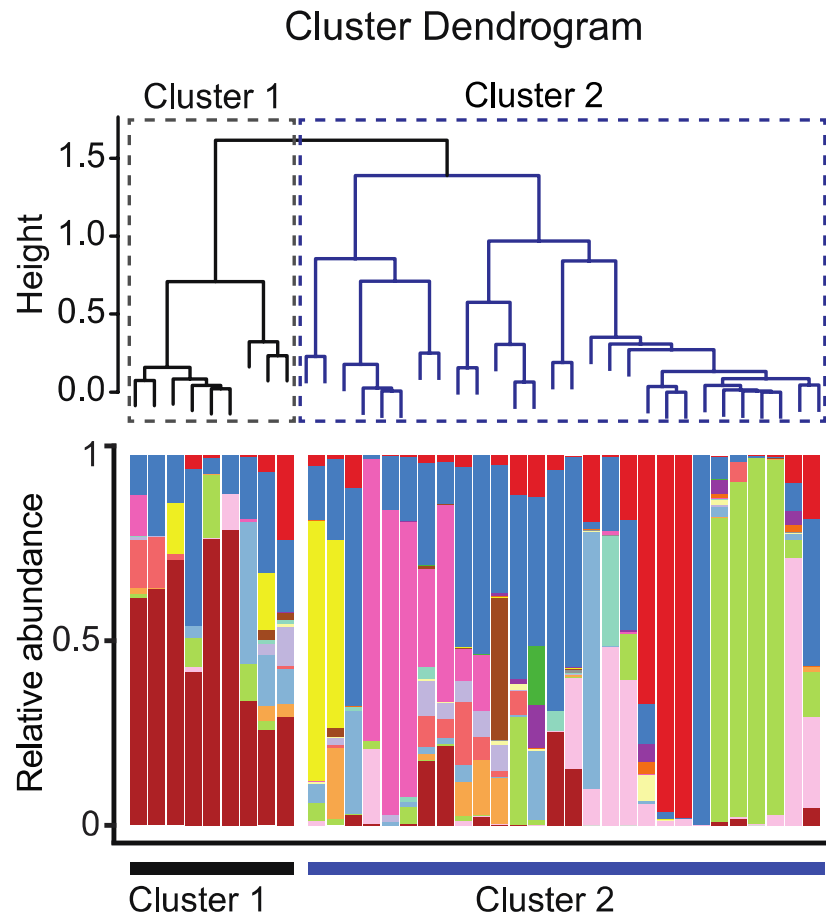
Does the microbiota impact  
GVHD response to treatment with  
corticosteroids?

# Evaluated Patients Presenting with New Lower GI GVHD (n=37)

THE UNIVERSITY OF TEXAS  
MD Anderson ~~Cancer~~ Center

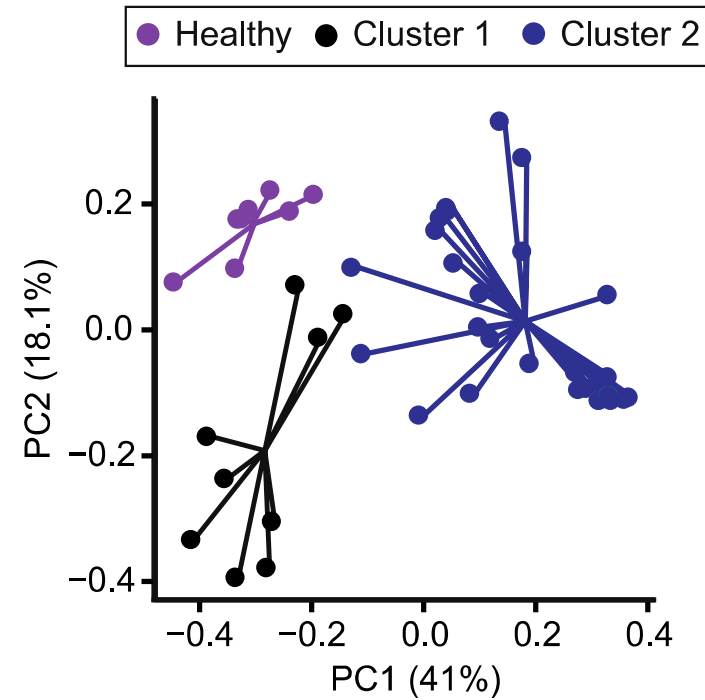
Making Cancer History®

# Patient Microbiome at GVHD Onset



## Genus

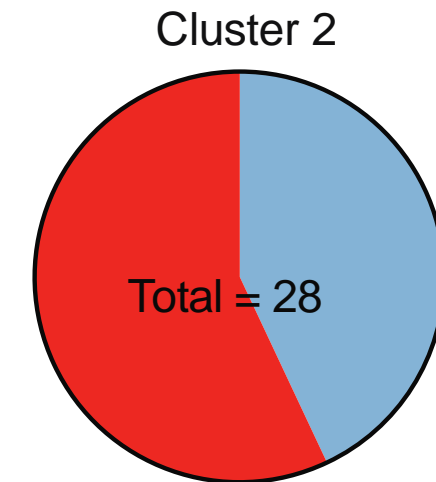
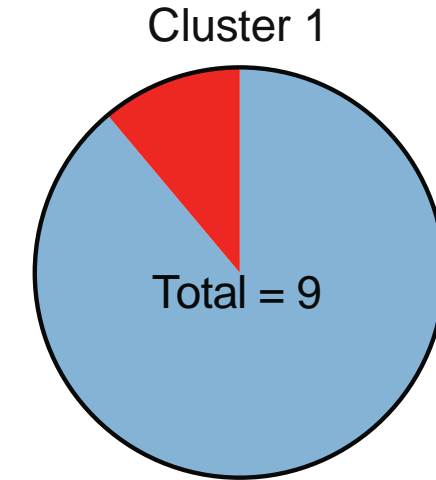
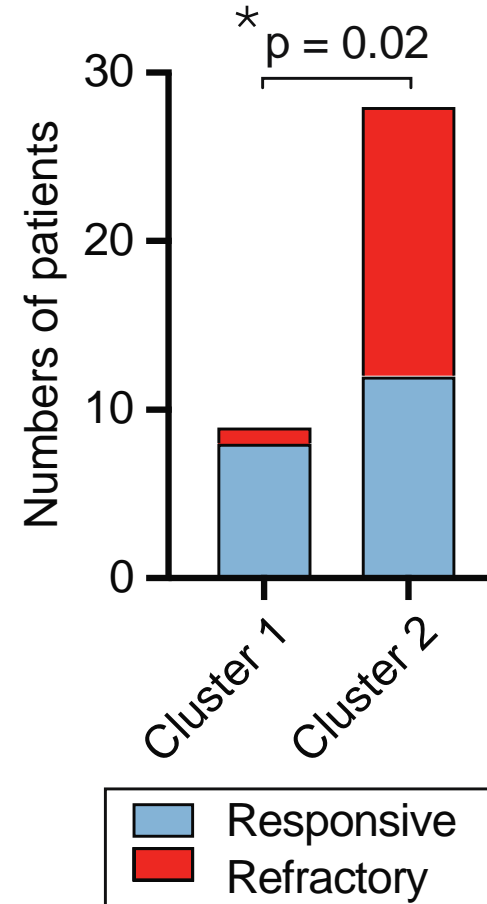
- *Streptococcus*
- *Other*
- *Butyricicoccus*
- *Granulicatella*
- *Rothia*
- *Akkermansia*
- *Anaerostipes*
- *Escherichia Shigella*
- *Muribaculaceae ge*
- *Bifidobacterium*
- *Actinomyces*
- *Lachnospiraceae ge*
- *Lachnoclostridium*
- *Veillonella*
- *Blautia*
- *Enterococcus*
- *Lactobacillus*
- *Bacteroides*



