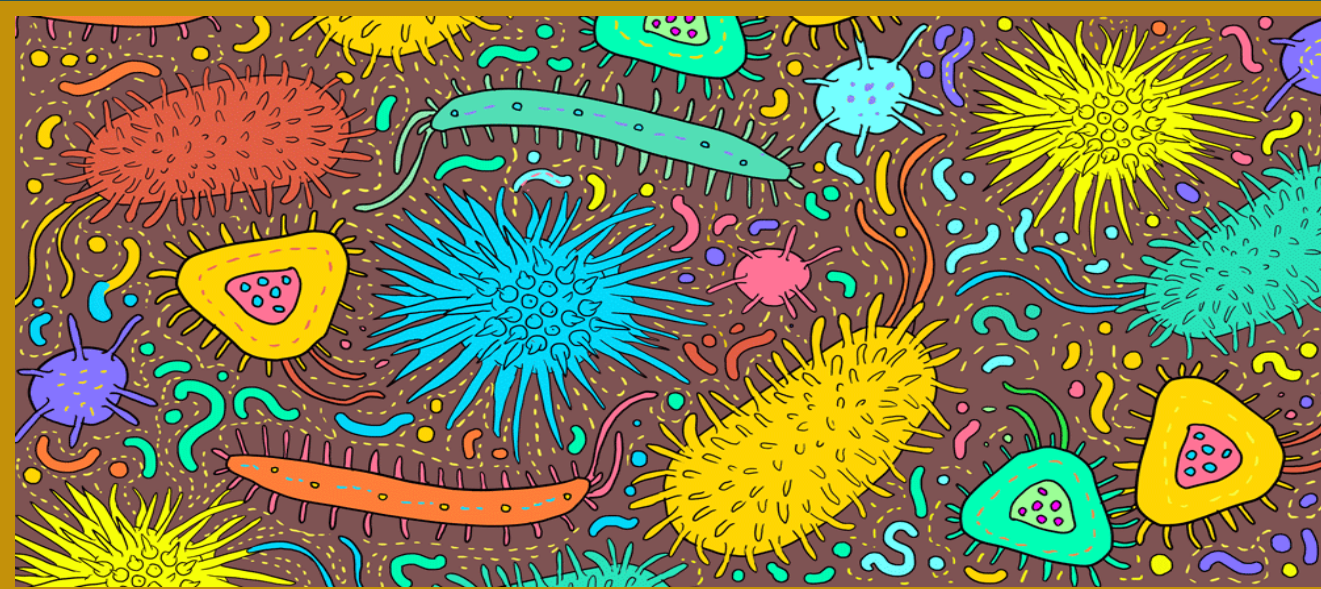


ECHO Discovery Series
Sept 11, 2024



“The early life microbiome and disrupting exposures”

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Henry Rutgers Professor of Microbiome and Health
CIFAR Fellow

Department of Biochemistry and Microbiology,
Department of Anthropology
Institute for Food, Nutrition and Health
Rutgers University, USA



Converging life-styles

Antropocene

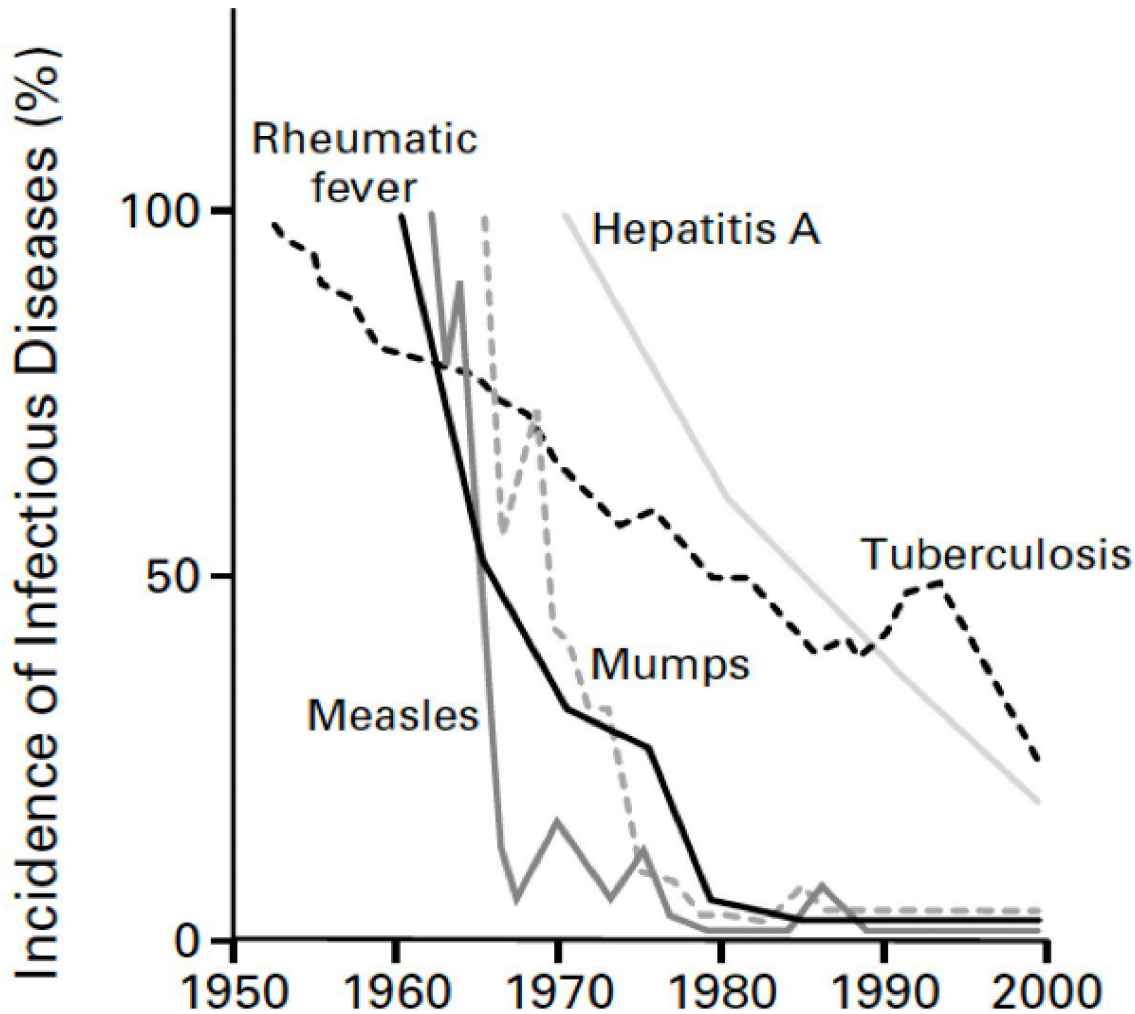
Cultural diversity



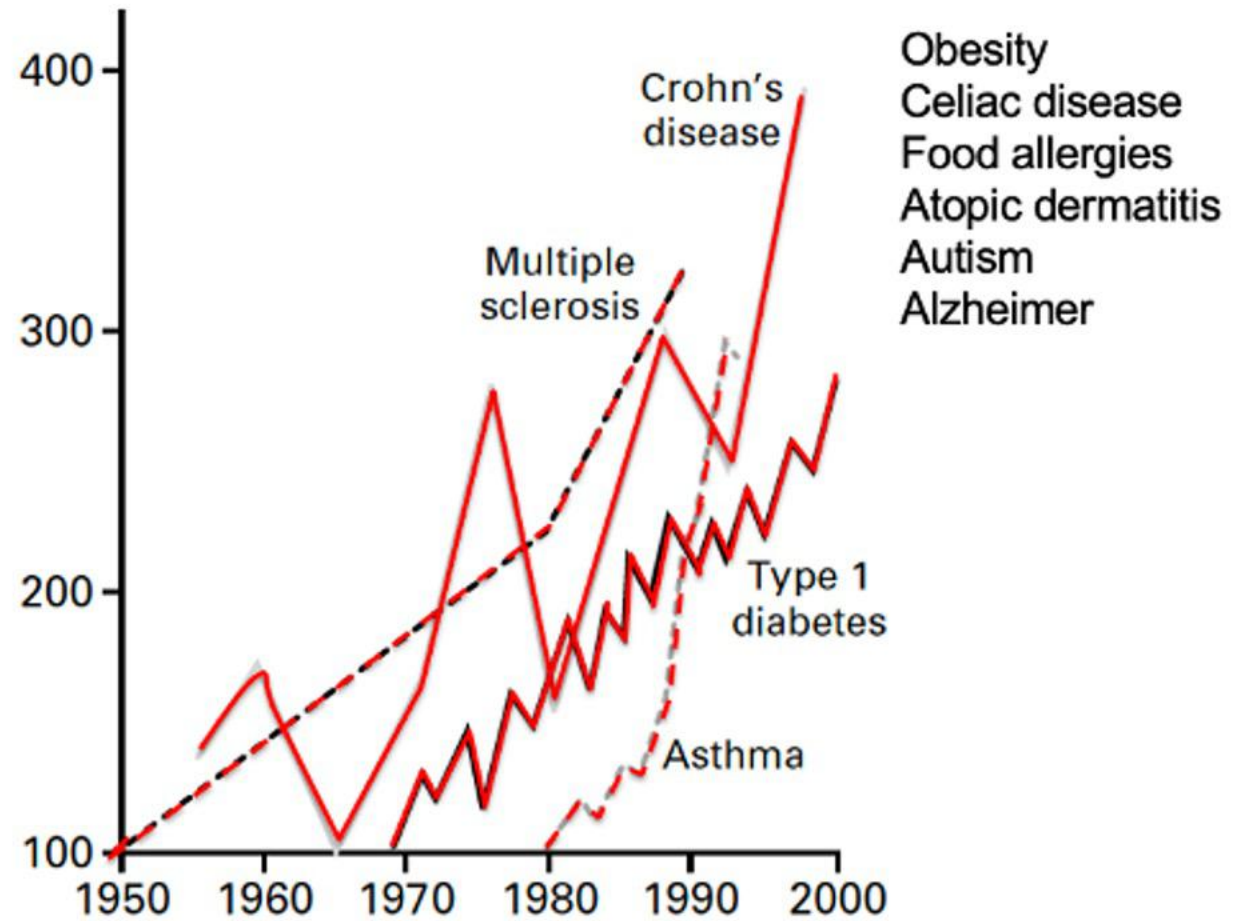
Social practices
Food production practices
Diet
Medicine



Disease trade of the 20th century



Bach 2001, NEJM



Bach et al. 2001, NEJM

Hypotheses for immune/metabolic diseases

Disappearing microbiota hypothesis



HUMAN-ADAPTED MICROBES
→ regulatory pathways in early life

Blaser 2006 EMBO Rep
Rook 2010 Clin Exp Immunol
von Hertzen et al. 2011 EMBO Rep.
Round et al. 2011 Science
Grainger, et al. 2010 J Exp Med
Atarashi et al. 2011 Science

Hygiene hypothesis



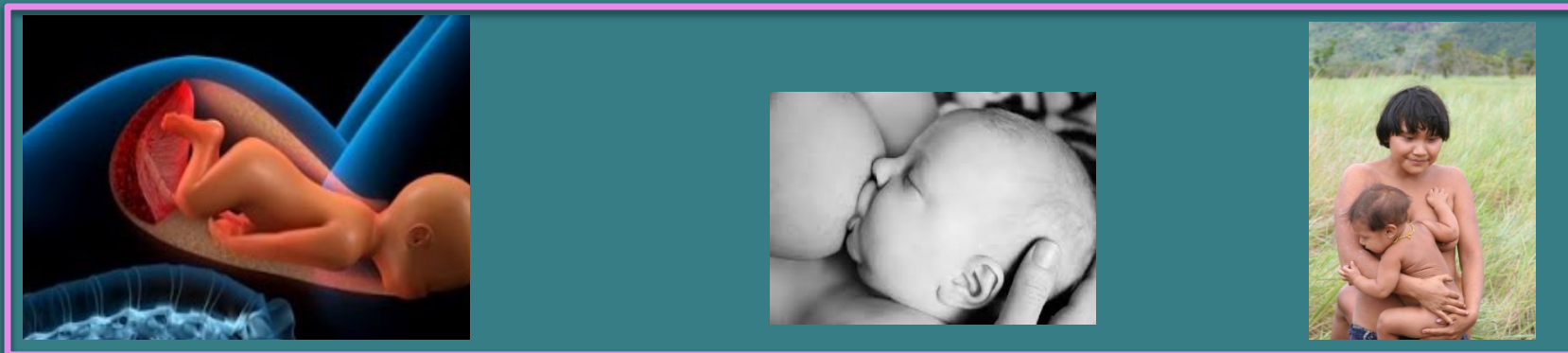
ENVIRONMENTAL MICROBES
< childhood asthma/allergies

Strachan 1989 Brit Med J
Rook 2009 Immunology
von Ehrenstein et al. 2000 Clin Exp Allergy
Riedler et al. 2001 Lancet
von Mutius, Vercelli, 2010 Nat Rev Immunol
Ownby et al. 2002 JAMA
Hanski I et al. 2012 PNAS

Natural microbial exposures

Human microbiome transmission

Environmental exposures



0-6 months

>6 months



Newborn



1 month



2 months



4 months

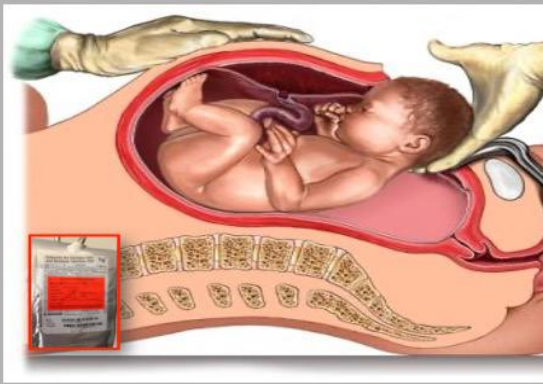


7 months

Microbiome stressors

Human microbiome transmission

Environmental exposures



0-6 months

>6 months



Newborn



1 month



2 months



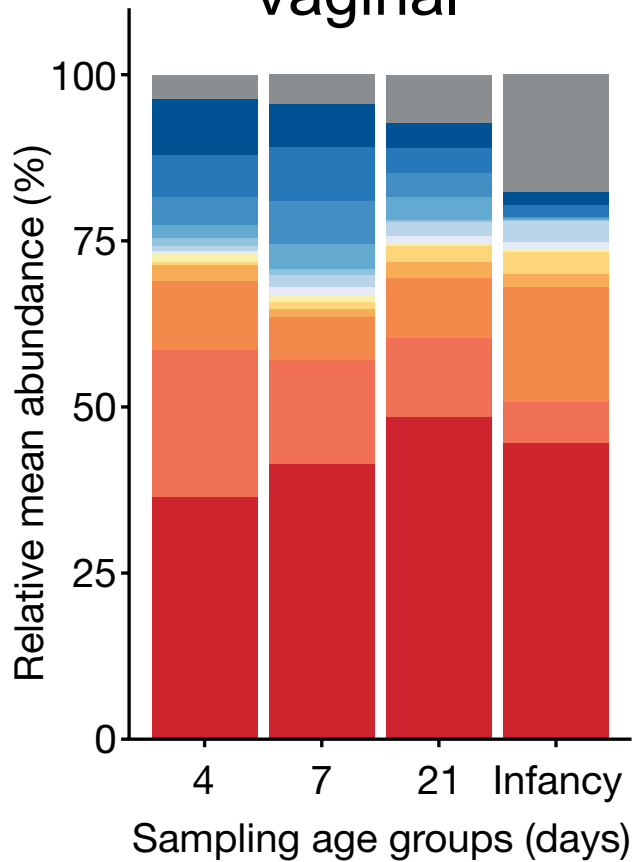
4 months



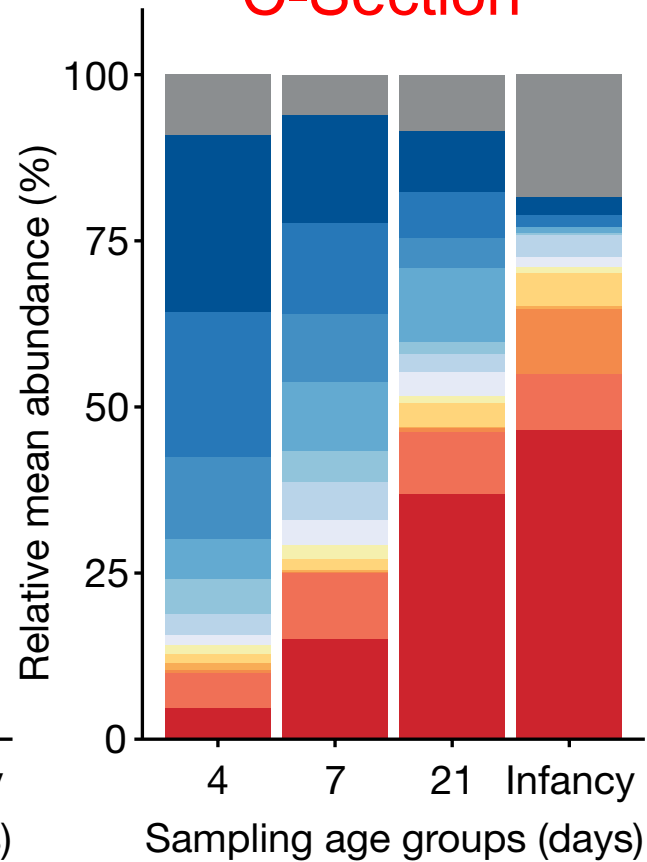
7 months

C-section alters the microbiome

Vaginal



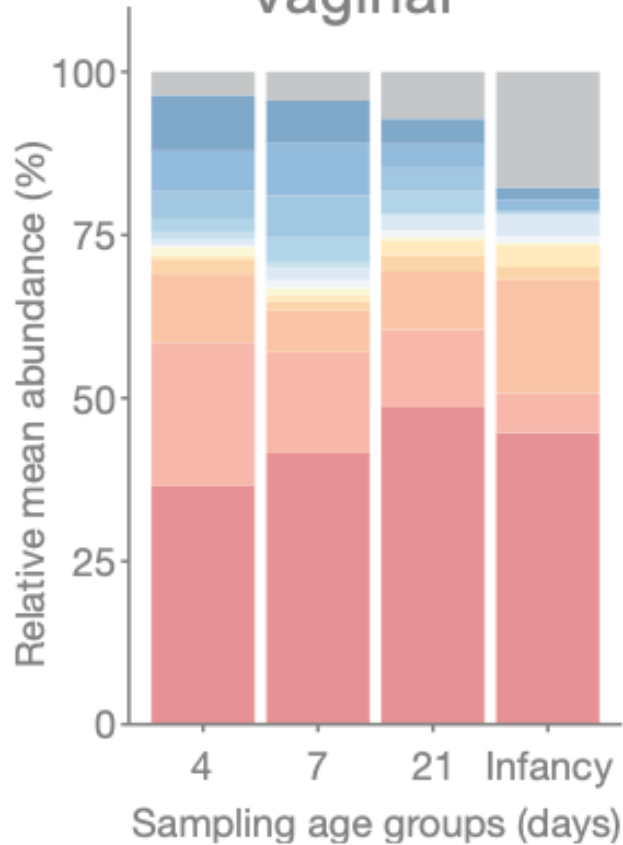
C-Section



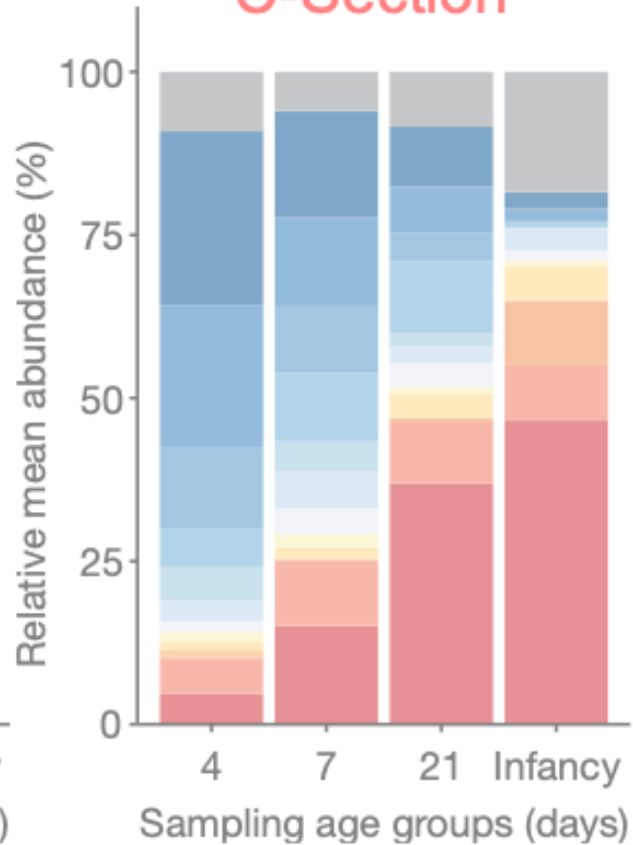
- Others
- Enterococcus*
- Streptococcus*
- Staphylococcus*
- Klebsiella*
- Enterobacter*
- Veillonella*
- Clostridium*
- Citrobacter*
- Lactobacillus*
- Parabacteroides*
- Bacteroides*
- Escherichia*
- Bifidobacterium*

