



# Diet, Microbes & Mood

Jane A. Foster, Ph.D.

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# “All diseases begin in the gut.”

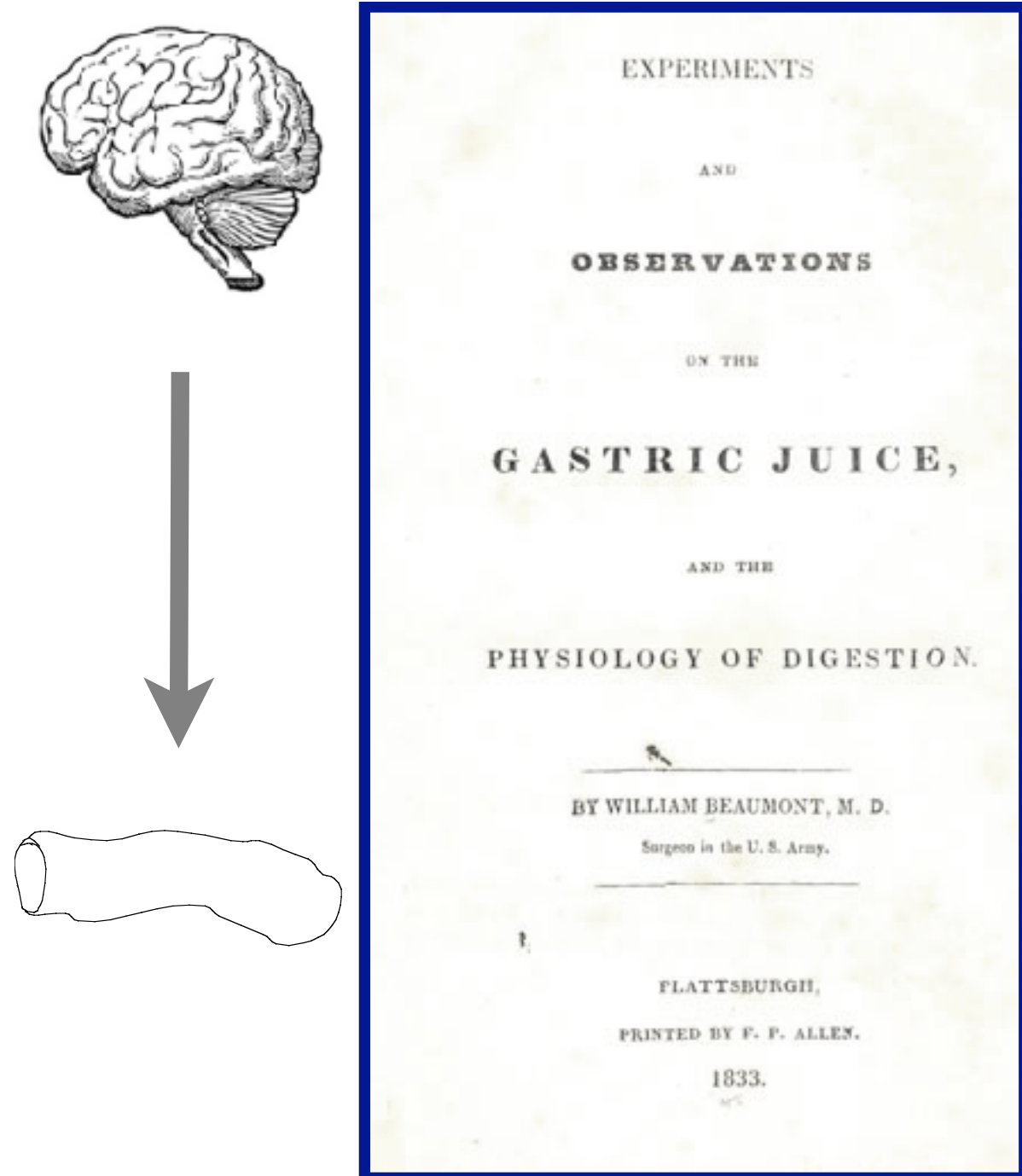
*-Hippocrates*



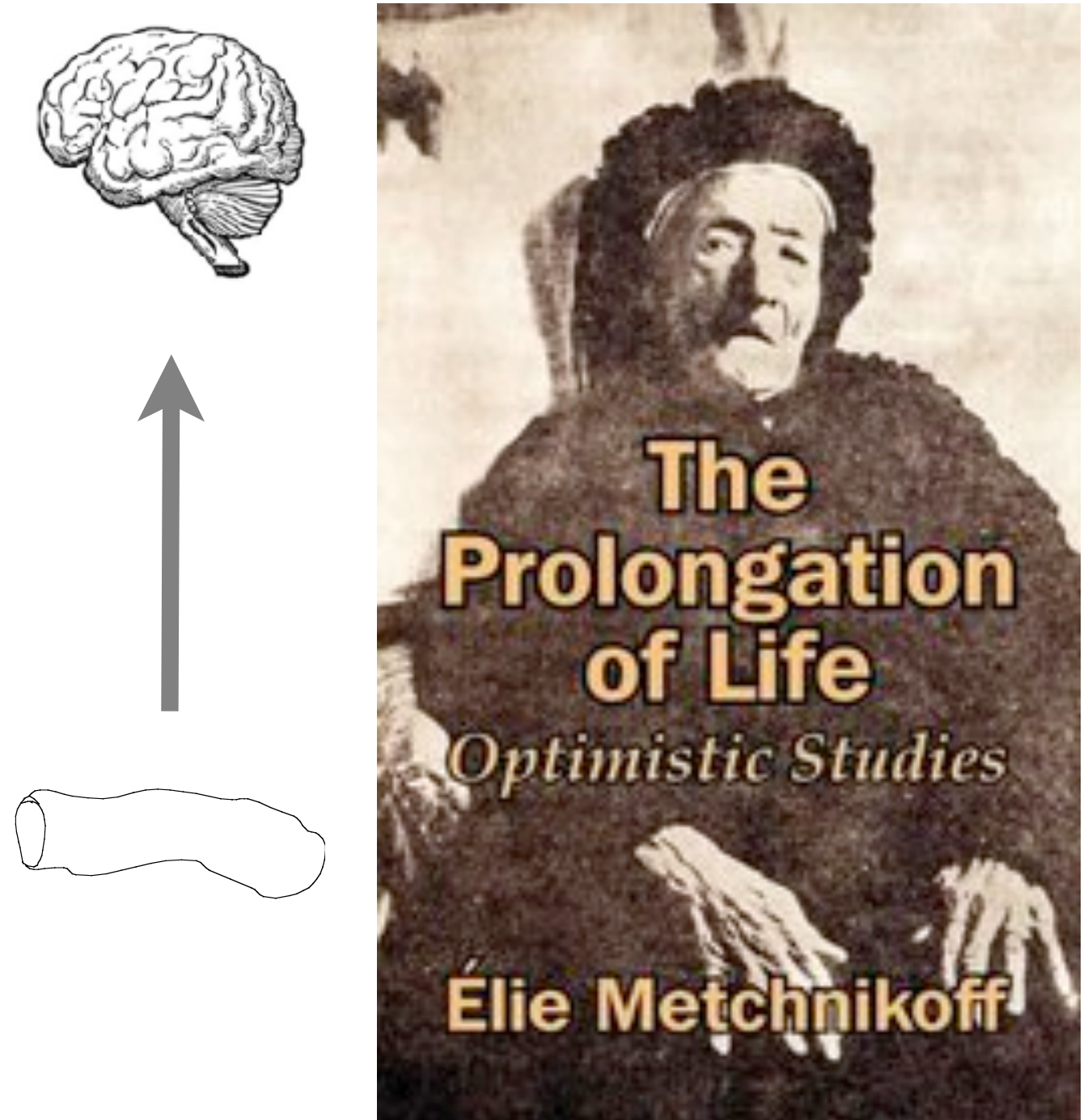


# Gut-brain Axis

1833 - Beaumont



1907 - Metchnikoff



# Fermented Foods to probiotics...



- Metchnikoff inspired Minoru Shirota to investigate the connection between bacteria and good GI health
- Shirota is the inventor of Yakult - the yogurt-like probiotic drink containing *Lactobacillus casei* strain Shirota - 1930