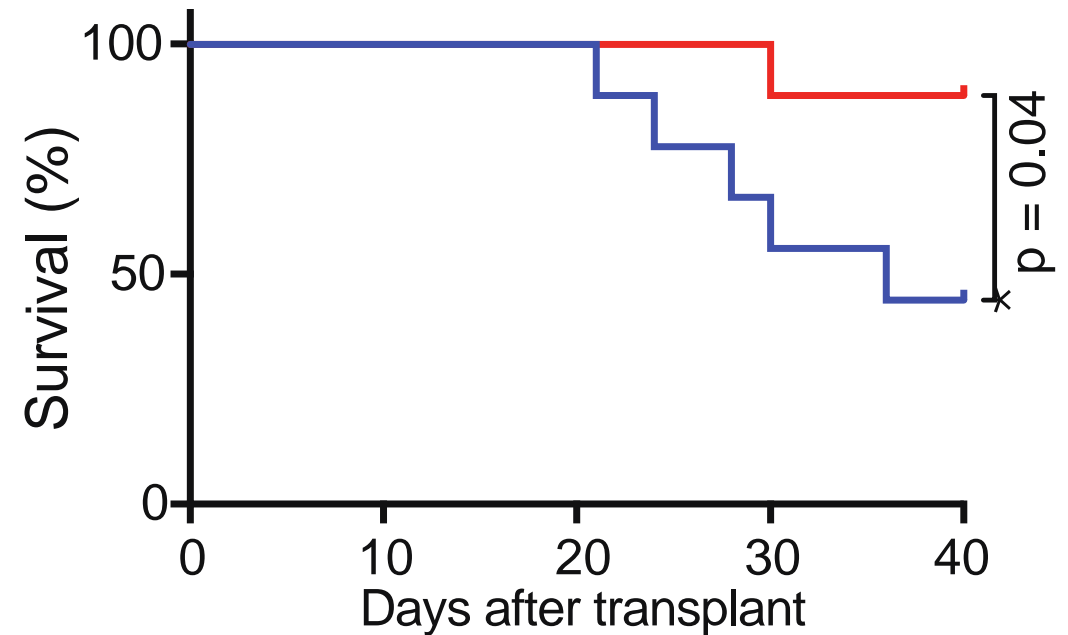
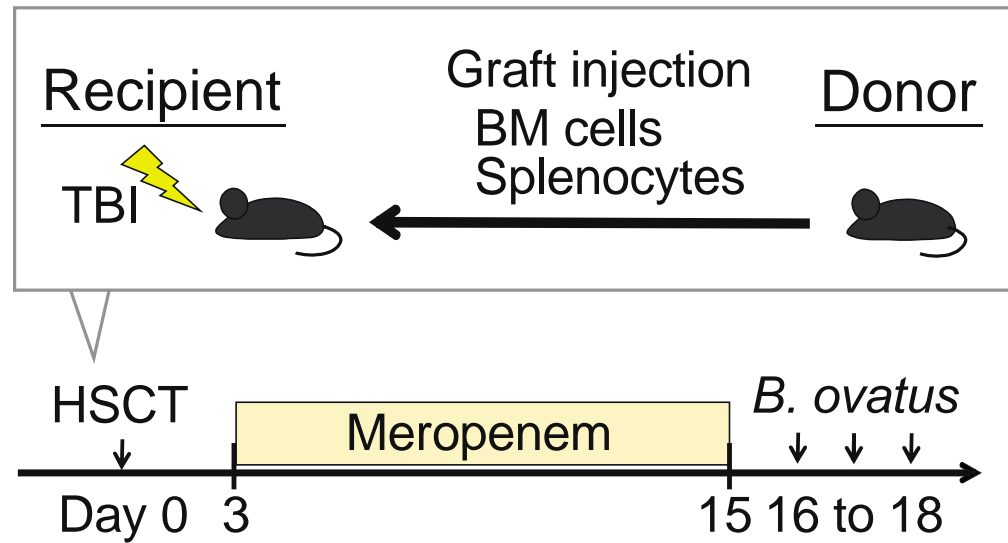


# Cluster 1 was Enriched in GVHD Treatment Responders

Treatment responsiveness

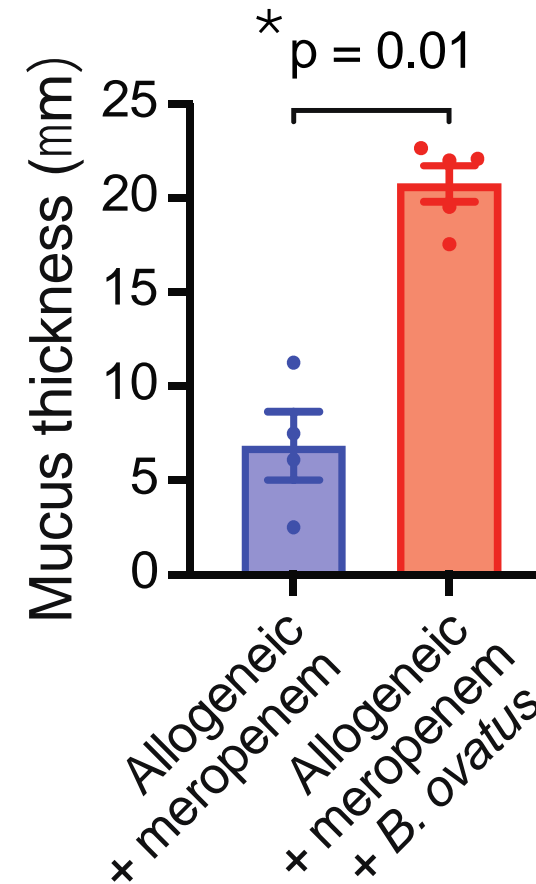
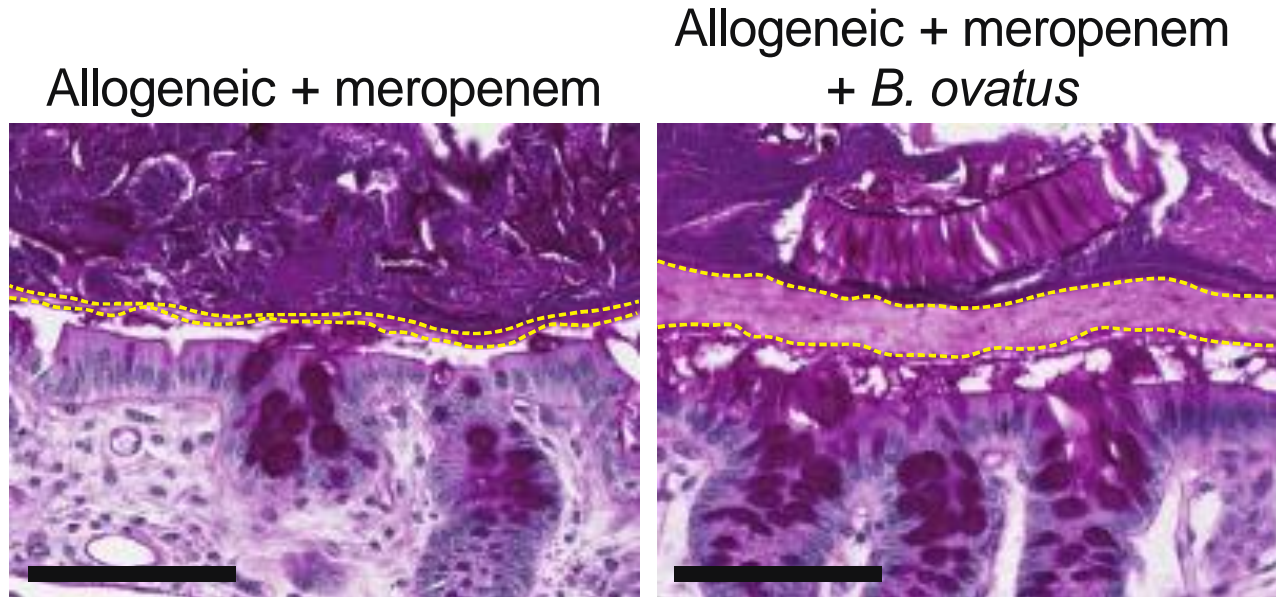