C-section alters the microbiome

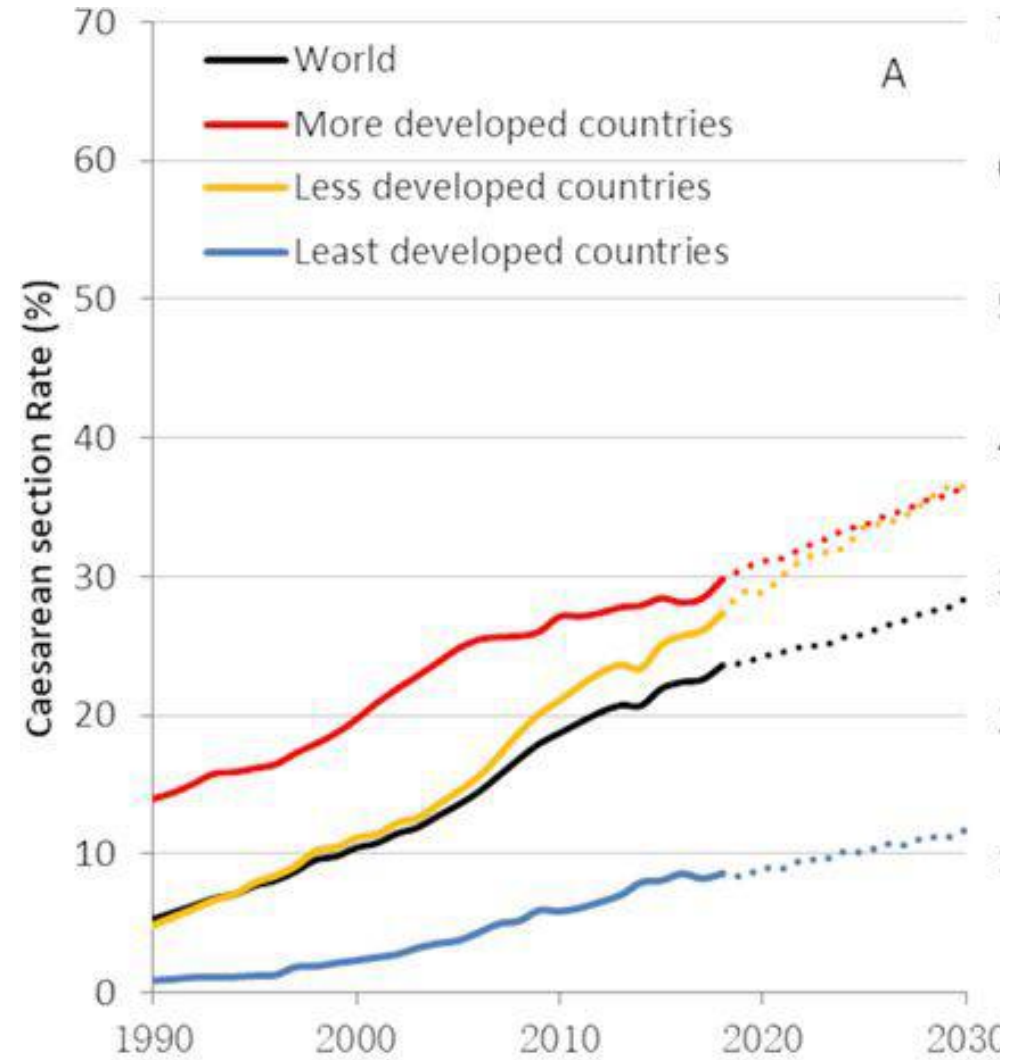
Vaginal



C-Section



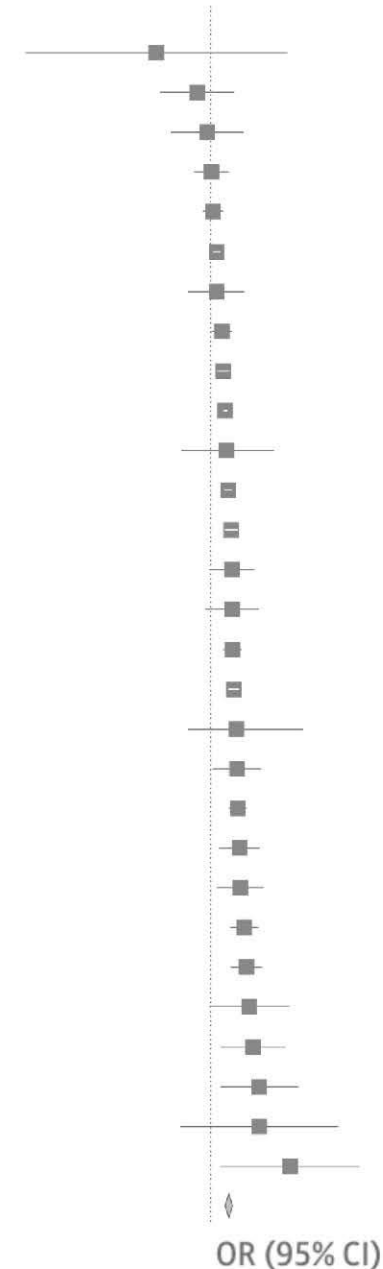
Shao 2019 Nature 574(7776):117-121



Betran AP, et al. BMJ Glob Health. 2021. PMID: 34130991.

CS and Autism Spectrum Diseases

Study	Outcome	Cesarean Delivery	Vaginal Delivery	OR (95% CI)
ASD				
Matsuishi et al, ³¹ 1999	ASD	1/18	25/205	0.42 (0.05-3.32)
Mrozek-Budzyn et al, ³² 2013	ASD	21/74	67/204	0.81 (0.45-1.45)
Hamadé et al, ⁸³ 2013	ASD	24/74	62/184	0.94 (0.53-1.68)
Maramara et al, ³³ 2014	ASD	73/31872	190/84023	1.01 (0.77-1.33)
Burstyn et al, ³⁴ 2010	ASD	304/49456	834/170572	1.04 (0.88-1.22)
Axelsson et al, ²⁰ 2019	ASD	1564/119433	6703/560440	1.10 (1.04-1.16)
Maimburg et al, ³⁵ 2006	ASD	76/633	371/4099	1.10 (0.70-1.70)
Dodds et al, ³⁶ 2011	ASD	218/26754	706/102948	1.19 (1.02-1.39)
Chien et al, ³⁷ 2015	ASD	684/174376	1166/362297	1.22 (1.11-1.34)
Curran et al, ²⁴ 2015	ASD	4330/340108	23960/2357206	1.26 (1.22-1.30)
Guisso et al, ³⁸ 2018	ASD	64/125	72/189	1.28 (0.63-2.70)
Yip et al, ³⁹ 2017 (Finland)	ASD	1331/156863	5753/891684	1.32 (1.24-1.40)
Schieve et al, ⁴⁰ 2014	ASD	NA	NA	1.38 (1.25-1.52)
Haglund et al, ⁴¹ 2011	ASD	36/7434	214/61530	1.39 (0.98-1.99)
Chen et al, ⁴² 2017	ASD	36/6285	52/12698	1.40 (0.91-2.15)
Yip et al, ³⁹ 2017 (Norway)	ASD	213/127922	1090/924475	1.41 (1.22-1.64)
Polo-Kantola et al, ⁴³ 2014	ASD	860/3349	3328/17117	1.43 (1.31-1.56)
Duan et al, ⁴⁴ 2014	ASD	152/287	134/285	1.50 (0.70-4.30)
Durkin et al, ⁴⁵ 2015	ASD	33/4624	127/26843	1.51 (1.03-2.22)
Yip et al, ³⁹ 2017 (Western Australia)	ASD	289/65953	802/279228	1.53 (1.34-1.75)
Eriksson et al, ⁴⁶ 2012	ASD	52/23286	134/94242	1.57 (1.14-2.17)
Hultman et al, ⁴⁷ 2002	ASD	87/352	321/2096	1.60 (1.10-2.30)
Kissin et al, ⁴⁸ 2015	ASD	328/27152	109/15231	1.70 (1.36-2.11)
Glasson et al, ⁴⁹ 2004	ASD	134/380	331/1398	1.76 (1.38-2.24)
Zhang et al, ⁵⁰ 2010	ASD	43/77	43/104	1.83 (0.98-3.43)
El-Baz et al, ⁵¹ 2011	ASD	39/89	61/213	1.94 (1.16-3.25)
Ji et al, ⁵² 2018	ASD	22/214	22/434	2.15 (1.16-3.97)
Winkler-Schwartz et al, ⁵³ 2014	ASD	7/35	5/48	2.15 (0.62-7.45)
Al-Jammas et al, ⁵⁴ 2012	ASD	14/19	36/81	3.50 (1.15-10.63)
Subtotal $I^2=69.5\%$, $P<.001$				1.33 (1.25-1.41)



Zhang, et al, 2019
JAMA Netw Open
2(8):e1910236

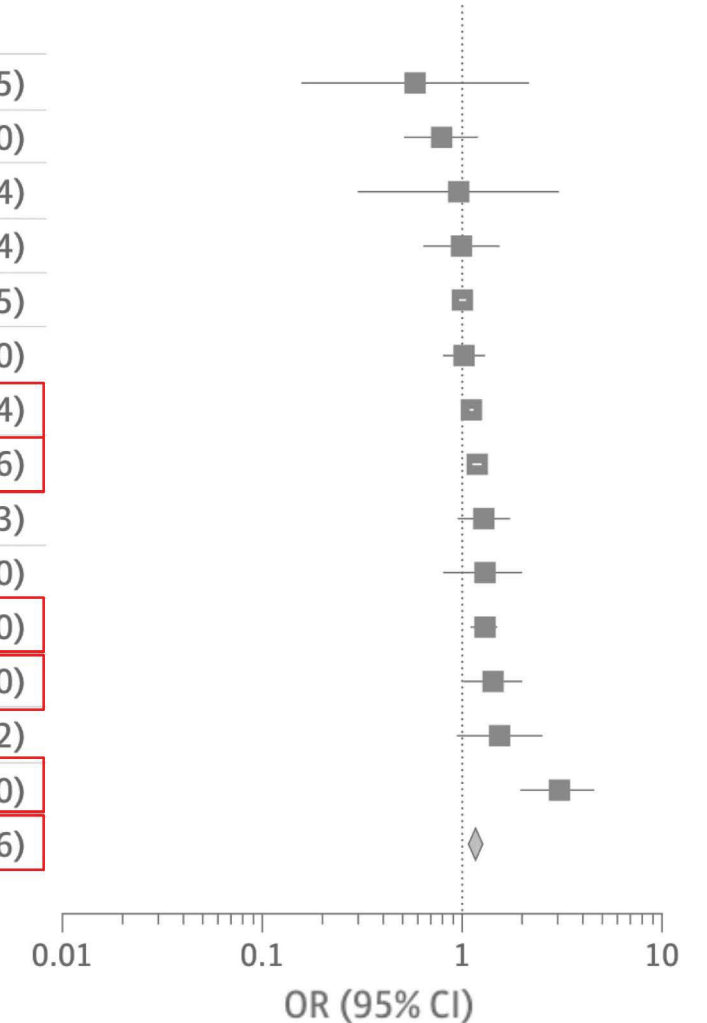
CS and Neurodevelopmental Diseases

ADHD

				ADHD
Çak et al, ⁵⁵ 2013	ADHD	18/80	4/12	0.58 (0.16-2.15)
Murray et al, ⁵⁶ 2016 (Pelotas)	ADHD	NA	NA	0.79 (0.51-1.20)
Yeo et al, ⁵⁷ 2015	ADHD	24/34	34/46	0.96 (0.30-3.04)
Gustafsson and Källén et al, ⁵⁸ 2011	ADHD	22/2996	215/29016	0.99 (0.64-1.54)
Axelsson et al, ¹⁹ 2019	ADHD	3167/117863	14804/553727	1.01 (0.97-1.05)
Silva et al, ⁵⁹ 2014	ADHD	2819/8863	10363/34829	1.02 (0.80-1.30)
Curran et al, ²⁴ 2016	ADHD	7230/238687	40548/1483861	1.11 (1.08-1.14)
Sucksdorff et al, ⁶⁰ 2018	ADHD	1892/8034	8428/40963	1.19 (1.12-1.26)
Chen et al, ⁴² 2017	ADHD	71/6320	112/12758	1.28 (0.95-1.73)
Ketzer et al, ⁶¹ 2012	ADHD	NA	NA	1.30 (0.80-2.00)
Halmøy et al, ⁶² 2012	ADHD	186/69051	2137/1103345	1.30 (1.10-1.50)
Ji et al, ⁵² 2018	ADHD	75/267	113/525	1.42 (1.02-2.00)
Murray et al, ⁵⁶ 2016 (ALSPAC)	ADHD	NA	NA	1.54 (0.94-2.52)
Amiri et al, ⁶³ 2012	ADHD	103/162	61/168	3.06 (1.96-4.80)
Subtotal $I^2=79.2\%$, $P<.001$				1.17 (1.07-1.26)

ADHD

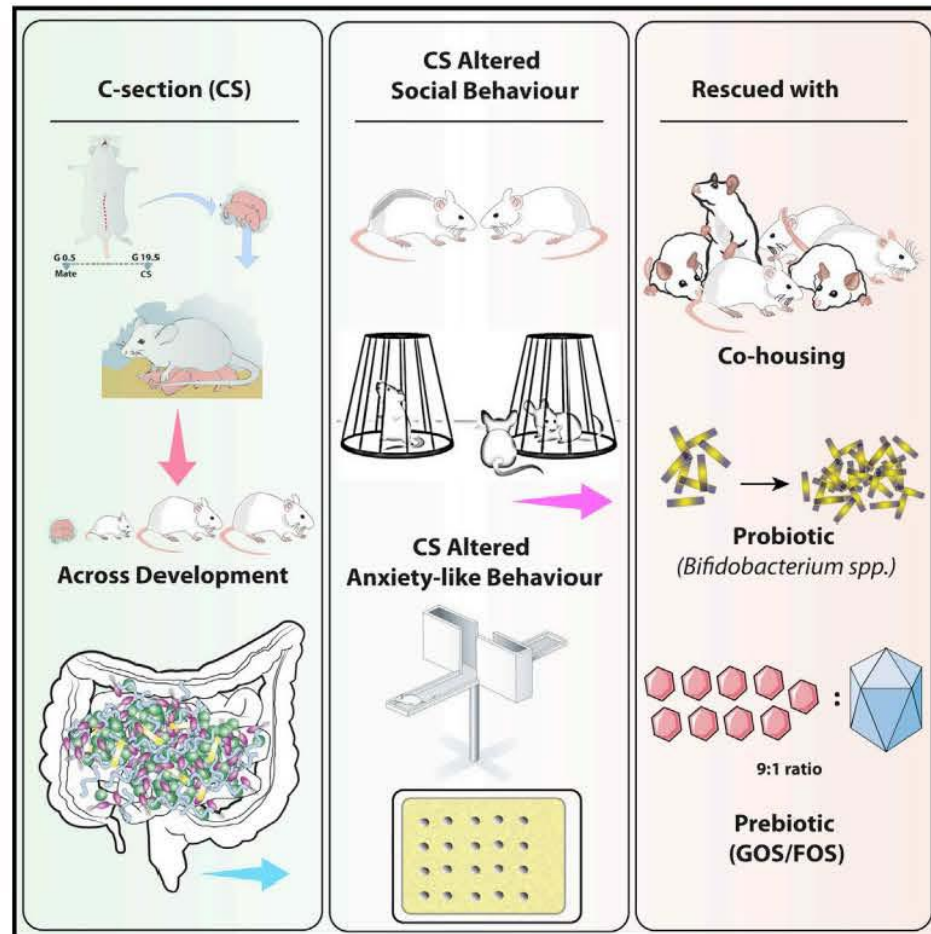
Attention Deficit
Hyperactivity Disorder



Current Biology

Enduring Behavioral Effects Induced by Birth by Caesarean Section in the Mouse

Graphical Abstract



Authors

Livia H. Morais, Anna V. Golubeva,
Gerard M. Moloney, ...,
Catherine Stanton, Timothy G. Dinan,
John F. Cryan

Correspondence

j.cryan@ucc.ie

In Brief

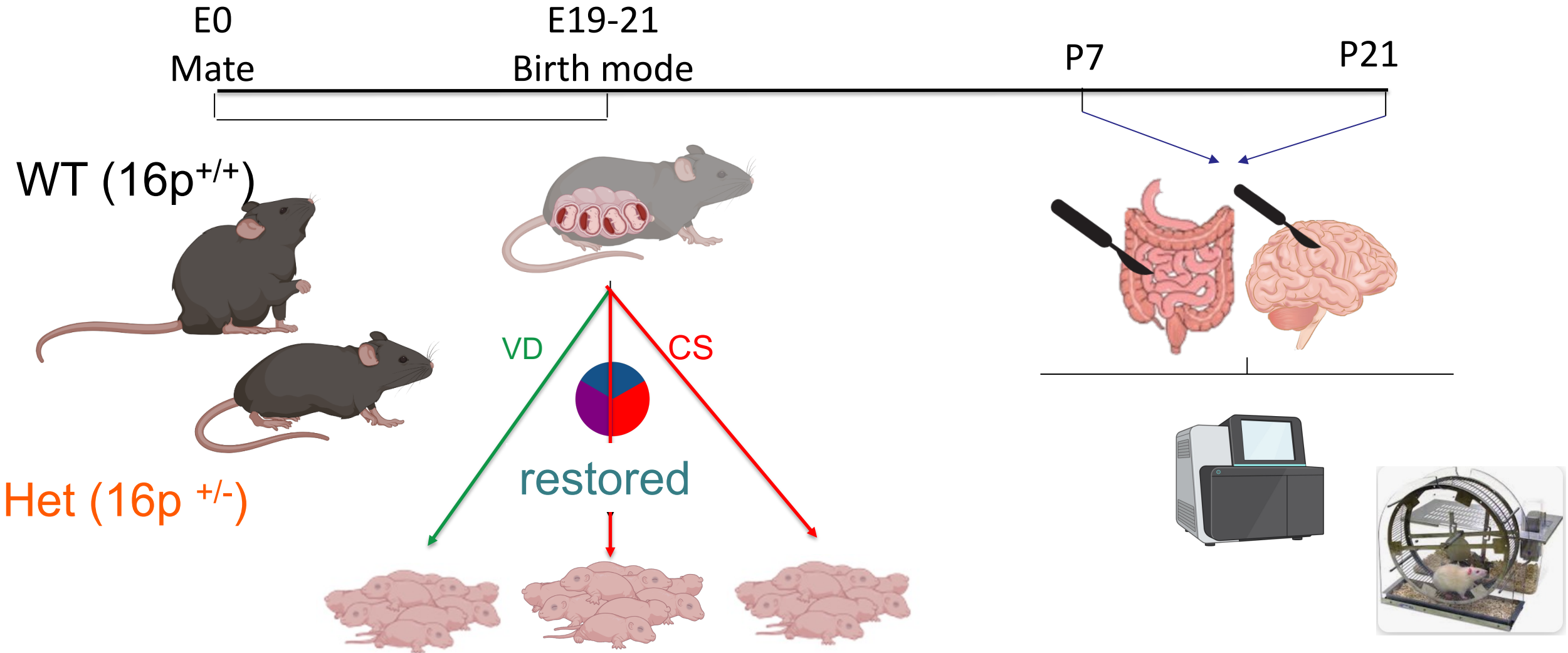
Recent evidence points to an important role for the microbiome in regulating brain function and behavior. Here, Morais *et al.* show that birth by C-section results in a different pattern of microbiota colonization with long-term behavioral consequences in the mouse. Targeting the gut microbiota reverses social behavioral effects of C-section.

Morais et al., 2020, Current Biology

Experimental Design: brain and the gut microbiome

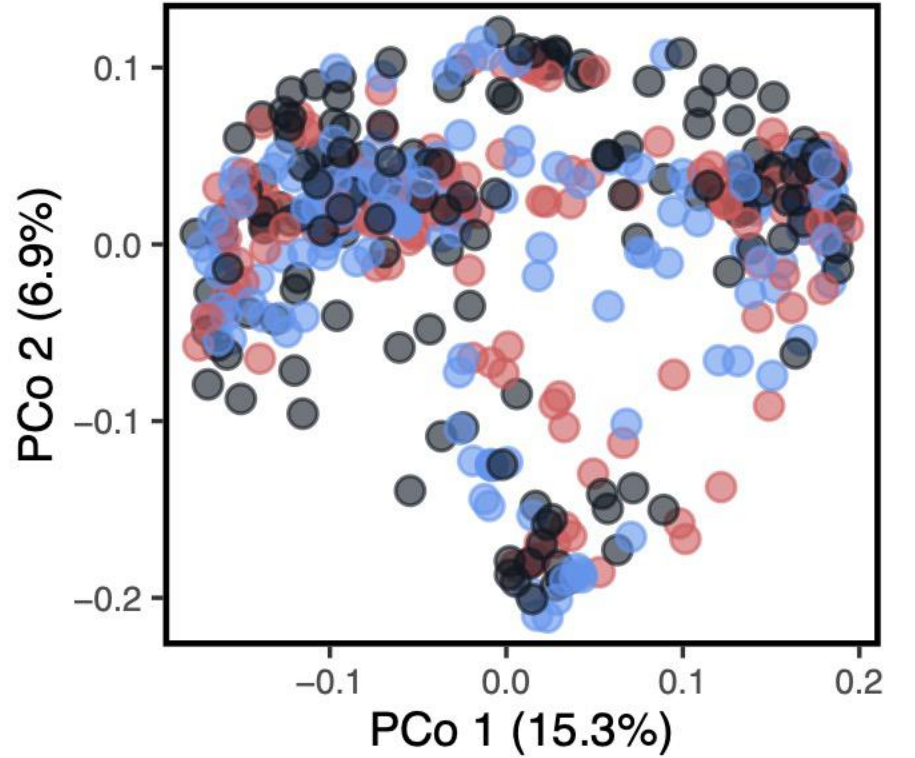


Jeremy Lessing



Beta Diversity

Birth Mode



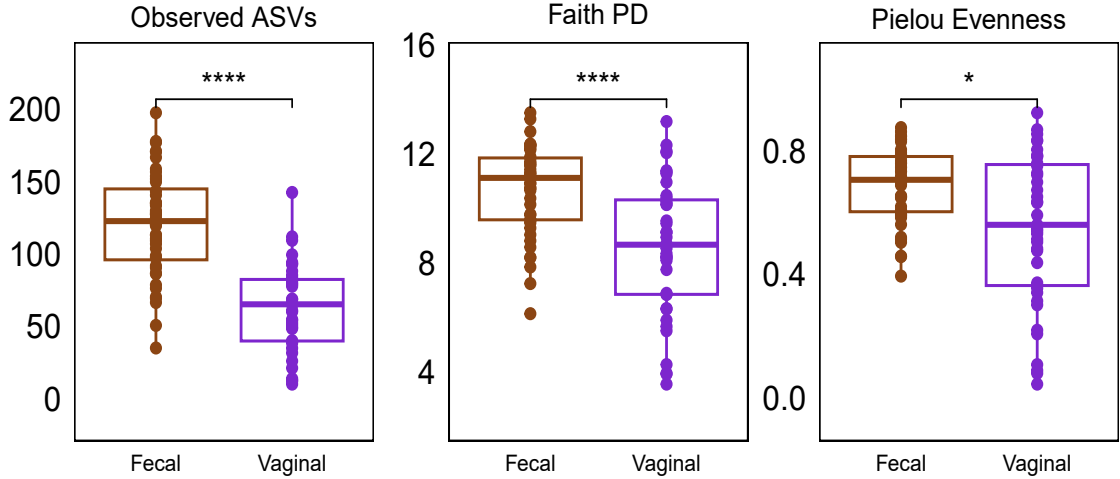
$p < 0.001$

$\omega^2 = 1\%$

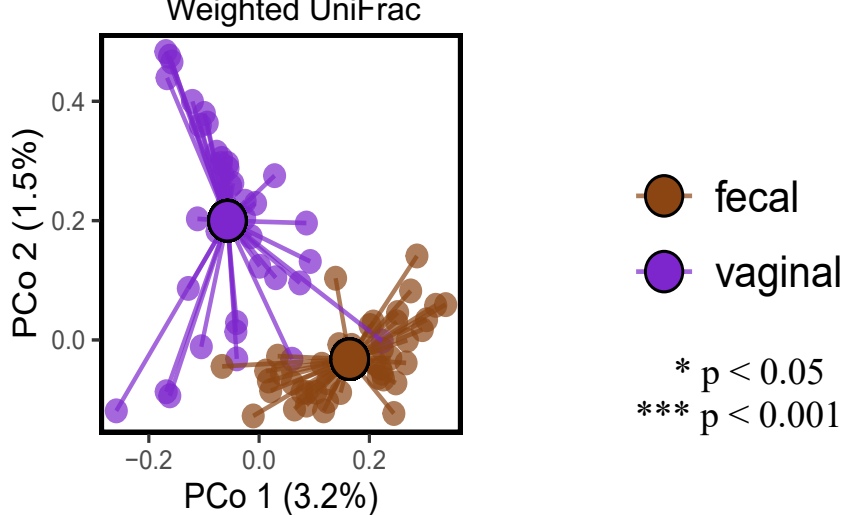
- VF
- CS
- CSR

Vaginal community is distinct...

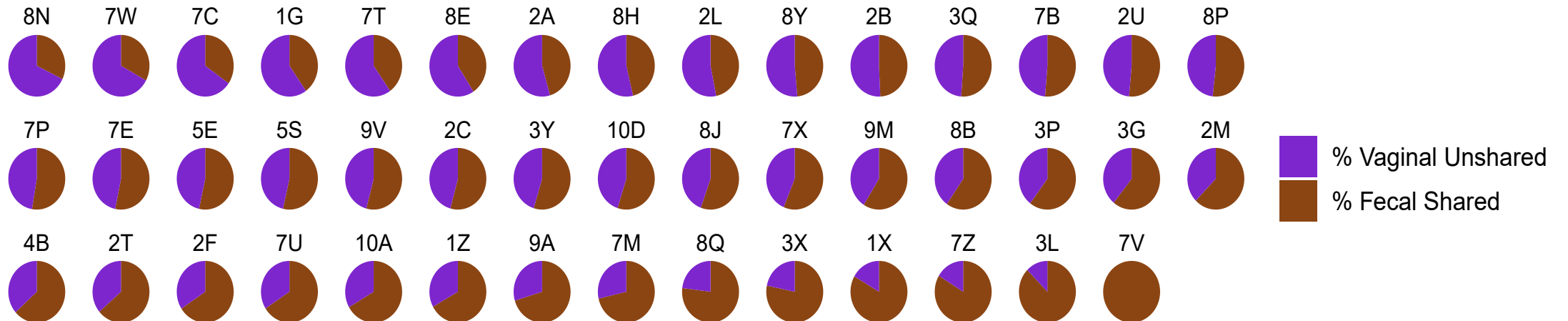
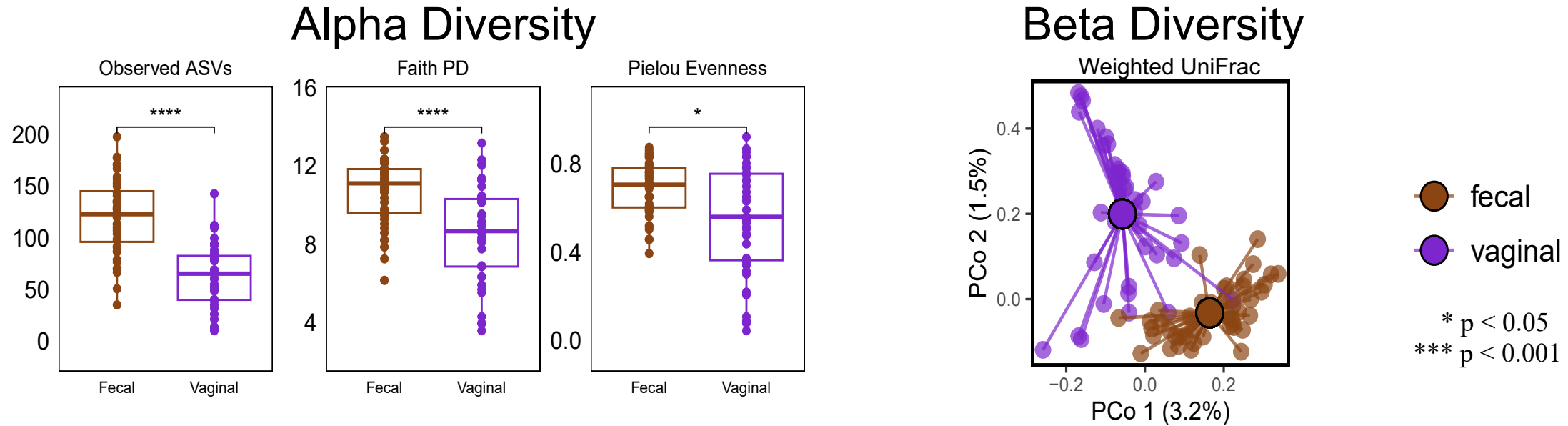
Alpha Diversity



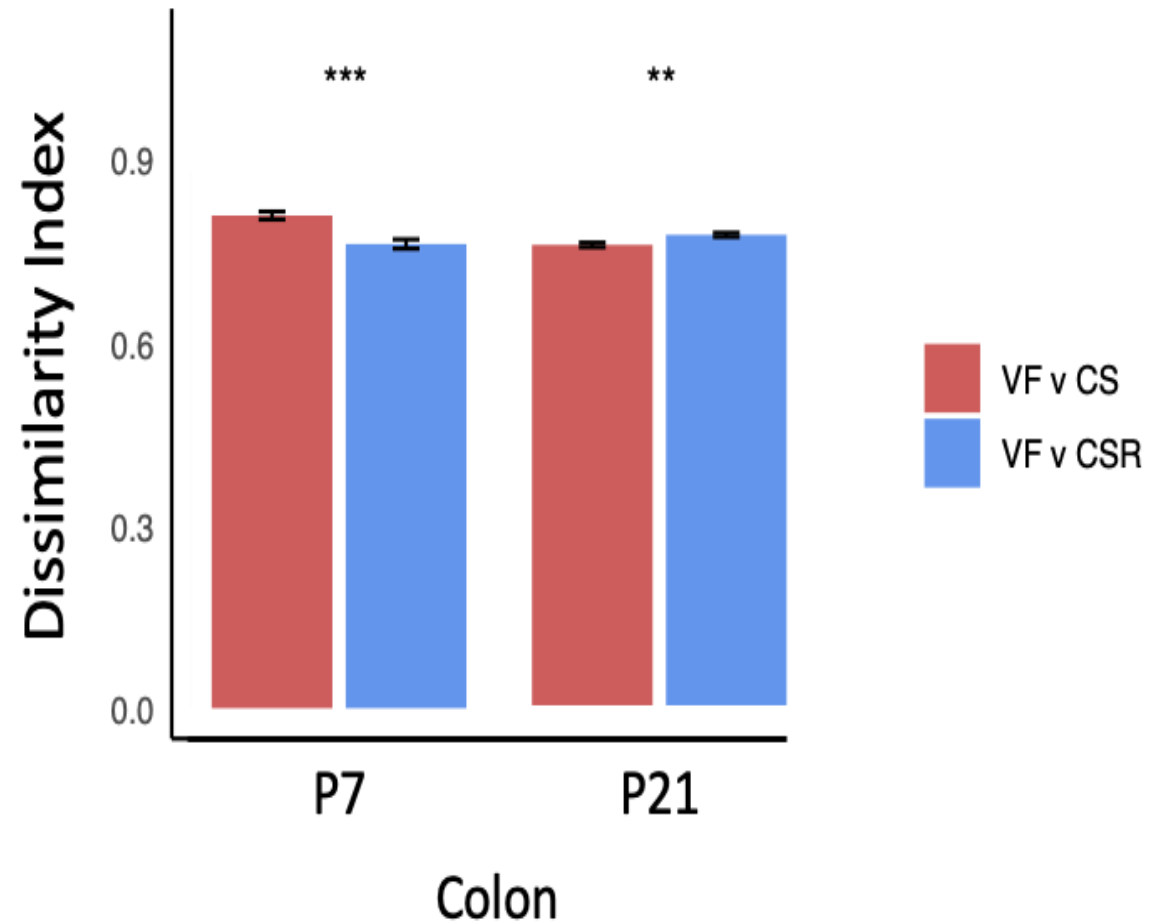
Beta Diversity



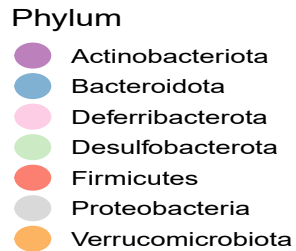
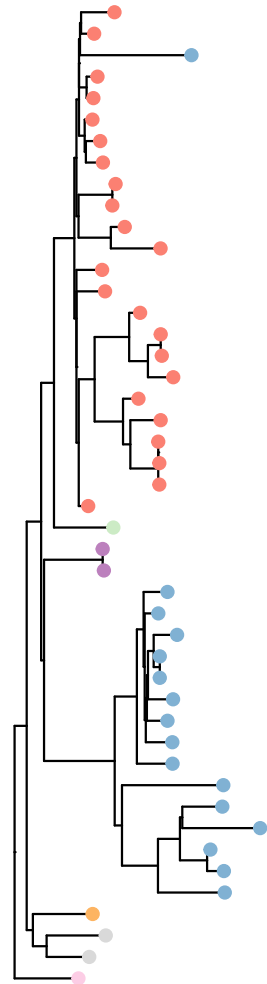
Vaginal community is distinct... but overlaps with Feces



Restoration decreases Bray Curtis distance between C-section and –vaginal birth



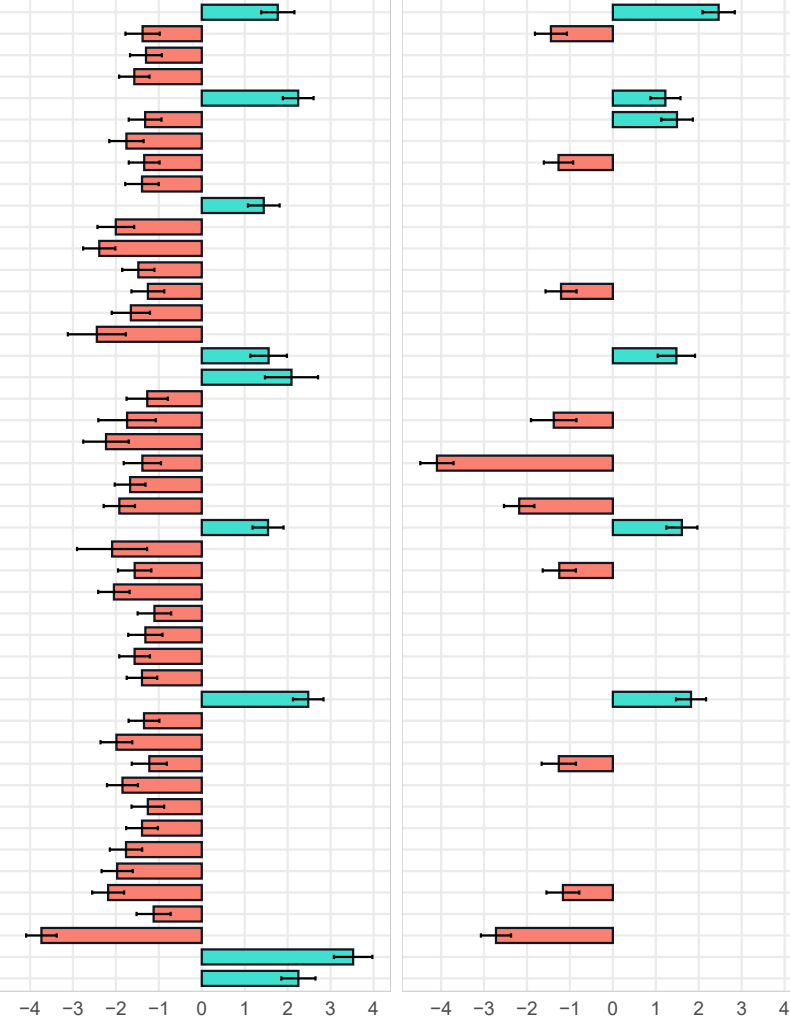
Restoration in the mouse colon at P7



Blautia sp. (ASV 9487)
Lachnospiraceae_NK4A136_group sp. (ASV 9282)
Muribaculaceae sp. (ASV 8854)
Lachnospiraceae_NK4A136_group sp. (ASV 8334)
Lachnospiraceae_NK4A136_group sp. (ASV 8197)
[Eubacterium]_xylanophilum_group sp. (ASV 8082)
[Eubacterium]_xylanophilum_group sp. (ASV 8043)
Lachnospiraceae_FCS020_group sp. (ASV 7987)
Lachnospiraceae_NK4A136_group sp. (ASV 7324)
Lachnospiraceae_NK4A136_group sp. (ASV 7317)
Lactobacillus sp. (ASV 6978)
Streptococcus danieliae (ASV 6891)
Lachnospiraceae_UCG-006 sp. (ASV 6639)
Roseburia sp. (ASV 6419)
Turicibacter sp. (ASV 6109)
Enterococcus sp. (ASV 6062)
Enterococcus sp. (ASV 6042)
Staphylococcus sp. (ASV 6005)
Lactobacillus intestinalis (ASV 5854)
Lactobacillus sp. (ASV 5731)
Lactobacillus sp. (ASV 5676)
Lactobacillus sp. (ASV 5669)
Lactobacillus sp. (ASV 5605)
Roseburia sp. (ASV 5272)
Bilophila sp. (ASV 4213)
Rothia nasimurium (ASV 3577)
Rothia nasimurium (ASV 3567)
Muribaculaceae sp. (ASV 2857)
Muribaculaceae sp. (ASV 2772)
Muribaculaceae sp. (ASV 2645)
Muribaculaceae sp. (ASV 2551)
Muribaculaceae sp. (ASV 2550)
Muribaculaceae sp. (ASV 2484)
Muribaculaceae sp. (ASV 2417)
Muribaculaceae sp. (ASV 2344)
Muribaculaceae sp. (ASV 2223)
Muribaculaceae sp. (ASV 2117)
Rikenellaceae_RC9_gut_group sp. (ASV 2011)
Odoribacter sp. (ASV 1980)
Alistipes sp. (ASV 1812)
Alistipes sp. (ASV 1784)
Bacteroides sartorii (ASV 1563)
Akkermansia sp. (ASV 810)
Parasutterella sp. (ASV 768)
Escherichia-Shigella sp. (ASV 515)
Mucispirillum sp. (ASV 386)