# Introduction of *B. Ovatus* Improved Survival in Meropenem-Induced GVHD Model

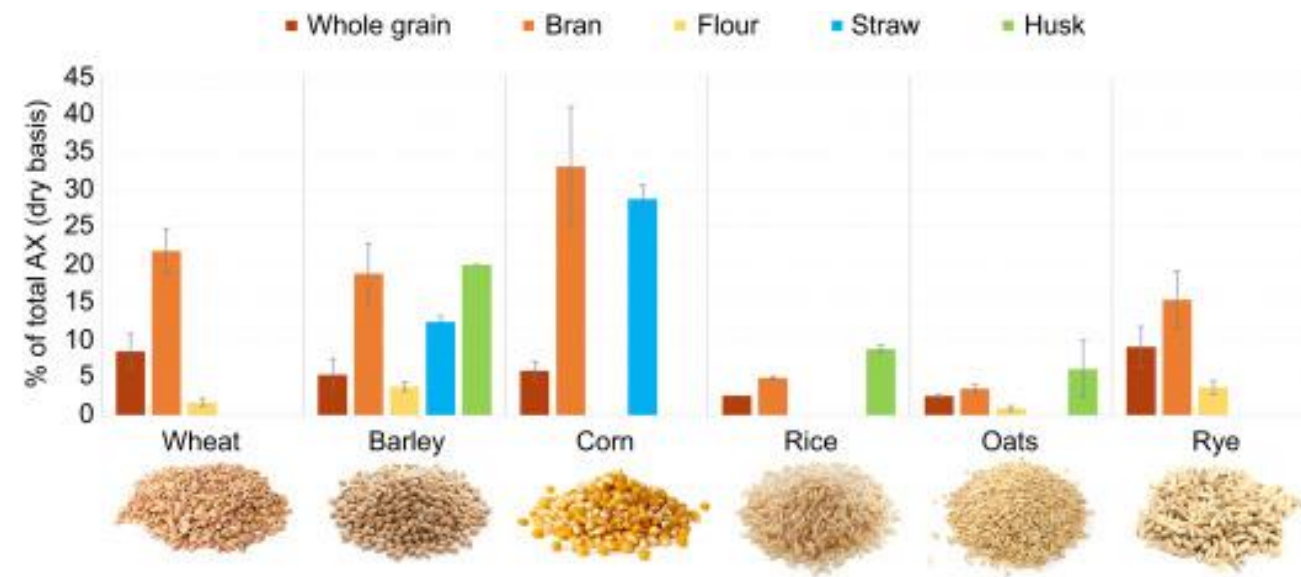
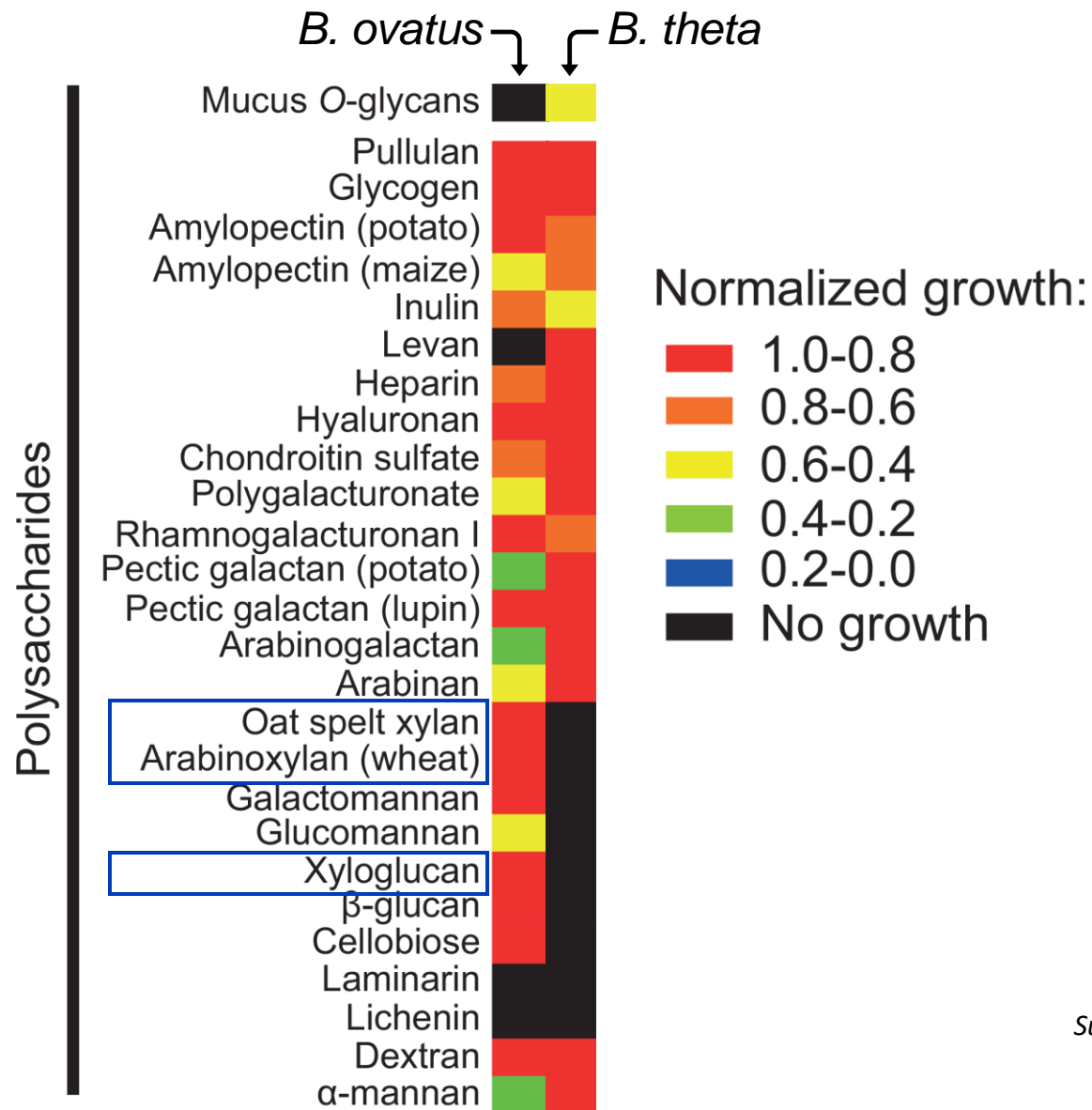