C-Section

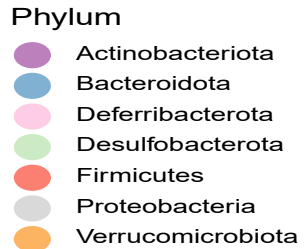
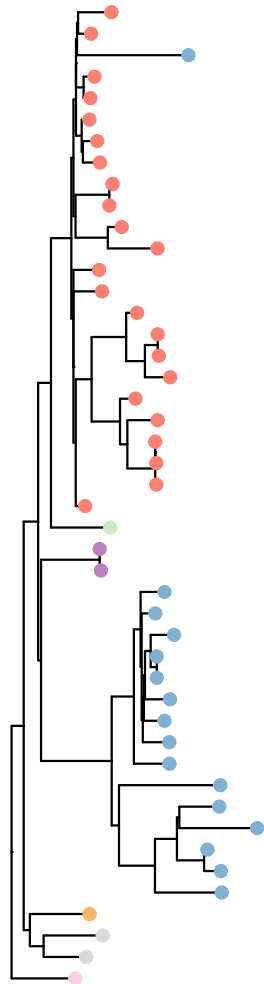
Restoration



Log Fold Change



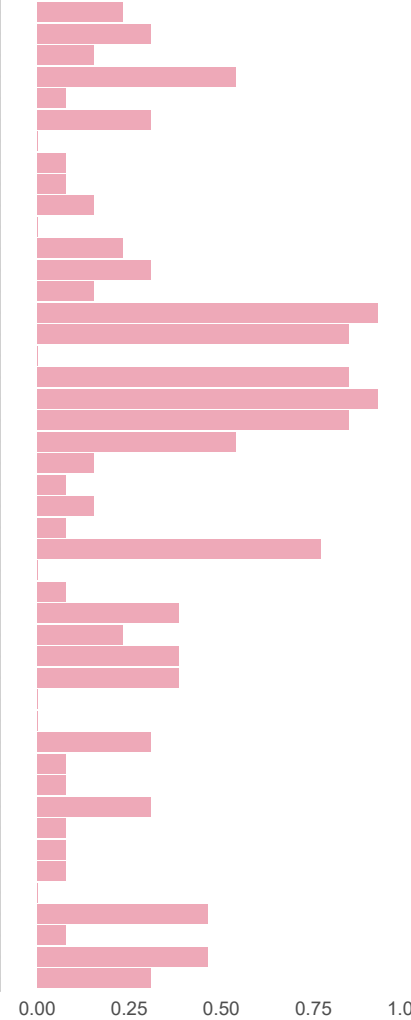
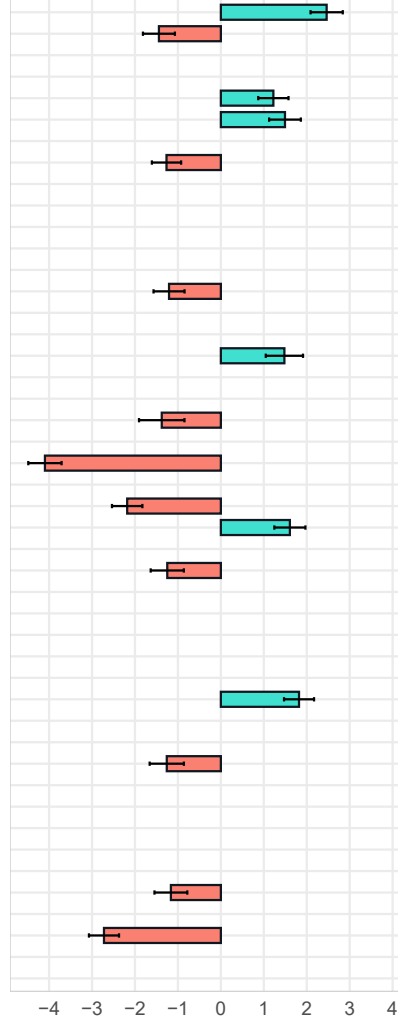
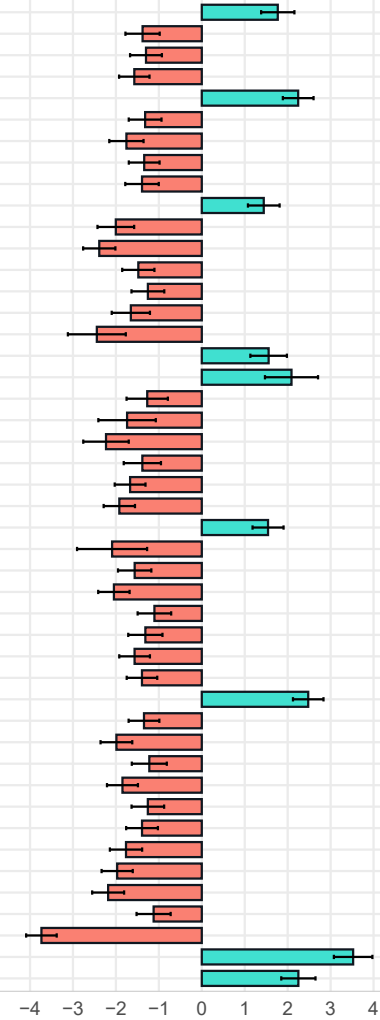
Restoration in the mouse colon at P7



C-Section

Restoration

Blautia sp. (ASV 9487)
Lachnospiraceae_NK4A136_group sp. (ASV 9282)
Muribaculaceae sp. (ASV 8854)
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Enterococcus sp. (ASV 6062)
Enterococcus sp. (ASV 6042)
Staphylococcus sp. (ASV 6005)
Lactobacillus intestinalis (ASV 5854)
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Lactobacillus sp. (ASV 5676)
Lactobacillus sp. (ASV 5669)
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Akkermansia sp. (ASV 810)
Parasutterella sp. (ASV 768)
Escherichia-Shigella sp. (ASV 515)
Mucispirillum sp. (ASV 386)

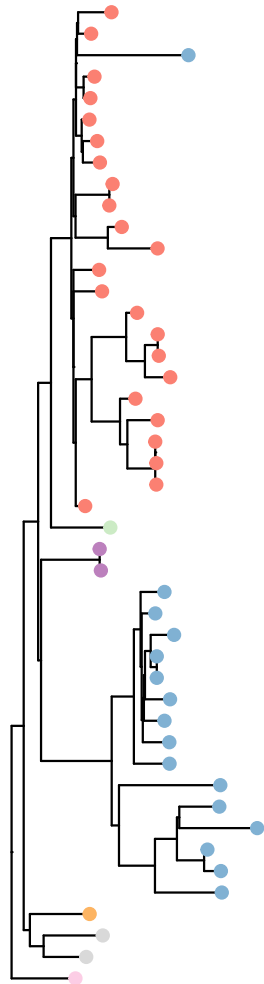


Log Fold Change

Present in
Vaginal Inoculum



Vaginal inocula is responsible for colon restoration at P7



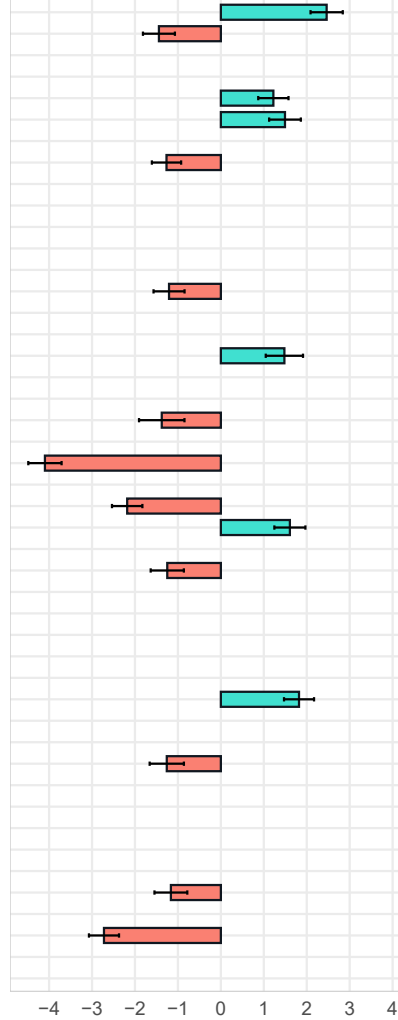
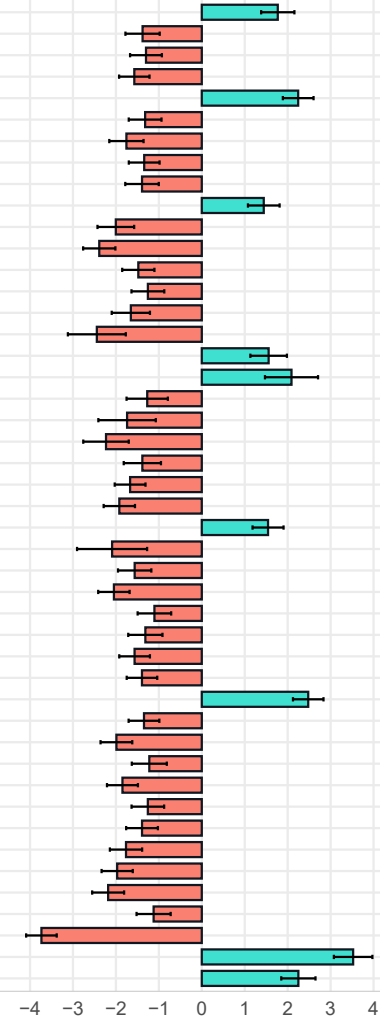
Phylum

- Actinobacteriota
- Bacteroidota
- Deferribacterota
- Desulfobacterota
- Firmicutes
- Proteobacteria
- Verrucomicrobiota

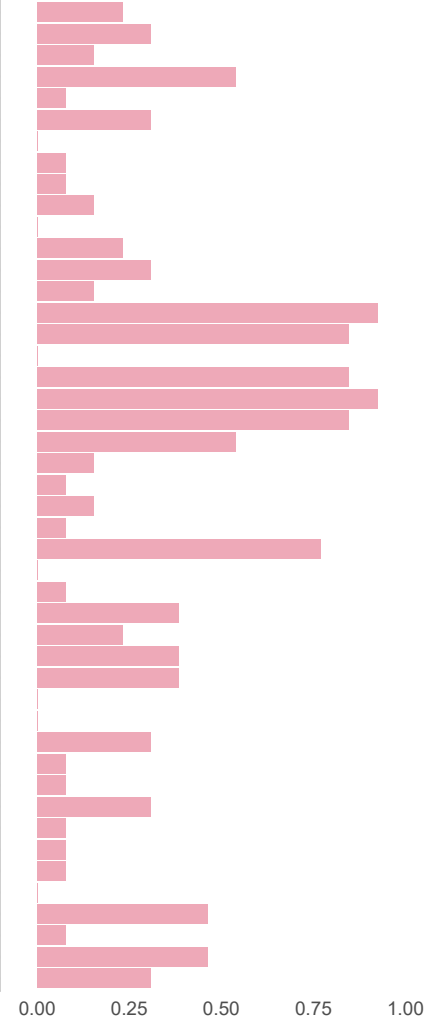
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Enterococcus sp. (ASV 6042)
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Muribaculaceae sp. (ASV 2550)
Muribaculaceae sp. (ASV 2484)
Muribaculaceae sp. (ASV 2417)
Muribaculaceae sp. (ASV 2344)
Muribaculaceae sp. (ASV 2223)
Muribaculaceae sp. (ASV 2117)
Rikenellaceae_RC9_gut_group sp. (ASV 2011)
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Alistipes sp. (ASV 1812)
Alistipes sp. (ASV 1784)
Bacteroides sartorii (ASV 1563)
Akkermansia sp. (ASV 810)
Parasutterella sp. (ASV 768)
Escherichia-Shigella sp. (ASV 515)
Mucispirillum sp. (ASV 386)

C-Section

Restoration

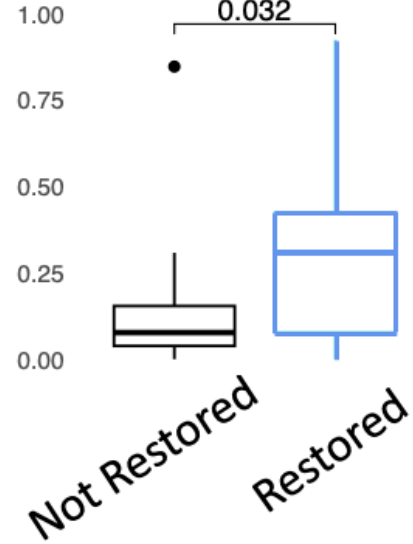


Log Fold Change



Present in vaginal inoculum

Presence in Vaginal Inocula



(Mann Whitney U Test)

Brain protein profile effects induced by C-section

Proliferation

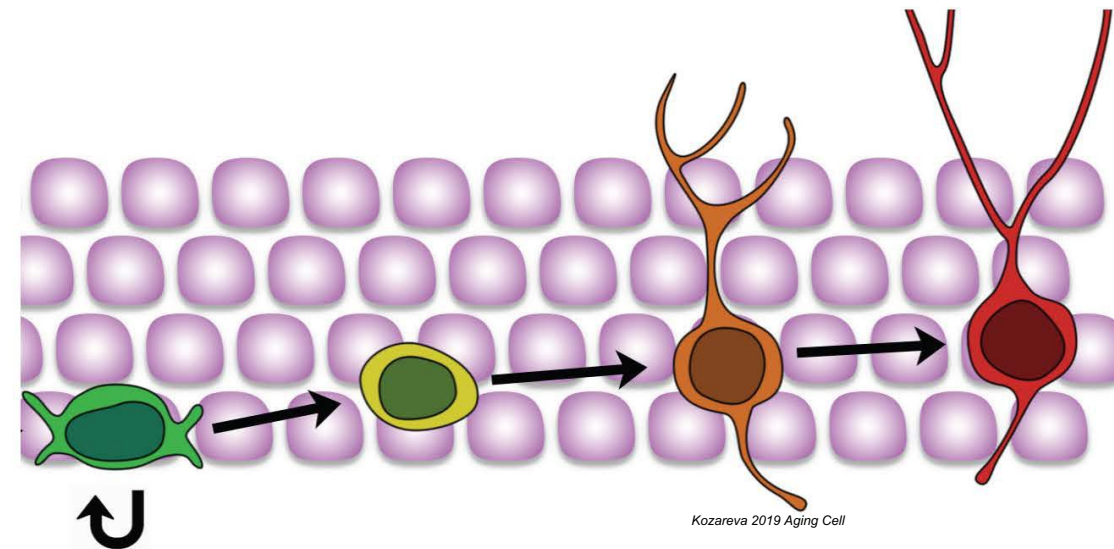
- Cyclin E G₁ -> S phase transition
- Sox2 SRY Box 2: Pluripotent neural stem-cells

Post-mitotic cytoskeleton

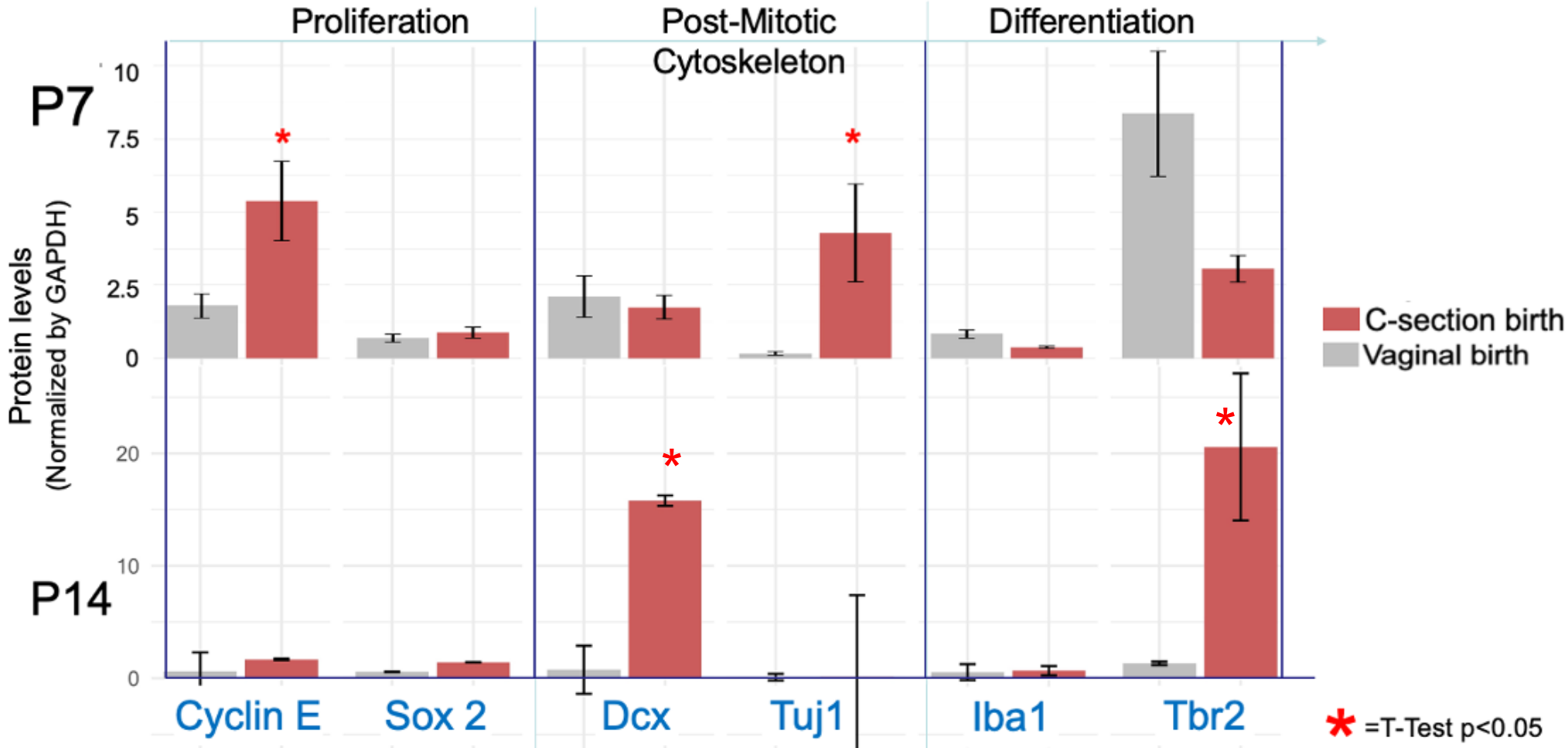
- Tuj1 B3 Tubulin: Neuron cytoskeleton
- Dcx Doublecortin: Developing brain

Differentiation

- GFAP Glial Fibrillary Acidic Protein: Astrocytes
- Tbr2 T-box Brain 2: Developing brain
- Iba1 Ionized Ca²⁺ Binding Adapter 1: Microglia

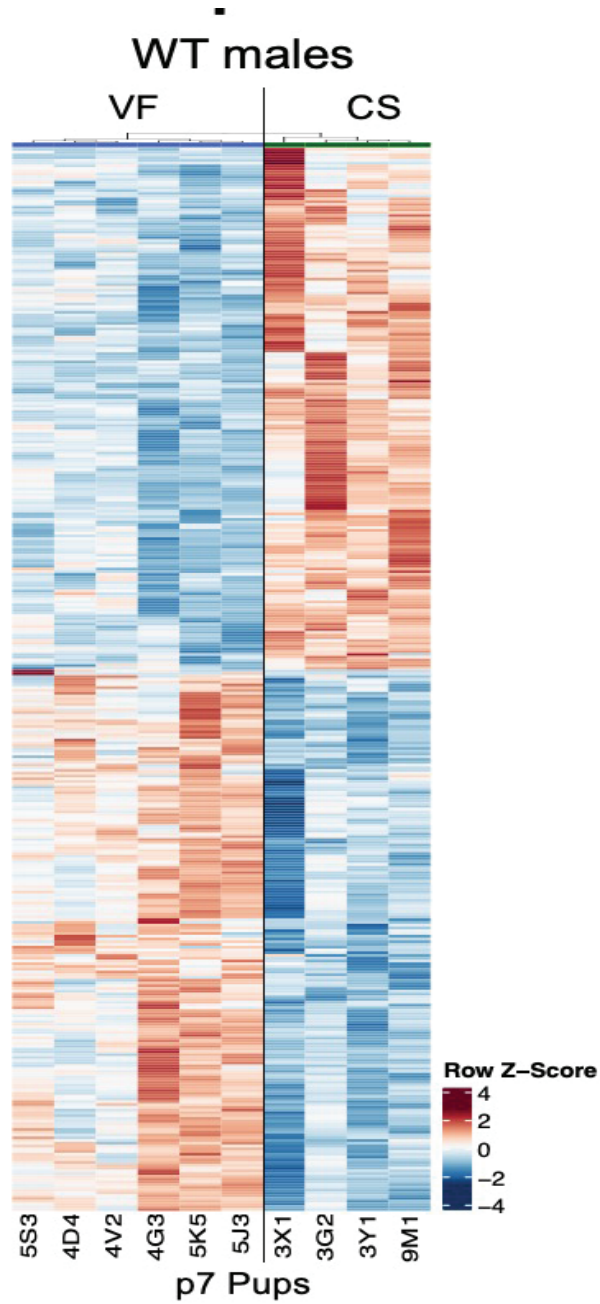


Cerebellum protein profile effects induced by C-section



Changes also in hippocampus proteins

Unsupervised Heatmap: CS affects Hyppocampus transcriptome



189 Genes Up in CS

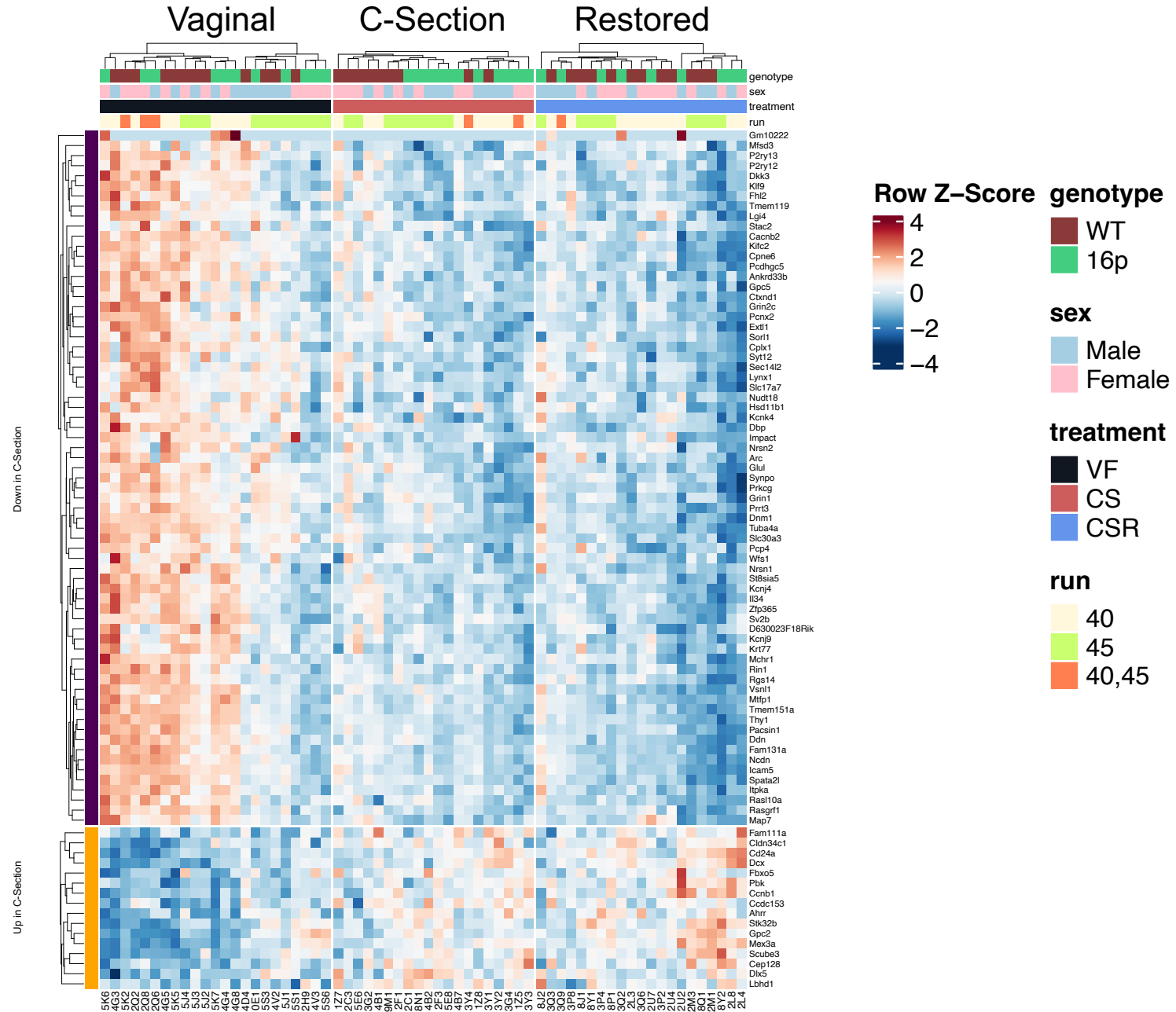
	Selected GO Biological Process	p Value
CS Upregulated	neural plate development	0.002
	ventricular system development	0.004
	central nervous system development	0.023
	brain development	0.024
	neural tube development	0.029
	neural plate morphogenesis	0.029

196 Genes Down in CS

	Selected GO Biological Process	p Value
CS Downregulated	synaptic vesicle recycling	< 0.001
	synapse organization	< 0.001
	regulation of neuronal synaptic plasticity	< 0.001
	trans-synaptic signaling	0.001
	synaptic vesicle endocytosis	0.001
	neurogenesis	0.001
	presynaptic endocytosis	0.001
	synaptic signaling	0.001
	synapse assembly	0.001
	neuron projection morphogenesis	0.001
	neuron projection development	0.001
	synaptic vesicle maturation	0.001
	postsynapse organization	0.002
	regulation of trans-synaptic signaling	0.003
	postsynaptic endocytosis	0.004
	postsynaptic neurotransmitter receptor internalization	0.004
	regulation of synapse organization	0.005
	neuron development	0.005
	regulation of synapse structure or activity	0.006
	neurotransmitter receptor internalization	0.006
	synaptic vesicle lumen acidification	0.006
	oligodendrocyte differentiation	0.006
	regulation of synaptic plasticity	0.007
	gliogenesis	0.008
	learning or memory	0.009
	axonogenesis	0.011
	cell morphogenesis involved in neuron differentiation	0.011
glial cell differentiation	0.015	
regulation of synaptic vesicle endocytosis	0.016	
neuron differentiation	0.018	
axon development	0.023	
regulation of nervous system development	0.026	
myelination	0.027	
regulation of postsynaptic membrane neurotransmitter receptor levels	0.028	
generation of neurons	0.029	
postsynaptic specialization assembly	0.029	
regulation of postsynapse organization	0.029	
excitatory synapse assembly	0.029	
peripheral nervous system development	0.029	
synaptic vesicle transport	0.038	
postsynapse assembly	0.047	
Schwann cell differentiation	0.05	

CS-altered hippocampus transcriptomic not restored by vaginal seeding

P7 Hippocampus



Summary

C-section causes early alterations in brain development

Microbial restoration after C-section partially restored the microbiome

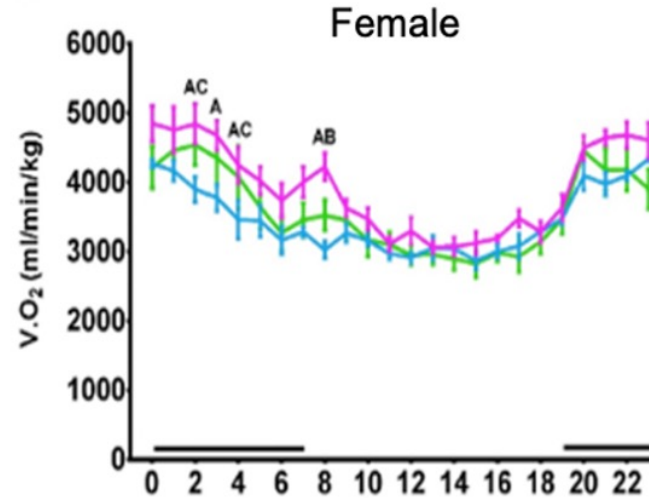
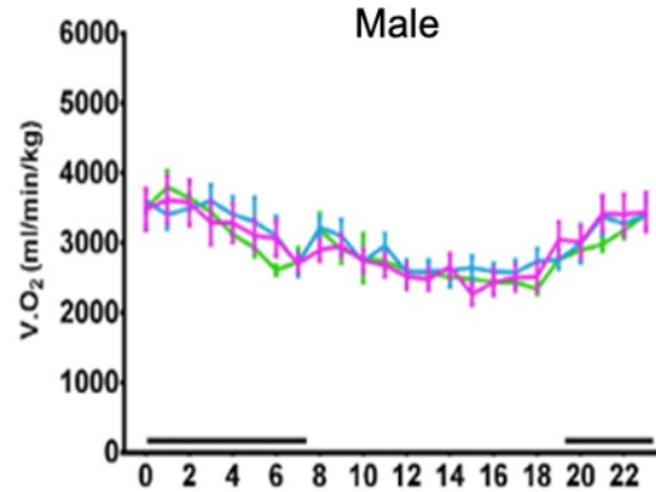
Microbial restoration did not fully rescue brain gene expression at P7

Metabolic/Behavioral effects induced by C-section birth in 11w old Swiss Webster mice

5-6 individuals from different litters per group

- Vaginal birth
- C-section birth
- C-section + vaginal restoration

-Lower nocturnal metabolic rate in CS females- partially restored

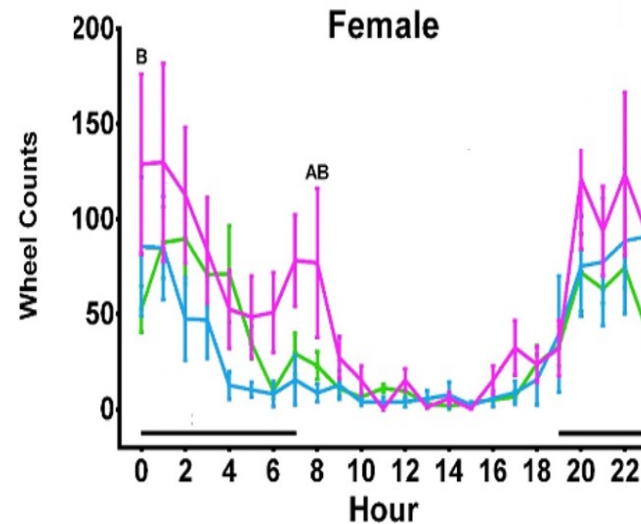
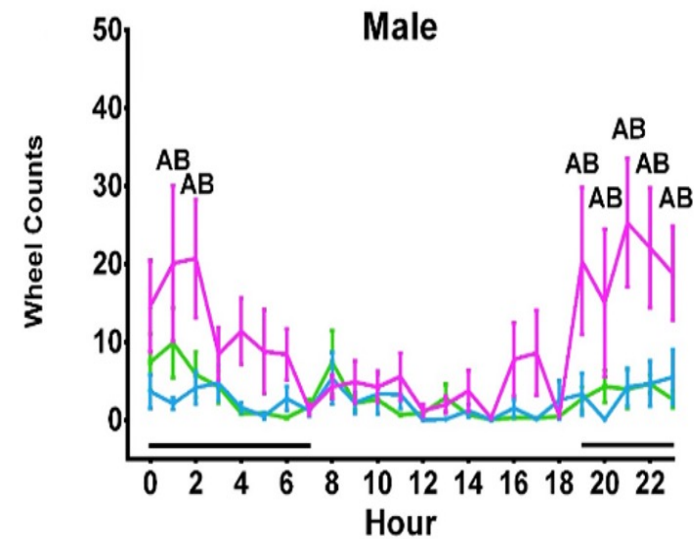
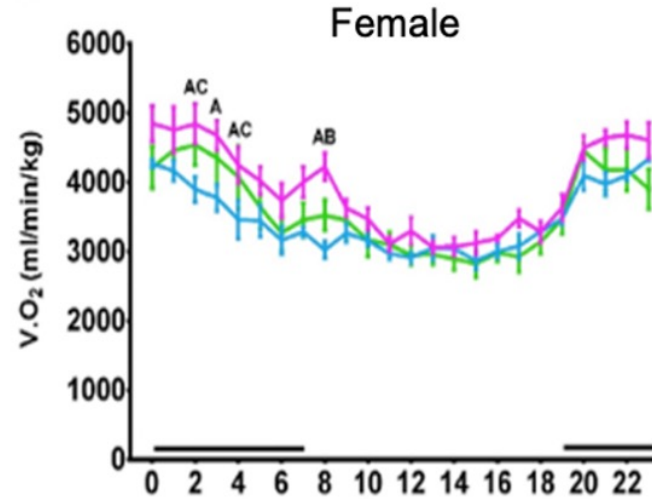
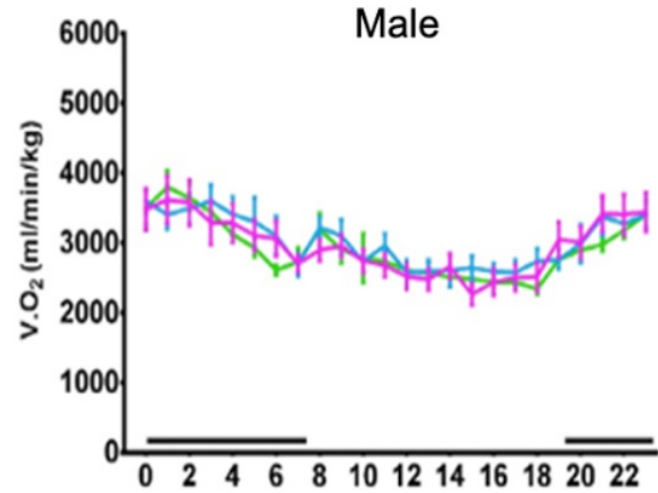


Metabolic/Behavioral effects induced by C-section birth in 11w old Swiss Webster mice

Vaginal birth
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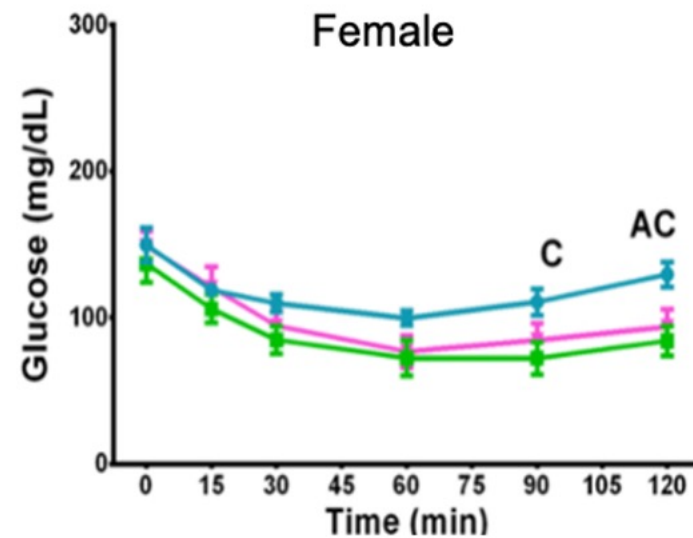
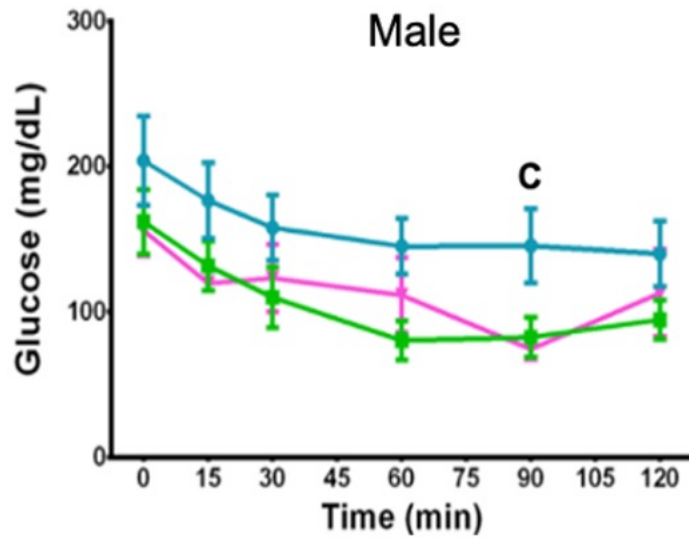
-Lower nocturnal activity in CS-restored



Metabolic effects induced by C-section birth in 11w old Swiss Webster mice

5-6 individuals from different litters per group

- Vaginal birth
- C-section birth
- C-section + vaginal restoration



-Glucose intolerance in CS restored



Lower energy expenditure in babies born to germ-free mice fed HFD

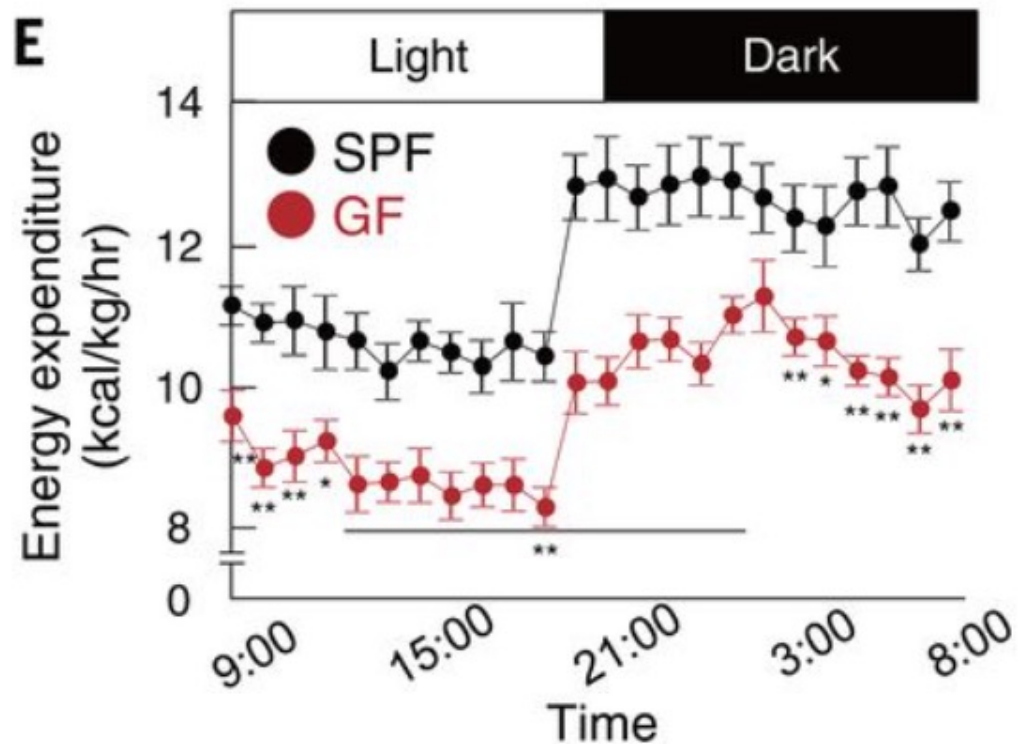


Fig. 1 Offspring from GF mothers exhibit severe obese phenotype when fed a HFD.