- + Allogeneic + meropenem (n = 9)
- + Allogeneic + meropenem + *B. ovatus* (n = 9)

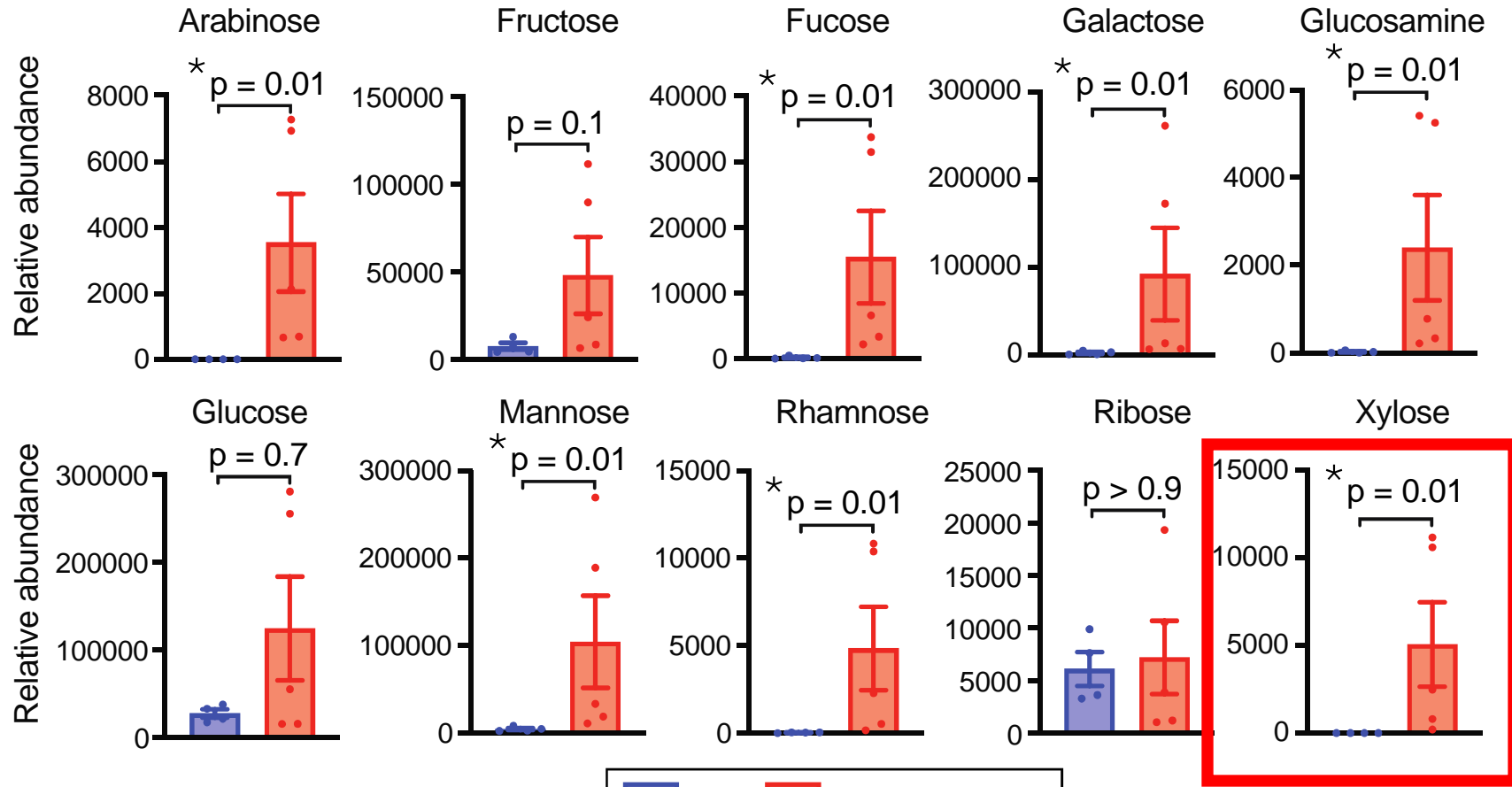
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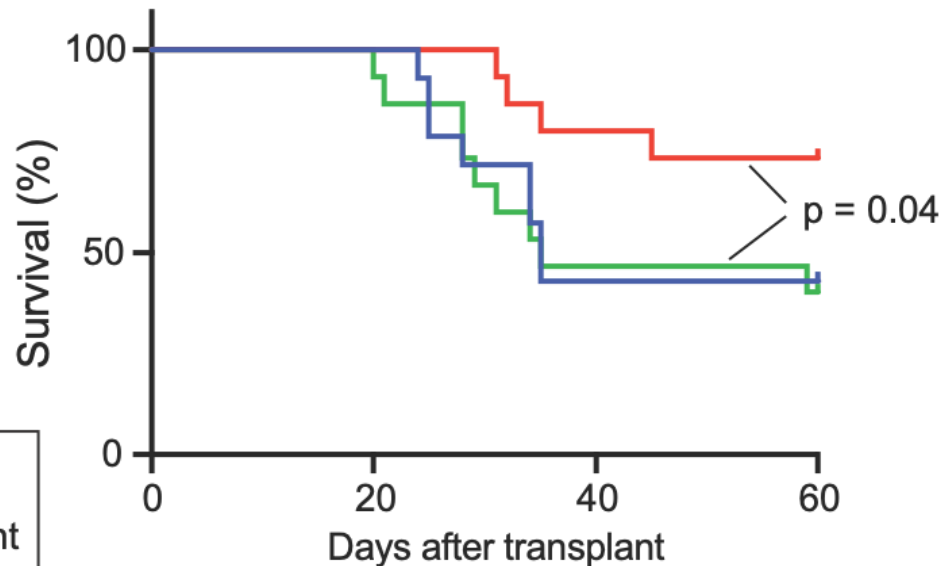
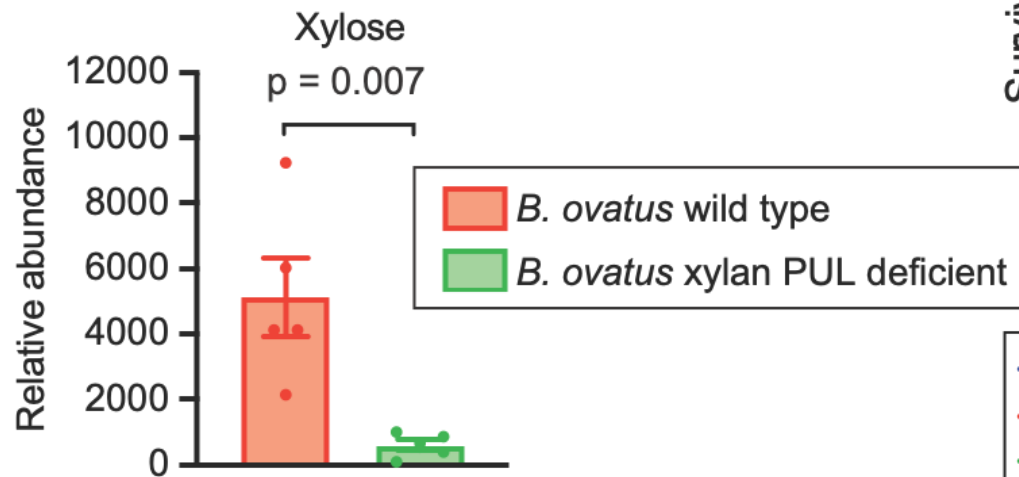
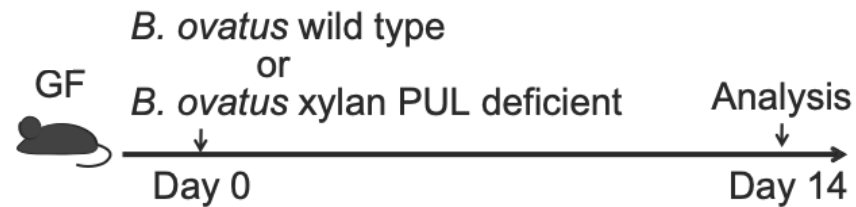
# *B. ovatus* and *B. theta* – A Tale of Two *Bacteroides*



# *B. ovatus* Introduction Increases Intestinal Sugar Concentrations



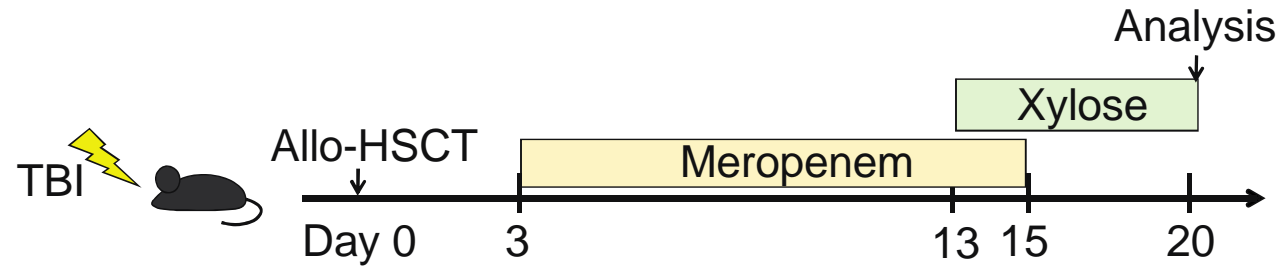
# An engineered mutant version of *B. ovatus* (xylanase insufficient) fails to rescue mice with GVHD after meropenem



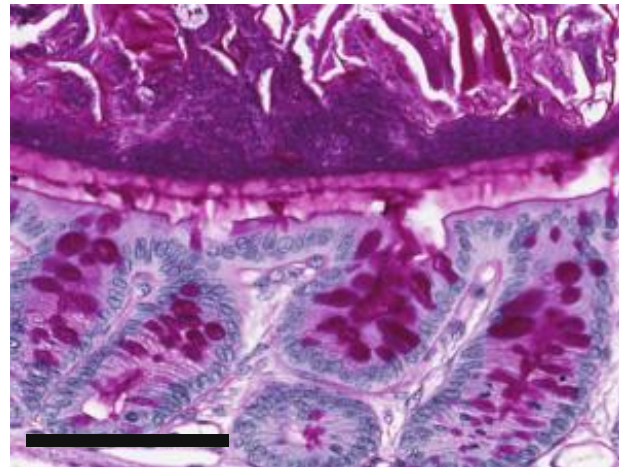
- Allogeneic + meropenem (n = 14)
- Allogeneic + meropenem + *B. ovatus* wild type (n = 15)
- Allogeneic + meropenem + *B. ovatus* xylan PUL deficient (n = 15)

courtesy of Eric Martens, U. Michigan

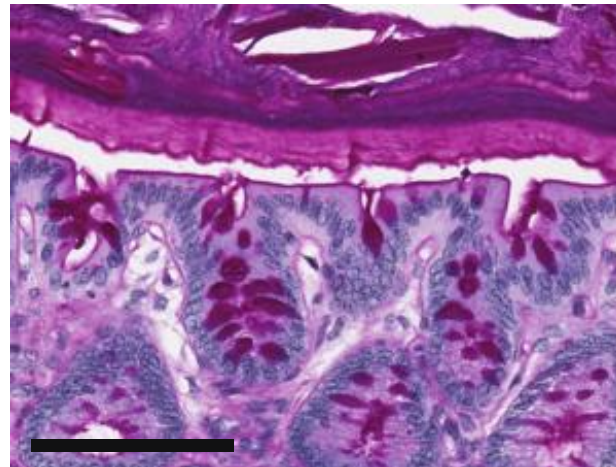
# Supplementation of Xylose Resulted in a Significantly Thicker Mucus Layer