Lower energy expenditure in babies born to germ-free mice fed HFD

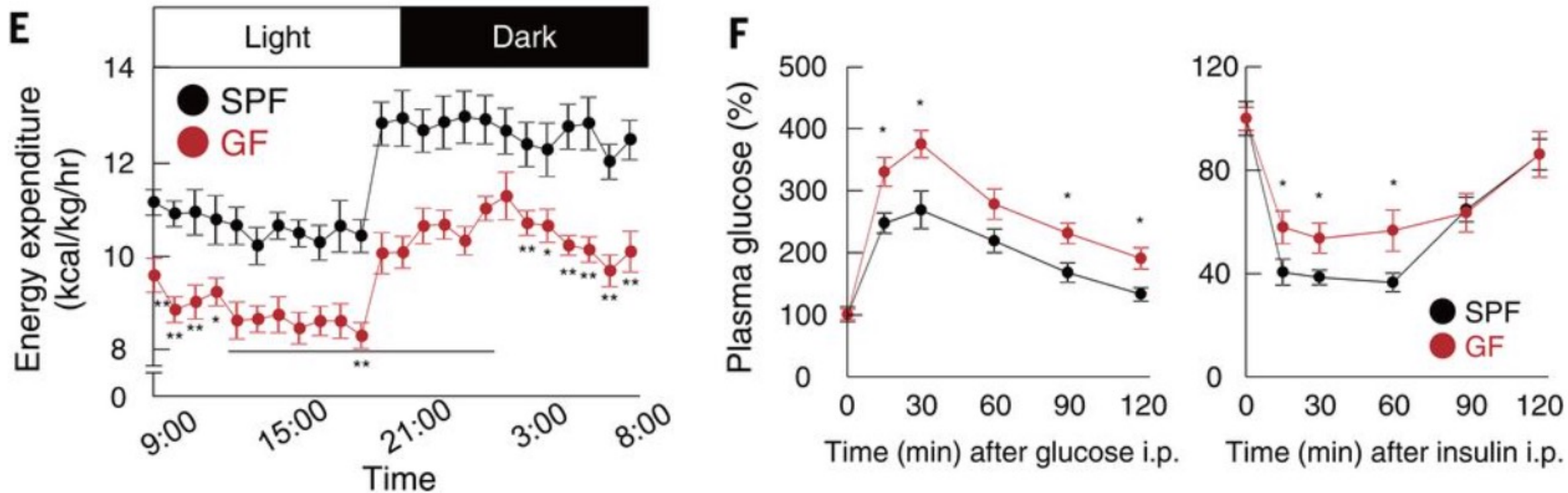
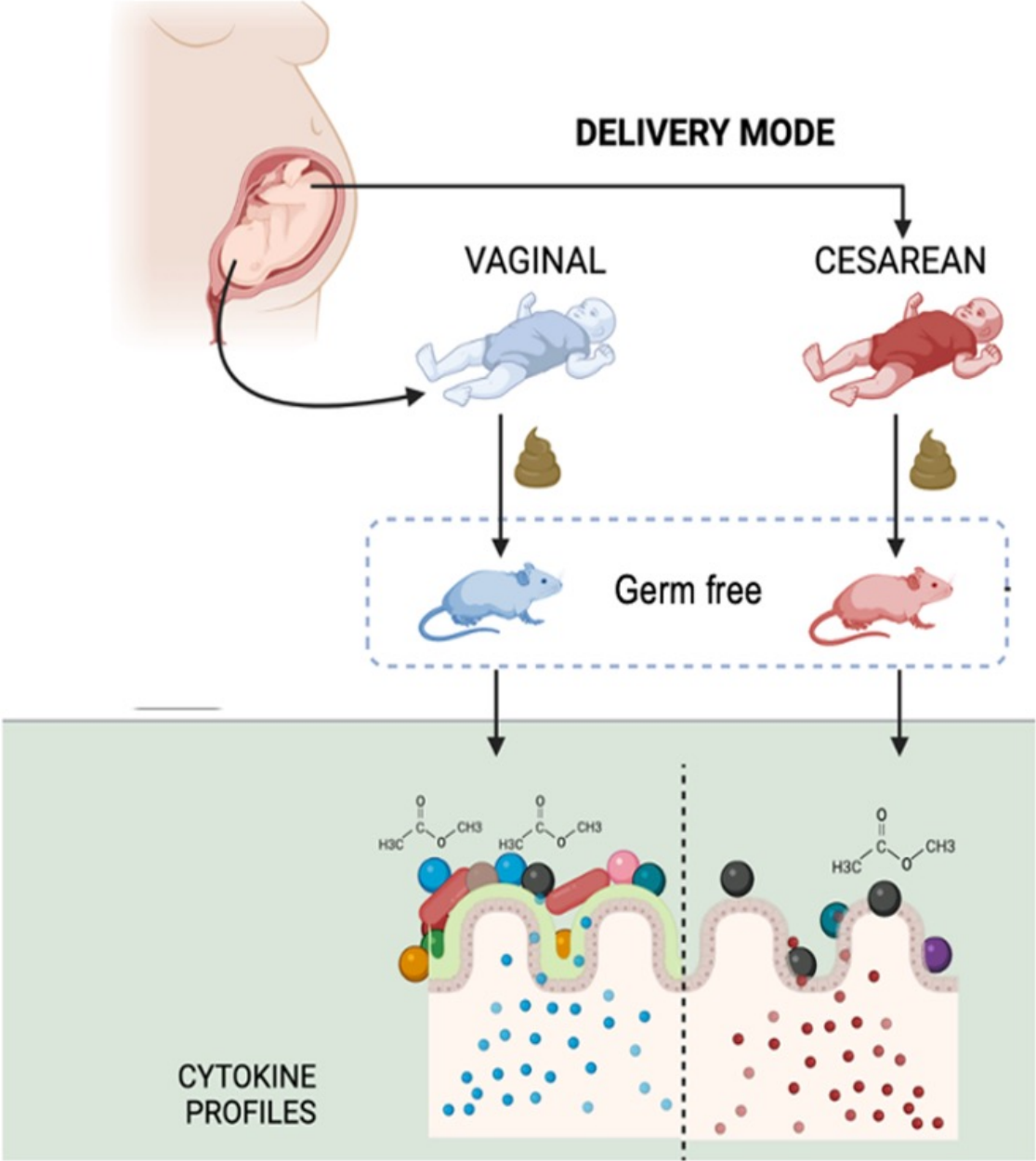
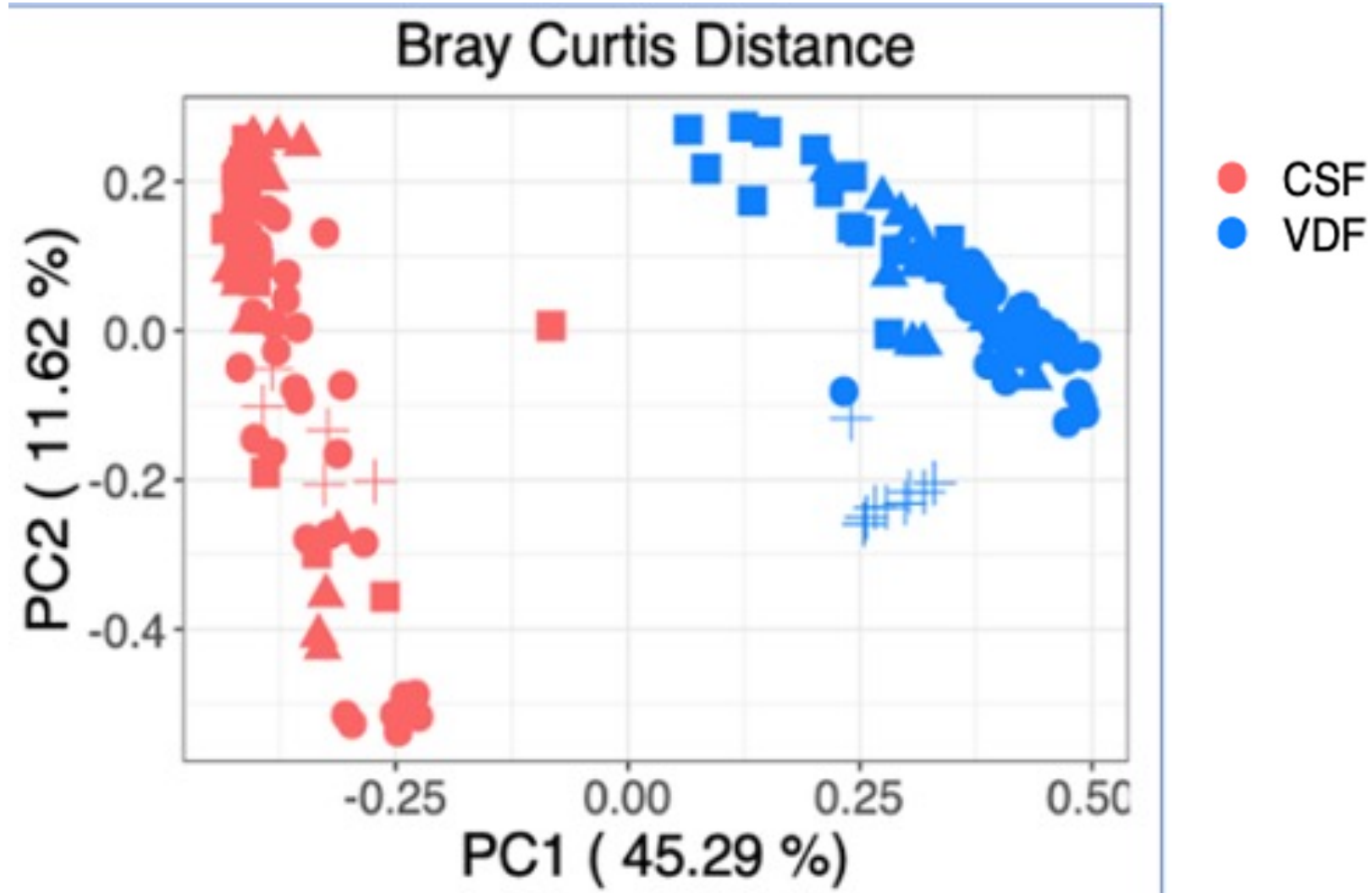


Fig. 1 Offspring from GF mothers exhibit severe obese phenotype when fed a HFD.

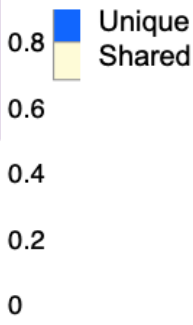
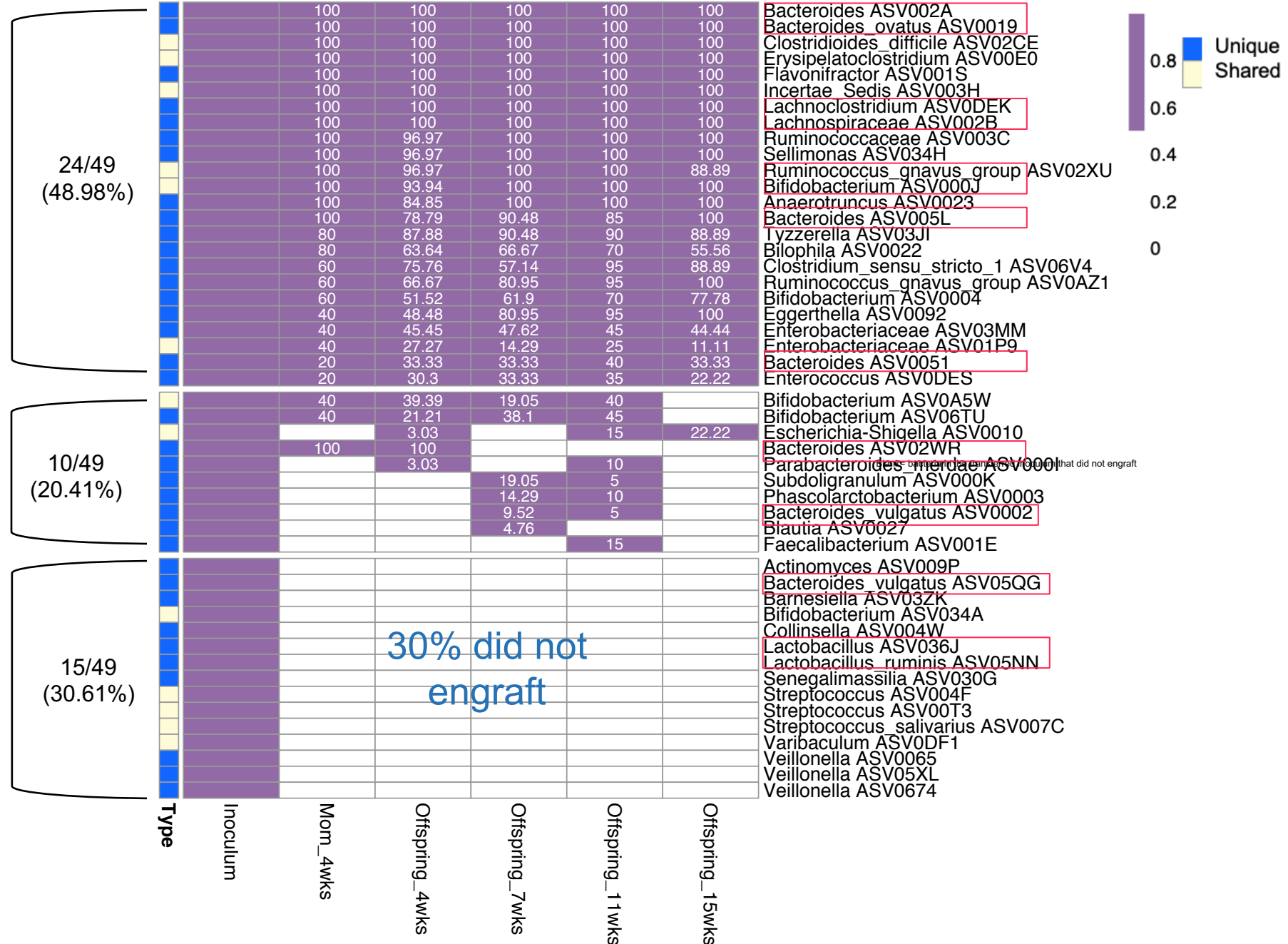
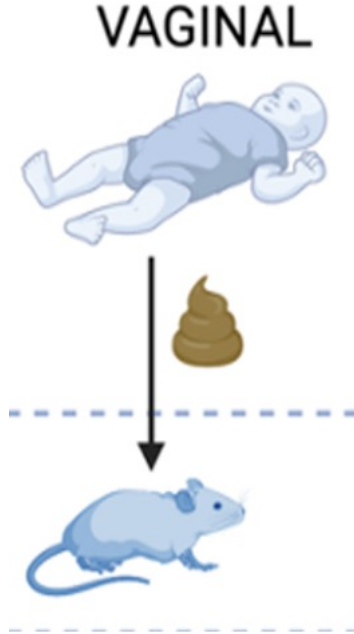
Experimental Design



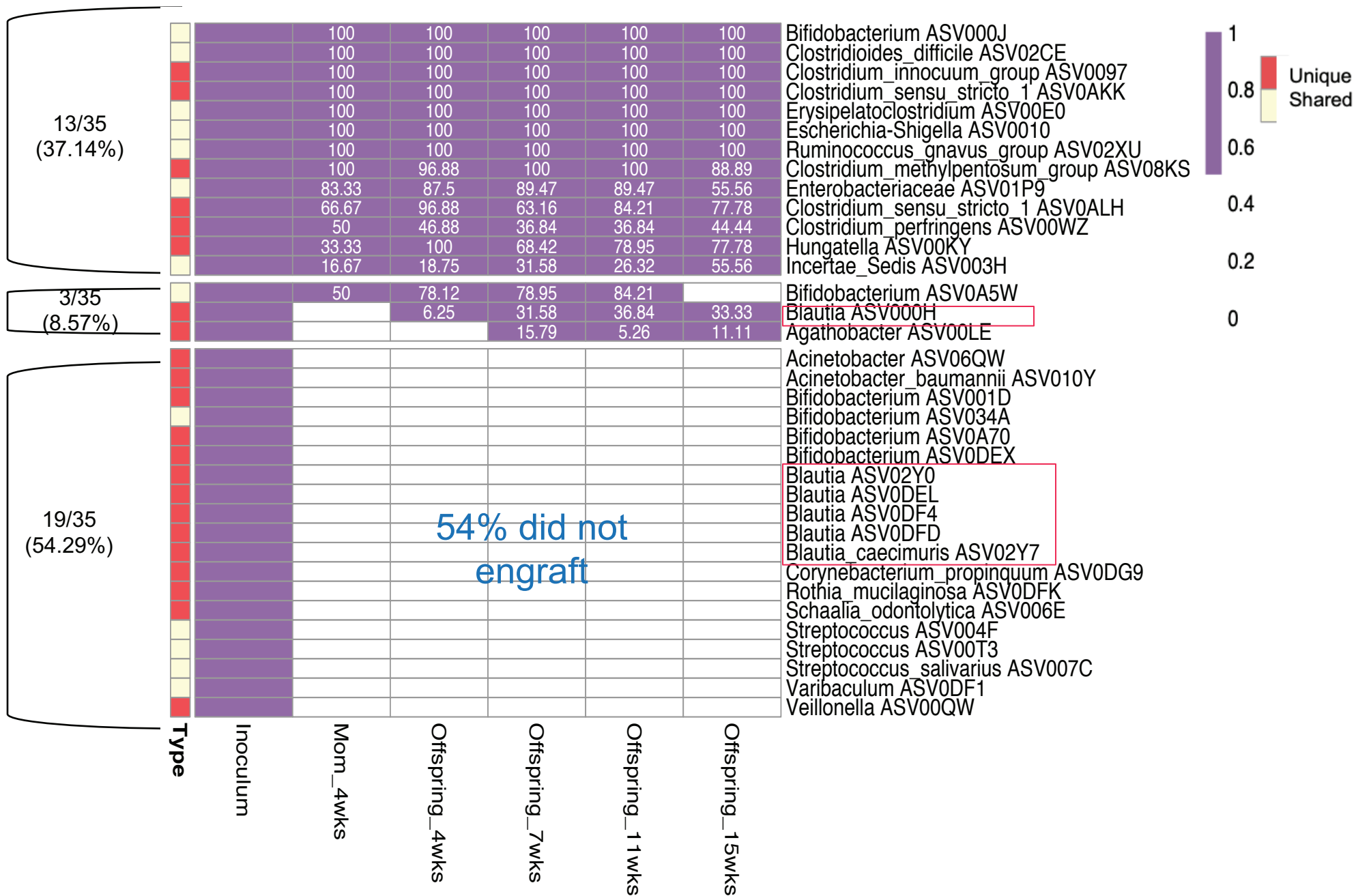
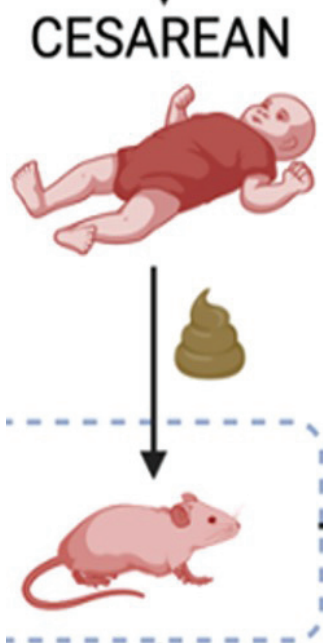
Fecal diversity in 4 week-old offspring mice transplanted with human baby feces



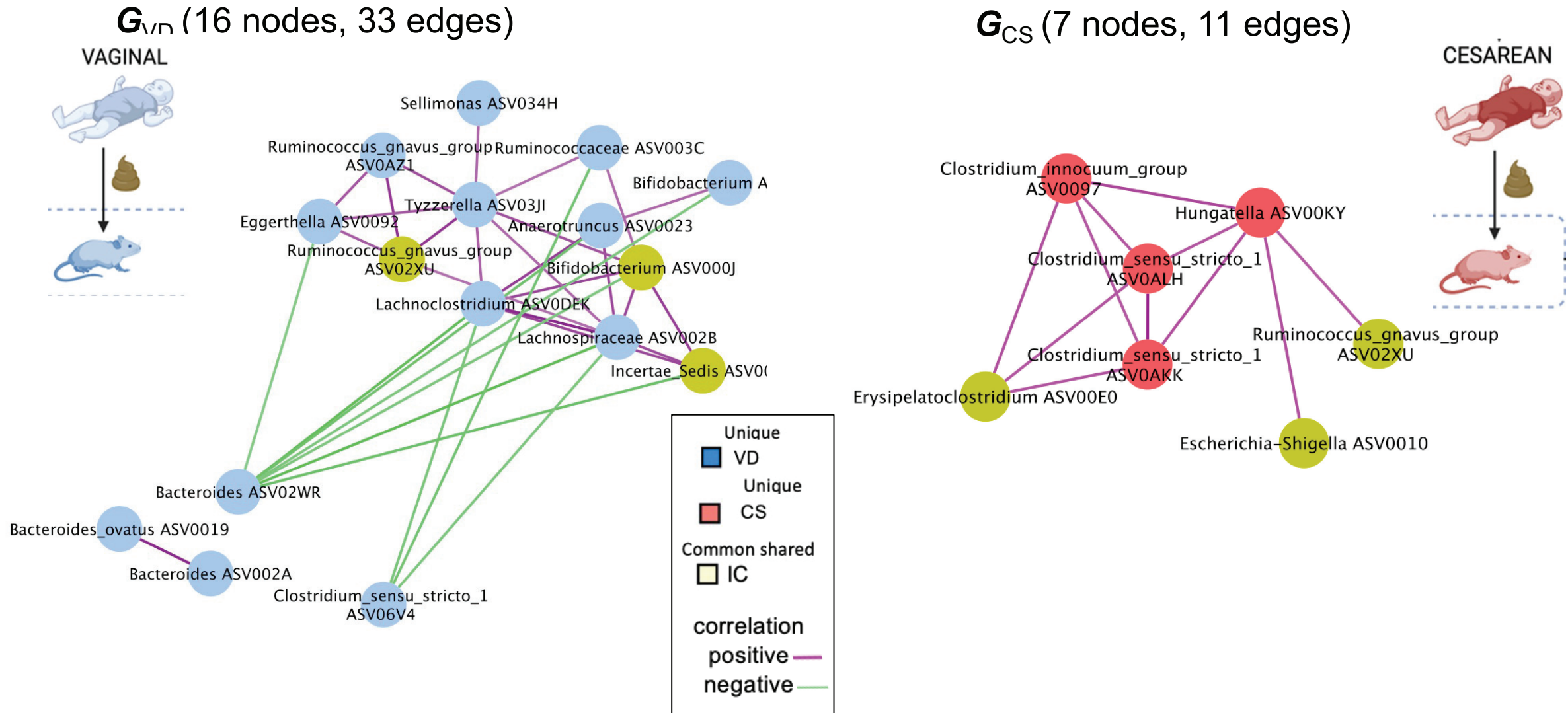
70% mouse bacterial engraftment from vaginally-born babies



46% mouse bacterial engraftment from C-section-born babies

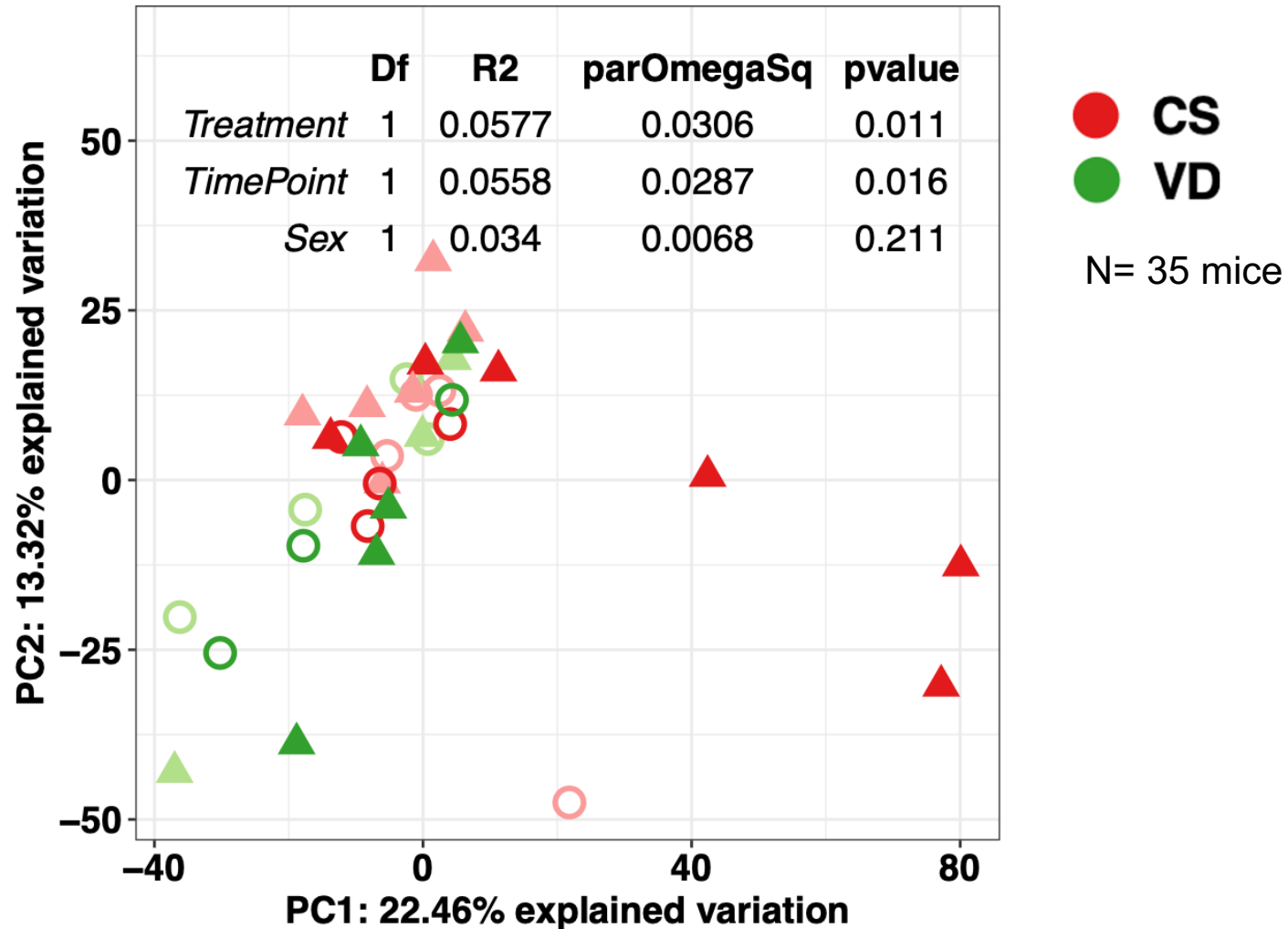


Simpler co-abundance network of human bacterial ASVs from the C-section feces mouse group



Repeated measurement correlation based on log-transformed ASV abundance. Correlations with BH adjusted $p < 0.05$ were kept in the network.

Ileum gene expression in mice receiving feces from vaginally or C-section born human babies



Inoculum has a significant effect

KEGG ileal gene pathways in mice at week 4 post-gavage

CESAREAN

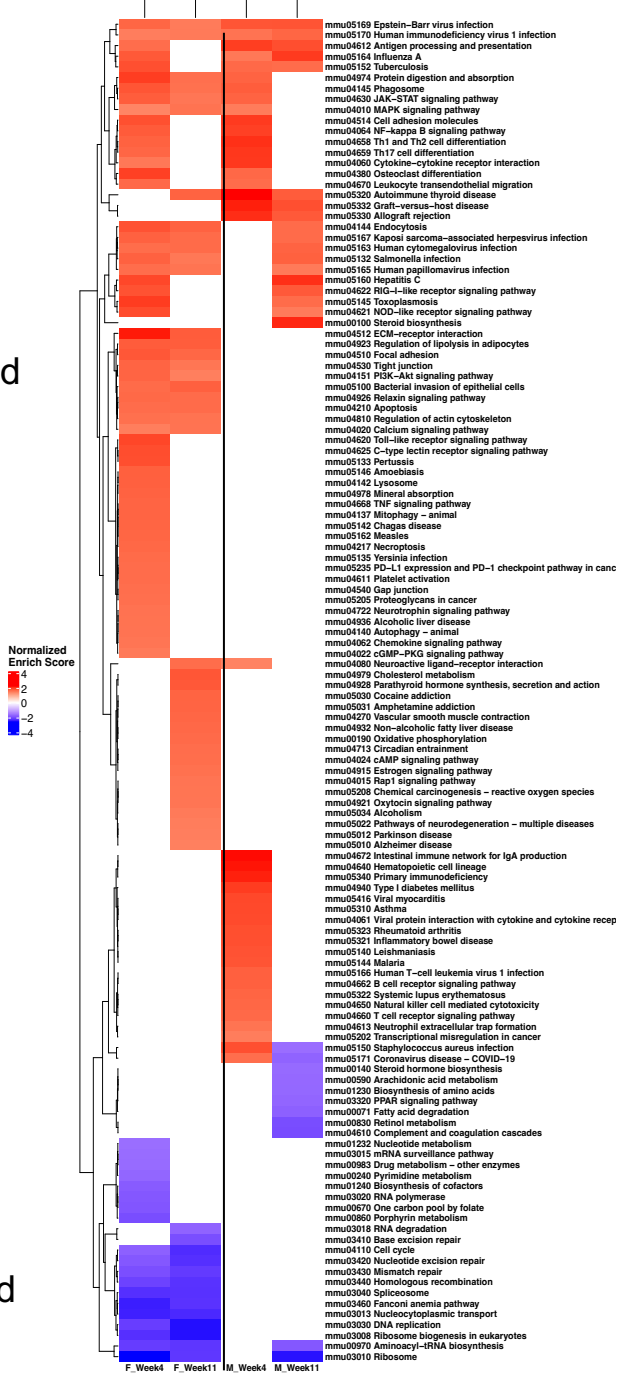


CS enriched

VAGINAL



VD enriched



Tuberculosis, Influenza A, **Antigen processing and presentation**, Endocytosis, Salmonella **infection**, Human papillomavirus **infection**, Kaposi sarcoma-associated herpesvirus **infection**, Human cytomegalovirus **infection**, Epstein-Barr virus **infection**, Human immunodeficiency virus 1 **infection**, **Phagosome**, Protein digestion and absorption, JAK-STAT signaling pathway, MAPK signaling pathway, Autoimmune thyroid disease

Cell cycle, DNA replication, Nucleocytoplasmic transport, Spliceosome, Fanconi anemia pathway, Ribosome biogenesis in eukaryotes, Homologous recombination, Mismatch repair, Nucleotide excision repair, Ribosome, Aminoacyl-tRNA biosynthesis

T cell profiles in mice transplanted baby microbiotas at w6

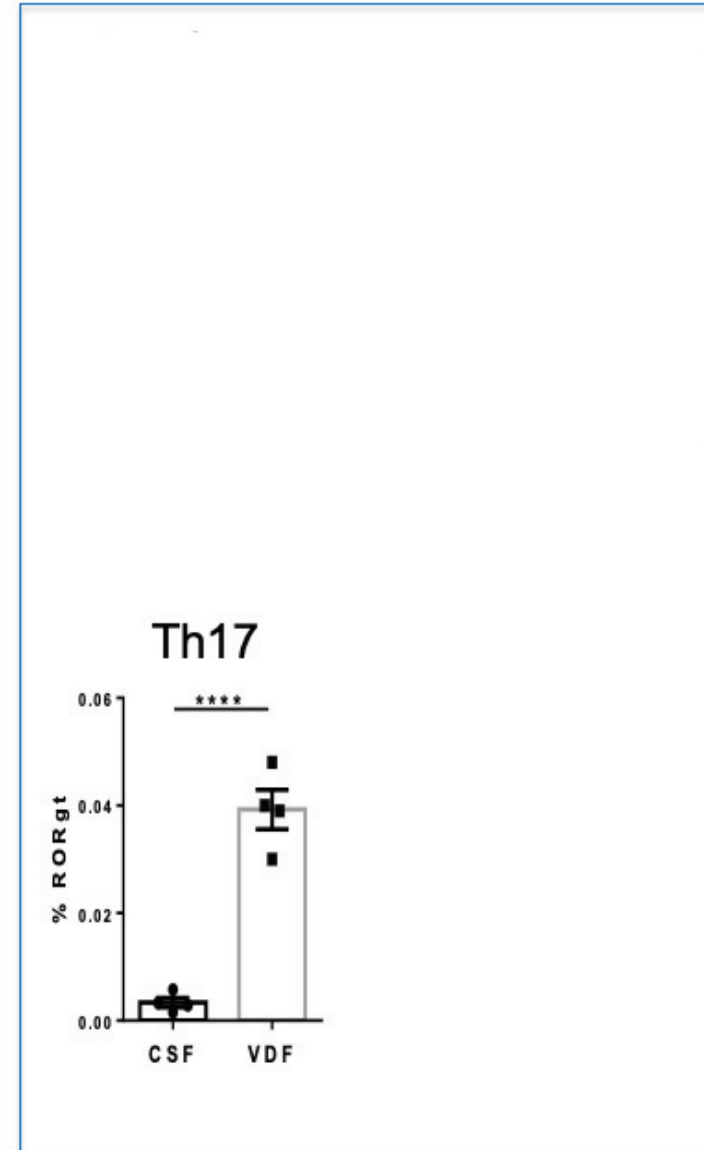
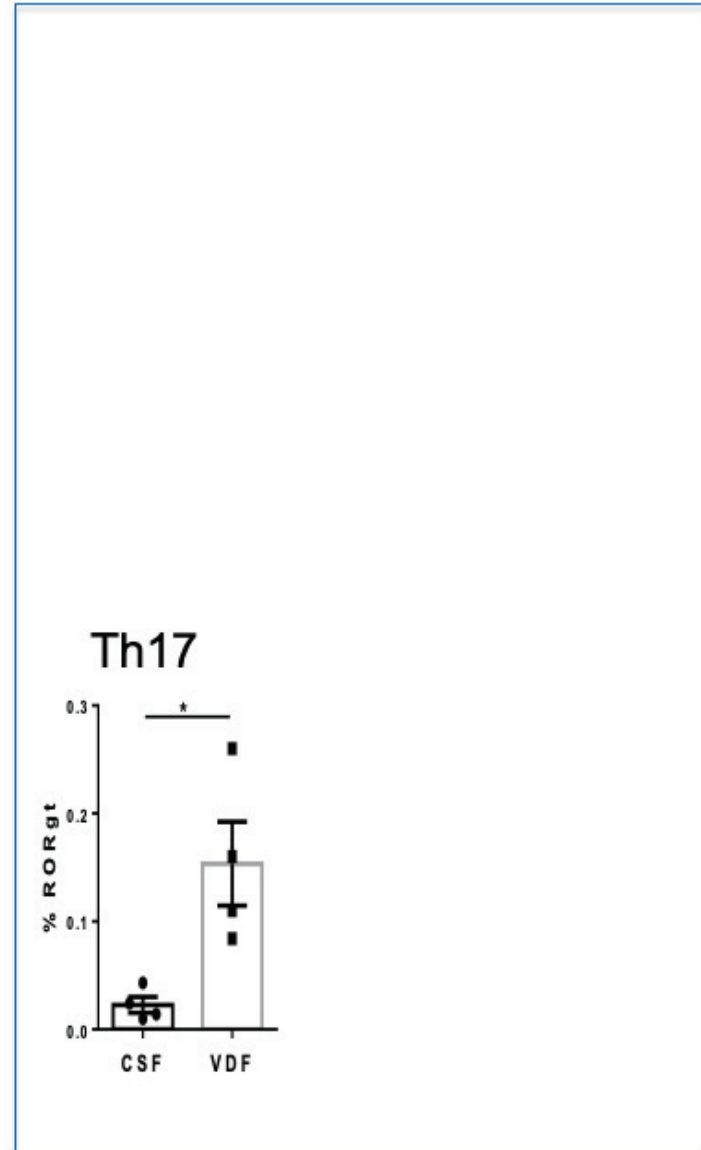
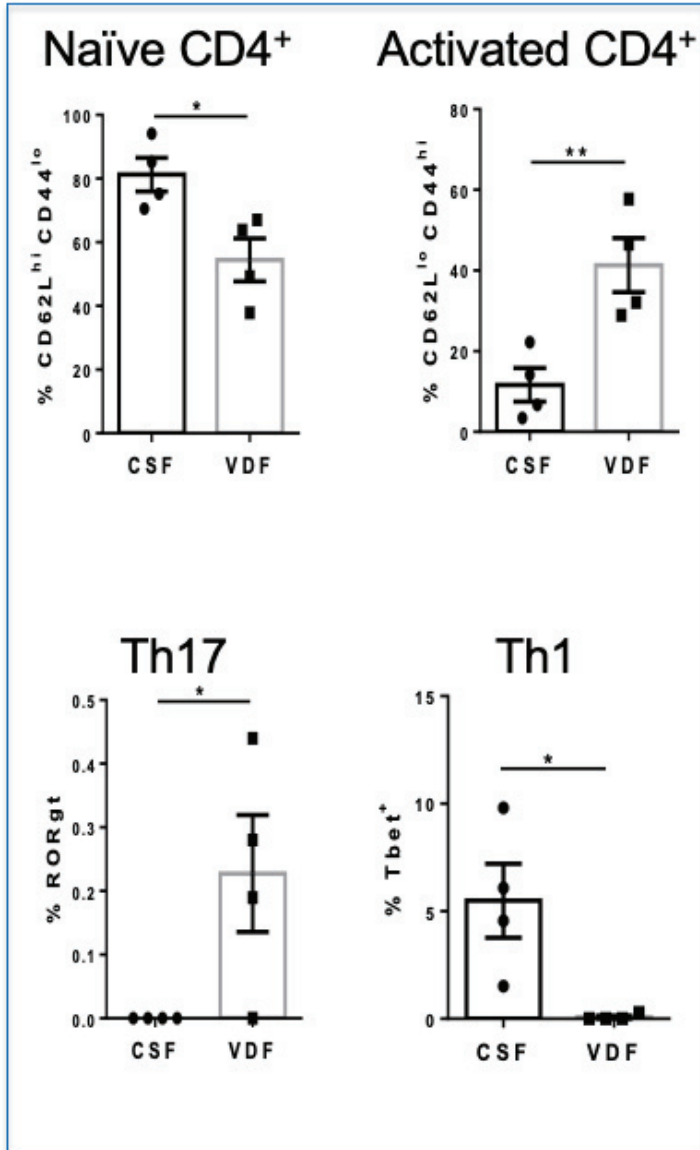