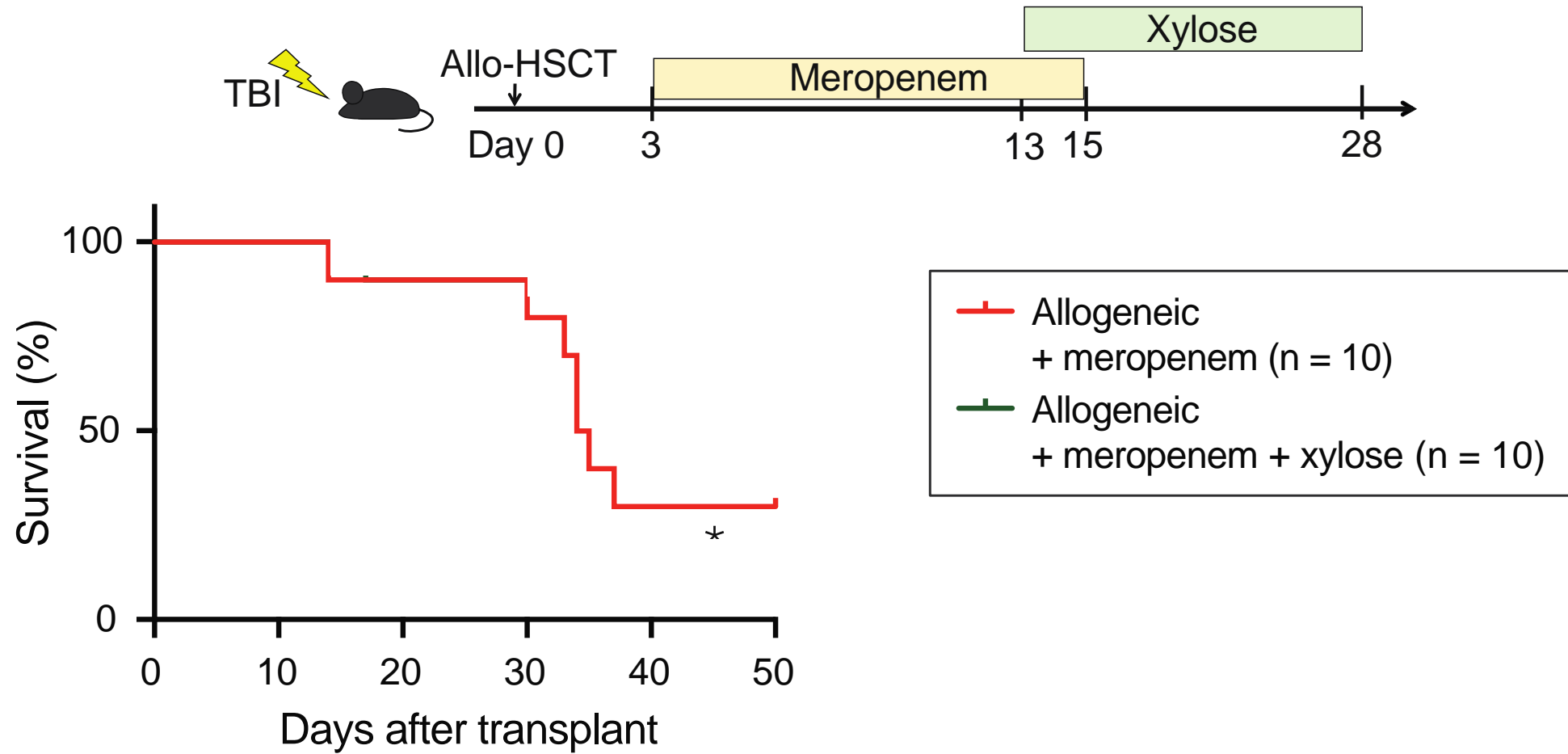
Allogeneic + meropenem



Allogeneic + meropenem + xylose

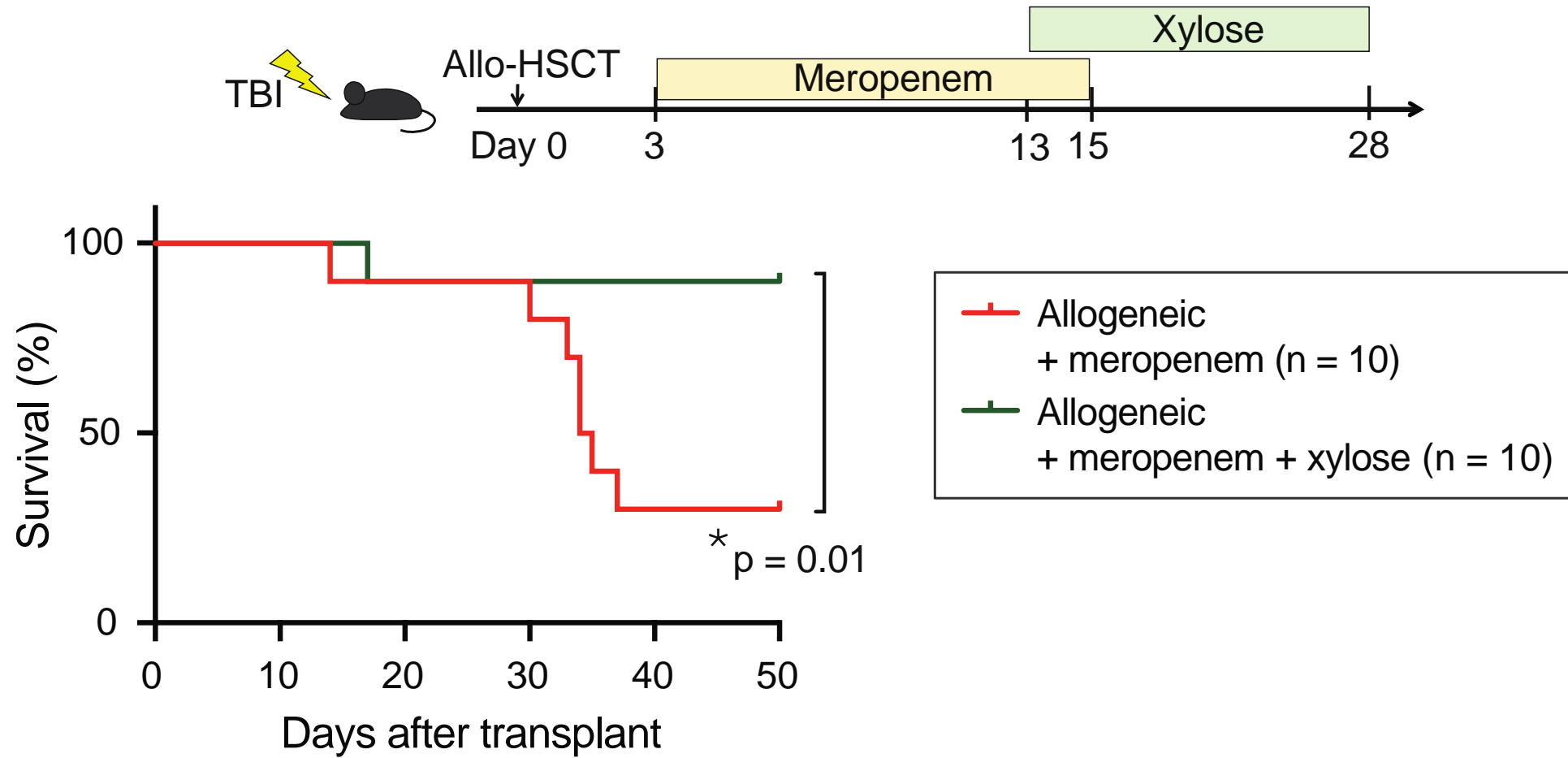


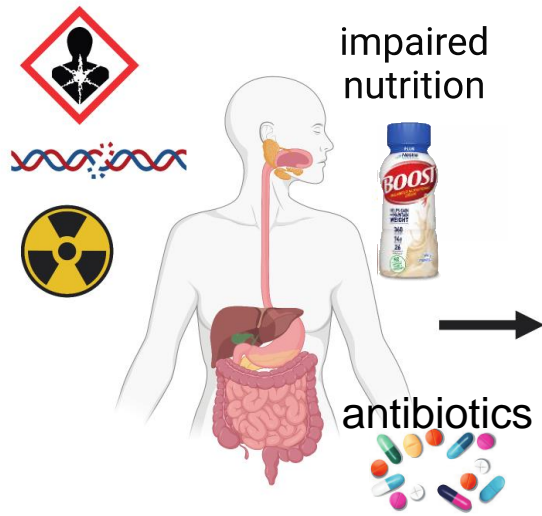
# Supplementation of Xylose Improved Survival in Meropenem-Treated Mice



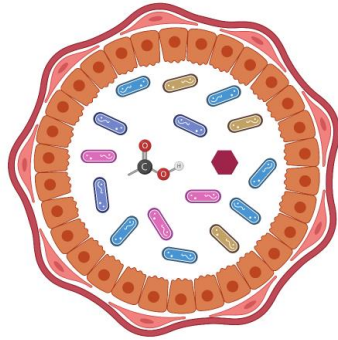


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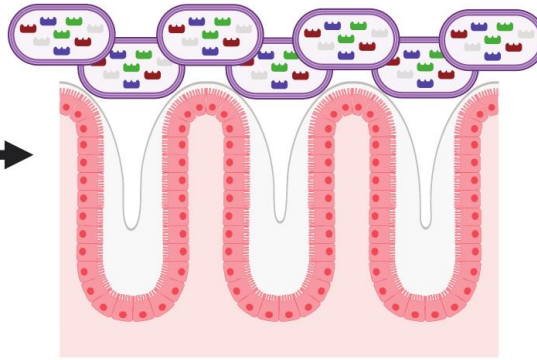


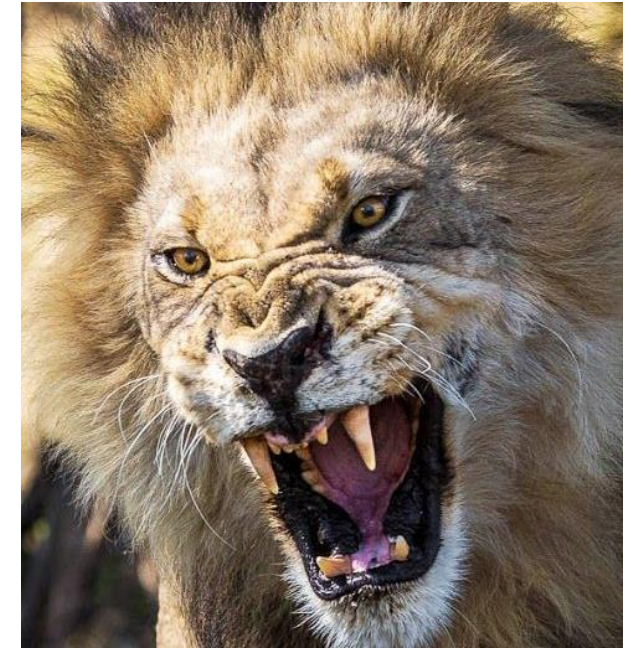
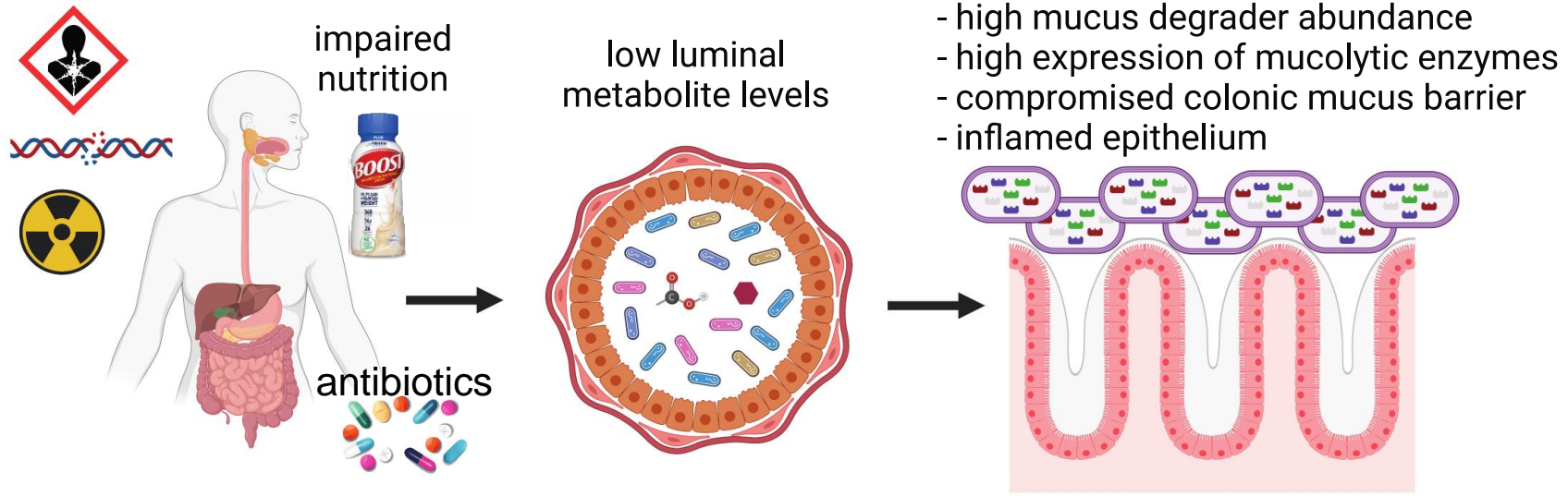


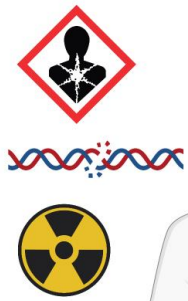
low luminal  
metabolite levels



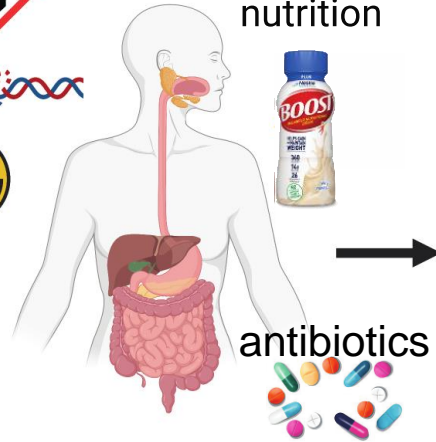
- high mucus degrader abundance
- high expression of mucolytic enzymes
- compromised colonic mucus barrier
- inflamed epithelium





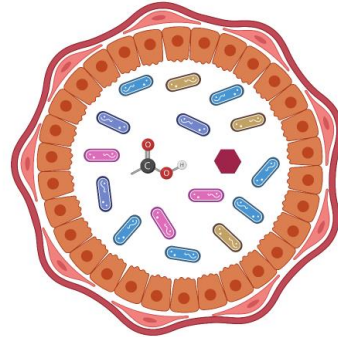


impaired nutrition

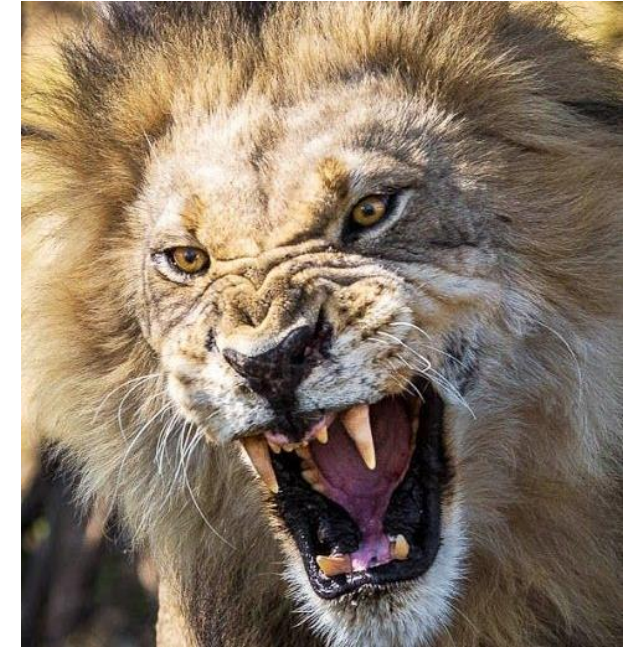
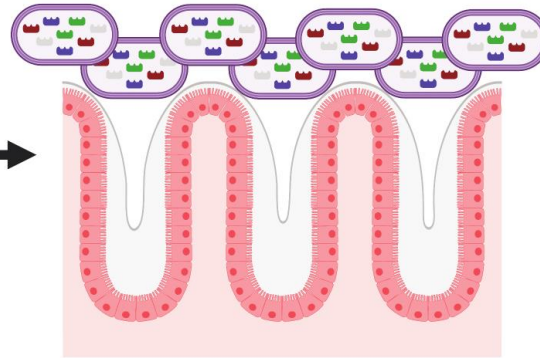