Derek Sant'Angelo

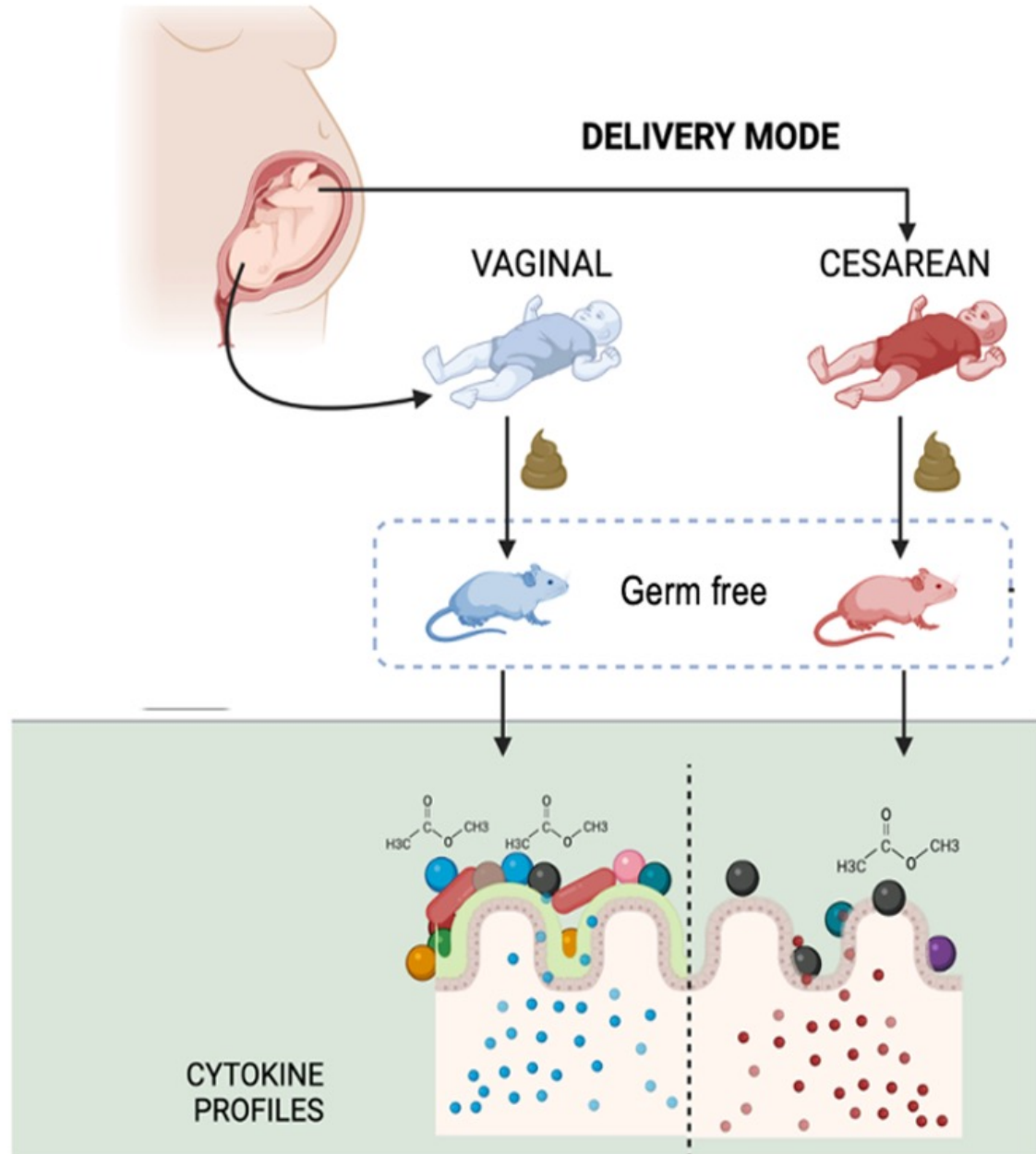
Colonic lamina propria

Inguinal lymph nodes

Spleen



Summary



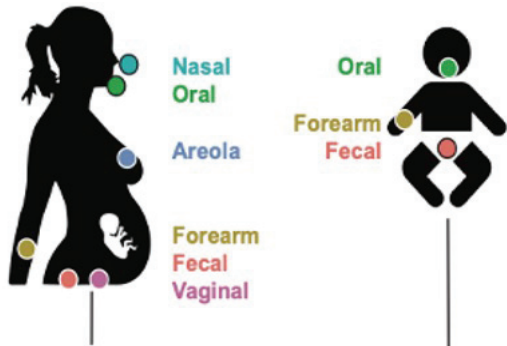
In relation to feces from vaginally born, feces from C-section human babies caused in transplanted mice:

- Different bacterial engraftment
- Altered intestinal gene expression
- Altered immune responses towards low tolerance/high inflammation

Vaginal seeding after C-section



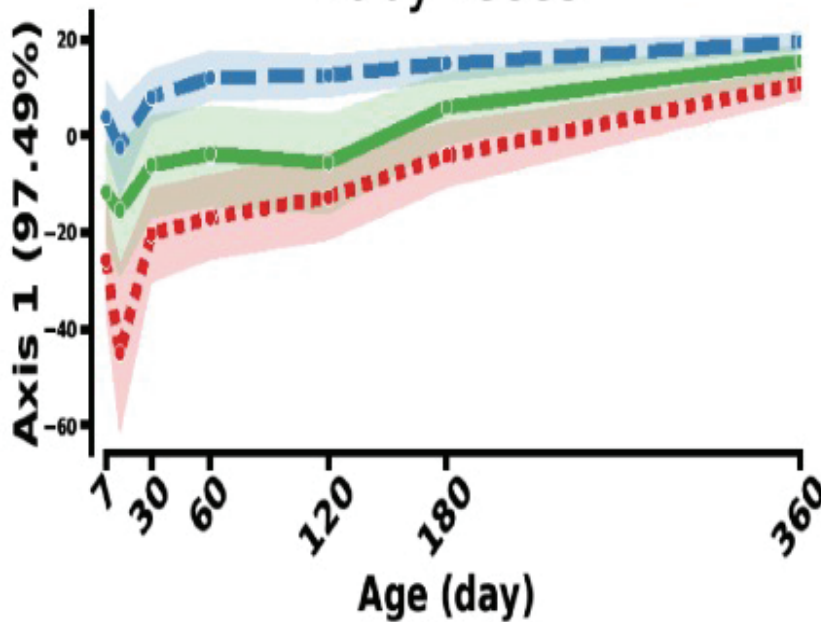
Microbiome trajectories by birth mode and exposure



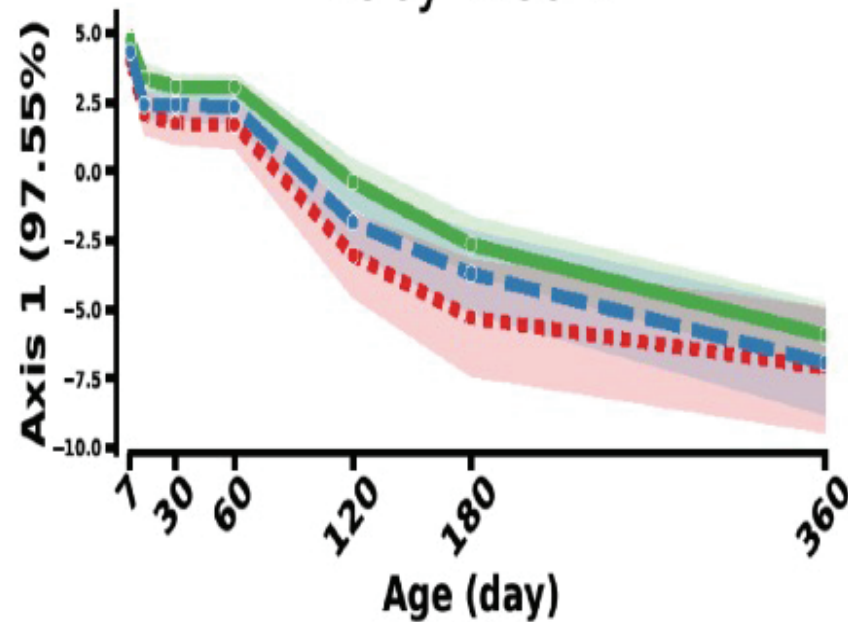
VD 
CS 
CSR 

Compositional Tensor Factorization (CFT) analyses

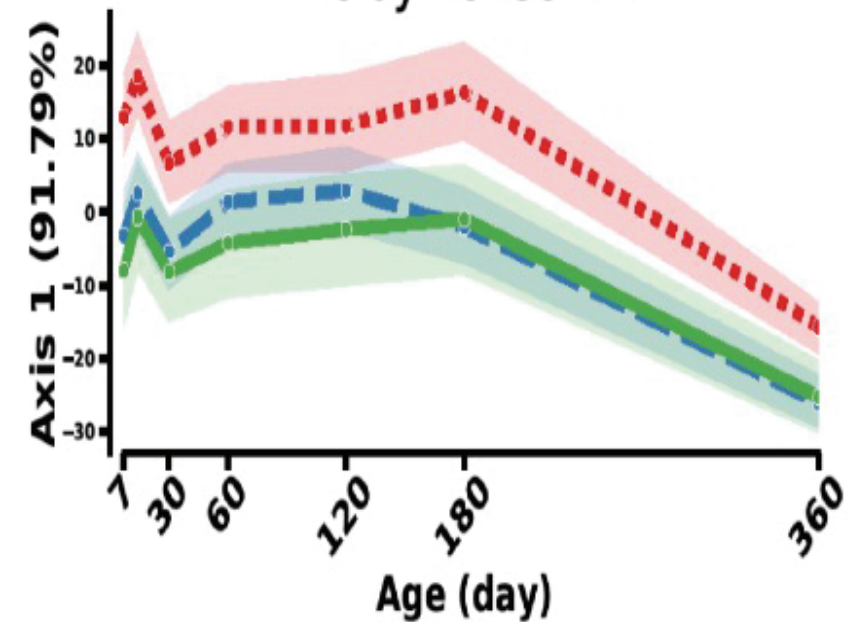
Baby-feces



Baby-mouth



Baby-forearm



Primordial Restoration of Infant Microbiome and Development

PRIMED

Effect of vaginal seeding in Cesarean-delivered infants
BMI-for-age z score over the first 3 years of life.

Randomized trial, 600 infants

PIs: Larry Appel (PI)

ClinicalTrials.gov Identifier: NCT03298334



Allergy Prevention in C-secTion Infants ViA Microbiome TransfEr

ACTIVATE

Effect of vaginal seeding in Cesarean
born infants on the microbiome and
allergic sensitization to food allergens

120 infants, 80 born by C-section with
randomized exposure (pilot)

PIs: Hugh Sampson and Jose
Clemente

ClinicalTrials.gov Identifier: NCT03567707



Randomized clinical trials

Restoration of Microbiota in Newborns

RoMaNs

Vaginal-fecal exposure at birth in CS born babies,
microbiota and systemic / mucosal immune
responses during infancy (up to 4 years).

Randomized trial, 200 infants

PIs: Lars Engstrand (PI)

ClinicalTrials.gov Identifier: NCT03928431



DATE: 10/10/2024

222

220

THE FUTURE IS

Restoration

10/10/2024

10/10/2024

10/10/2024

10/10/2024

10/10/2024

DATE:



DATE: 10/10/2024

DATE: 10/10/2024



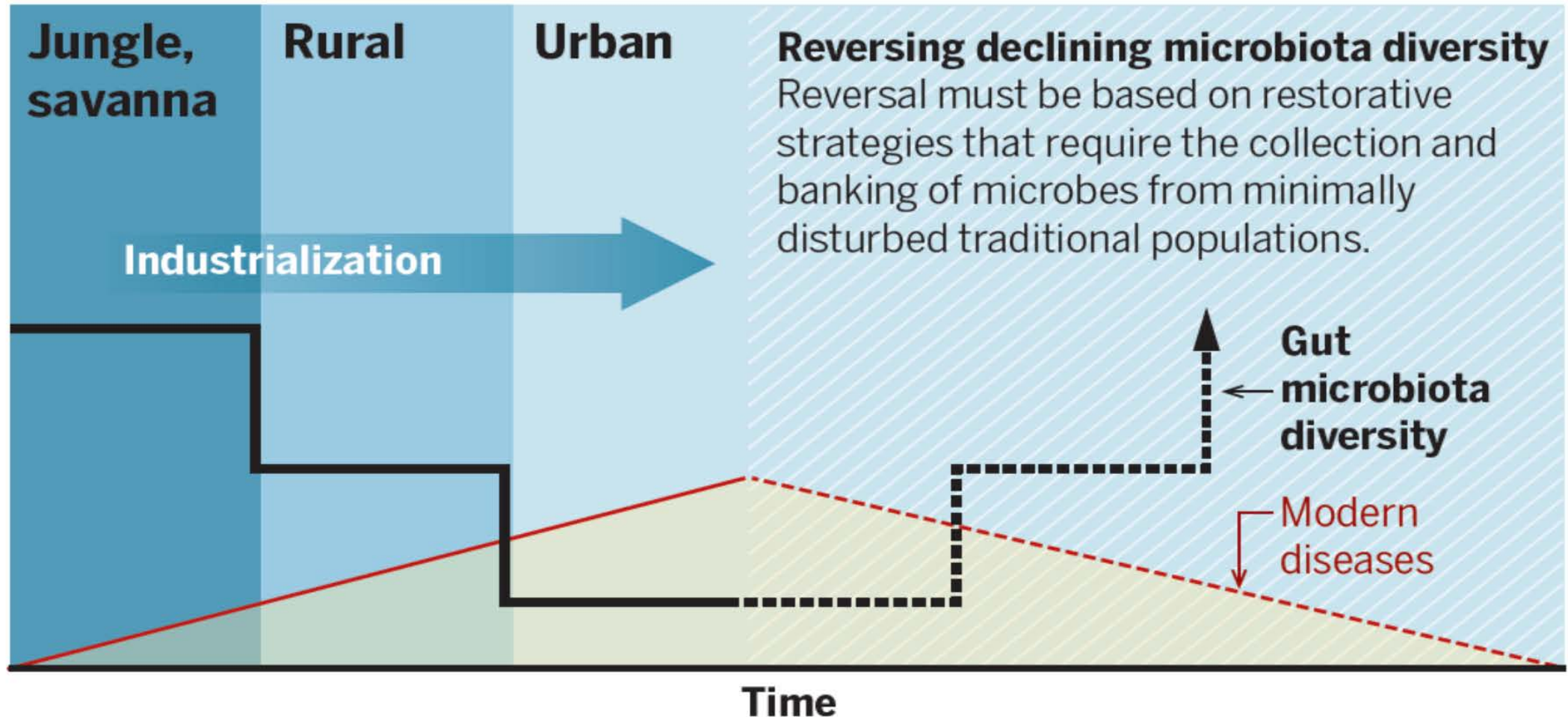
Created by AI

Preserving microbial diversity

Science 2018

Microbiota from humans of all cultures are needed to ensure the health of future generations

By Maria G. Dominguez Bello¹, Rob Knight²,
Jack A. Gilbert³, Martin J. Blaser⁴

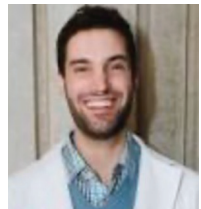


Thank you!

THE
MICROBIOTA
VAULT



Haipeng Sun

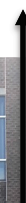


Jeremy Lessing

Jincheng Wang



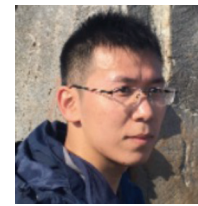
Melissa Woortman



Wessam Mohamed



Emanuel DiCicco



Gary Wu



Liping Zhao



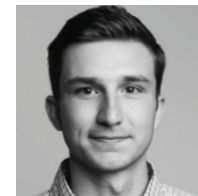
Martin Blaser



Troy Roepke



Pieter Dorrestein
UCSD



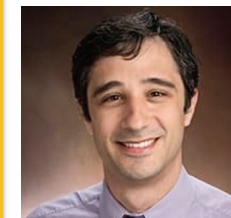
Cameron Martino
UCSD



Se Jin Song
Univ Colorado



Rob Knight
UCSD



Kyle Bittinger
PennCHOP



Dan Scott
PennCHOP



Oscar Noya-Alarcon
UCV



Monica Contreras
IVIC



Daniela Vargas
UNAM

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