antibiotics

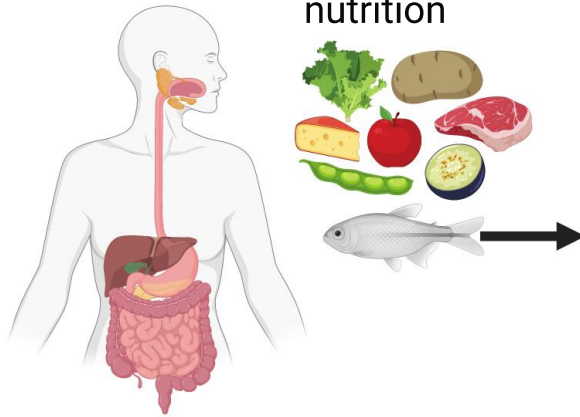
low luminal metabolite levels



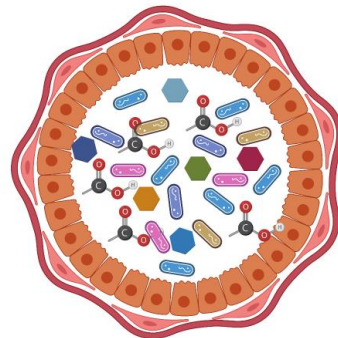
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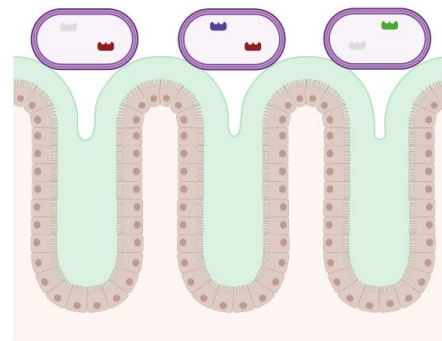
normal nutrition

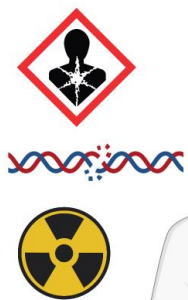


high luminal metabolite levels

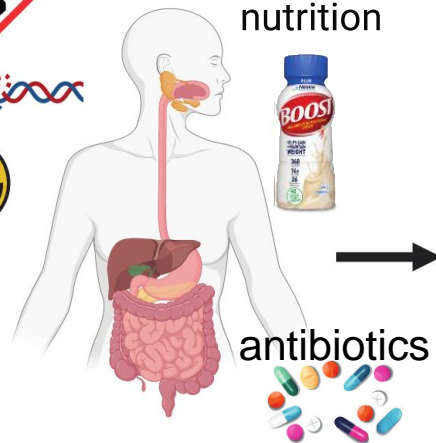


- low mucus degrader abundance
- low expression of mucolytic enzymes
- intact colonic mucus barrier
- healthy epithelium



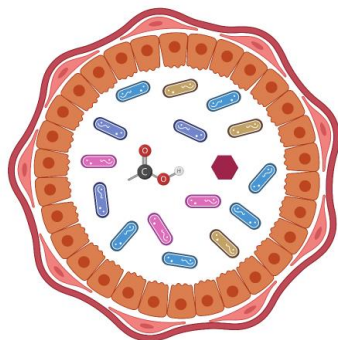


impaired nutrition

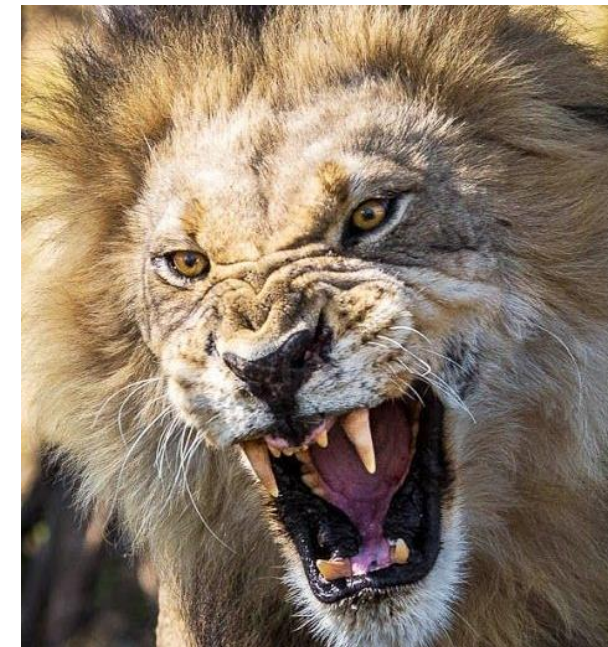
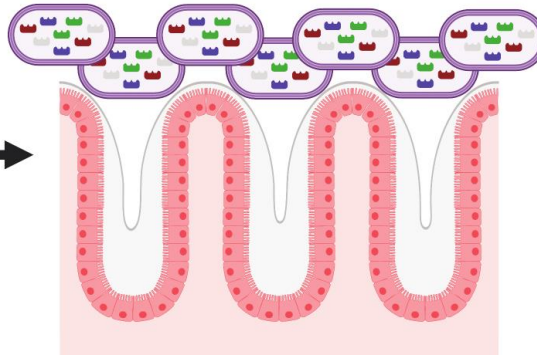


antibiotics

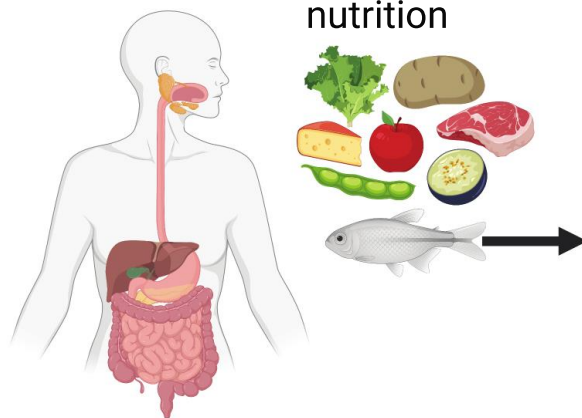
low luminal metabolite levels



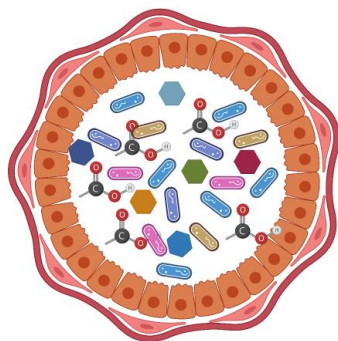
- high mucus degrader abundance
- high expression of mucolytic enzymes
- compromised colonic mucus barrier
- inflamed epithelium



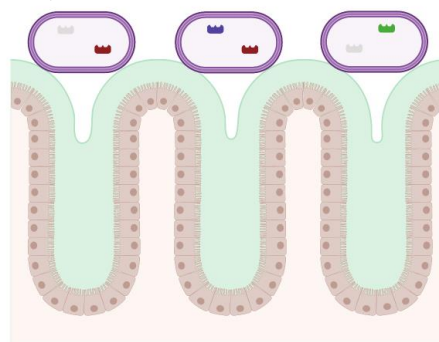
normal nutrition



high luminal metabolite levels



- low mucus degrader abundance
- low expression of mucolytic enzymes
- intact colonic mucus barrier
- healthy epithelium



# Lessons Learned

- Antibiotics are important tools for treating infections but can have side effects, including injuring the microbiome
- Diet and nutrition are important for supporting the microbiome, especially consuming a variety of different fruits and vegetables on a regular basis
- Probiotics that are currently available in pharmacies and health food stores are probably not beneficial in restoring the microbiome – they don't include the right beneficial bacteria
- Potentially more effective strategies are in clinical trials, including fecal transplantation and mixtures of cultured bacteria

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# Acknowledgements

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# Questions?



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