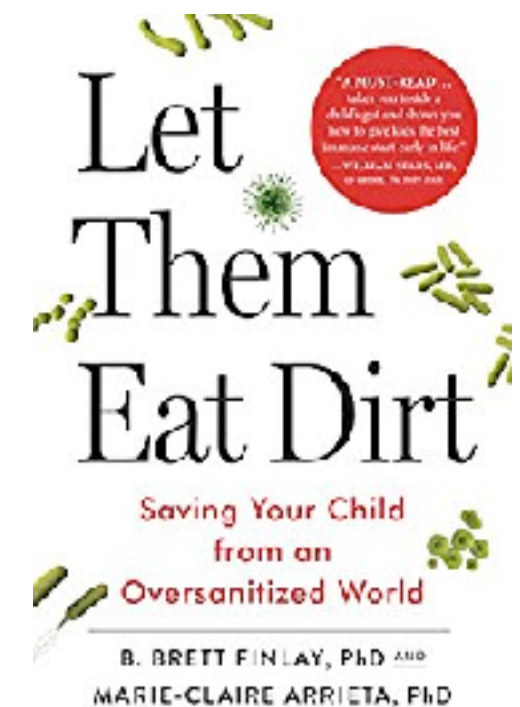
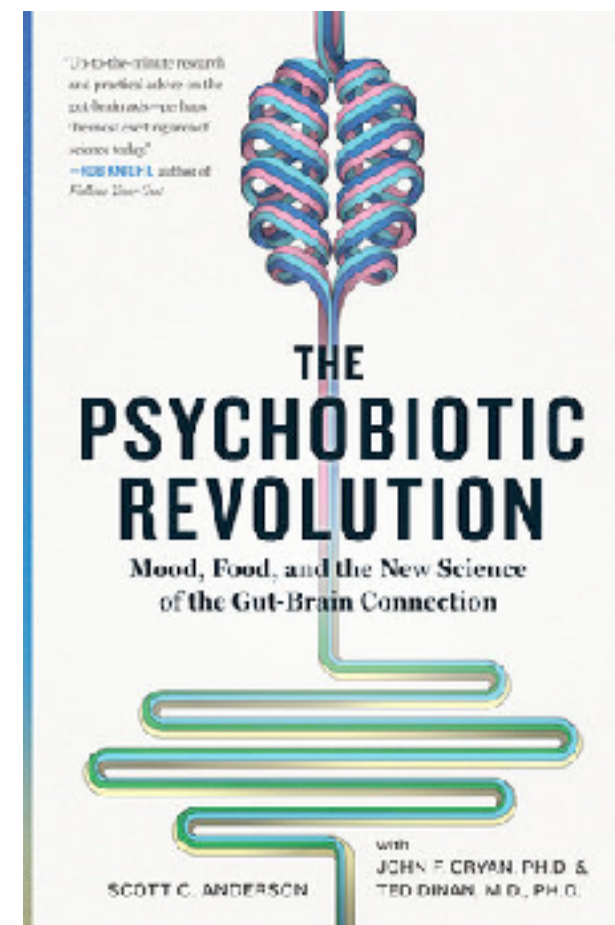
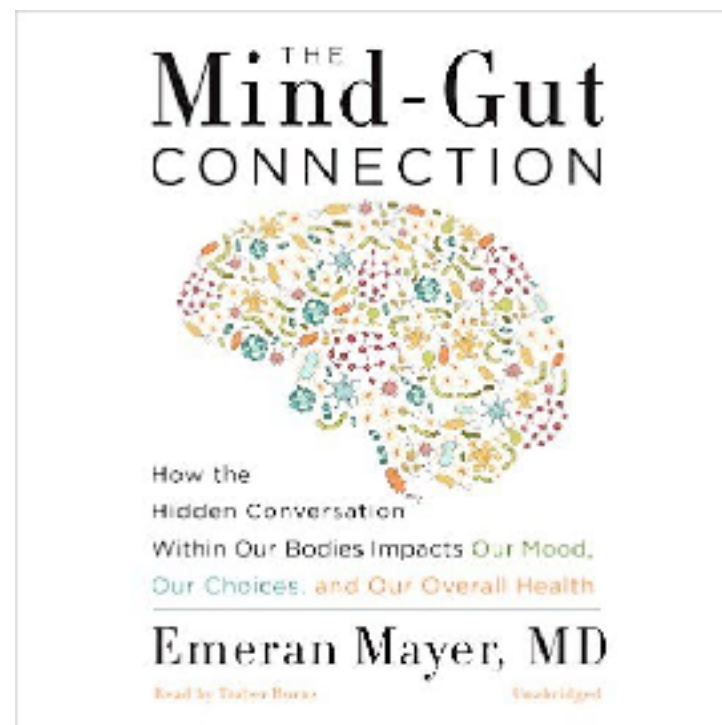
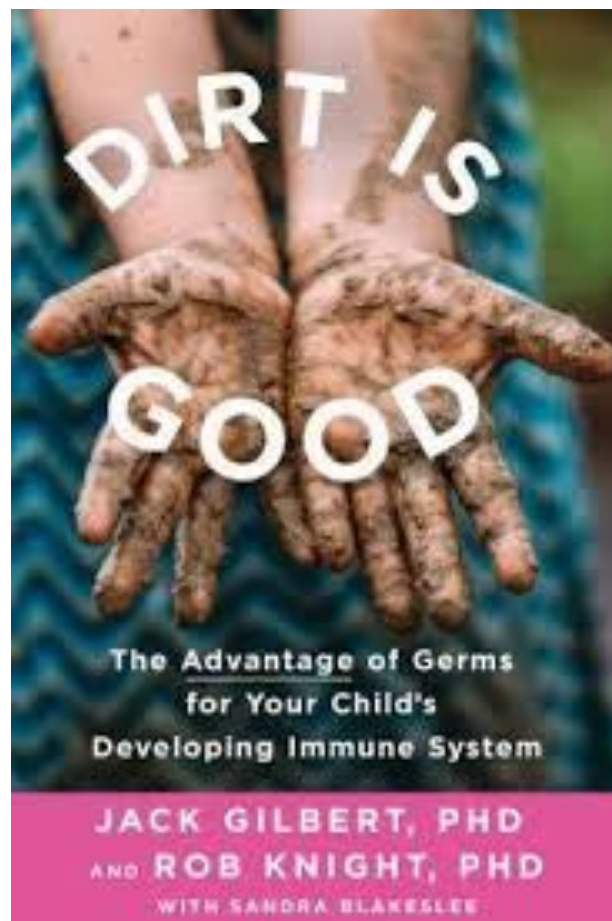
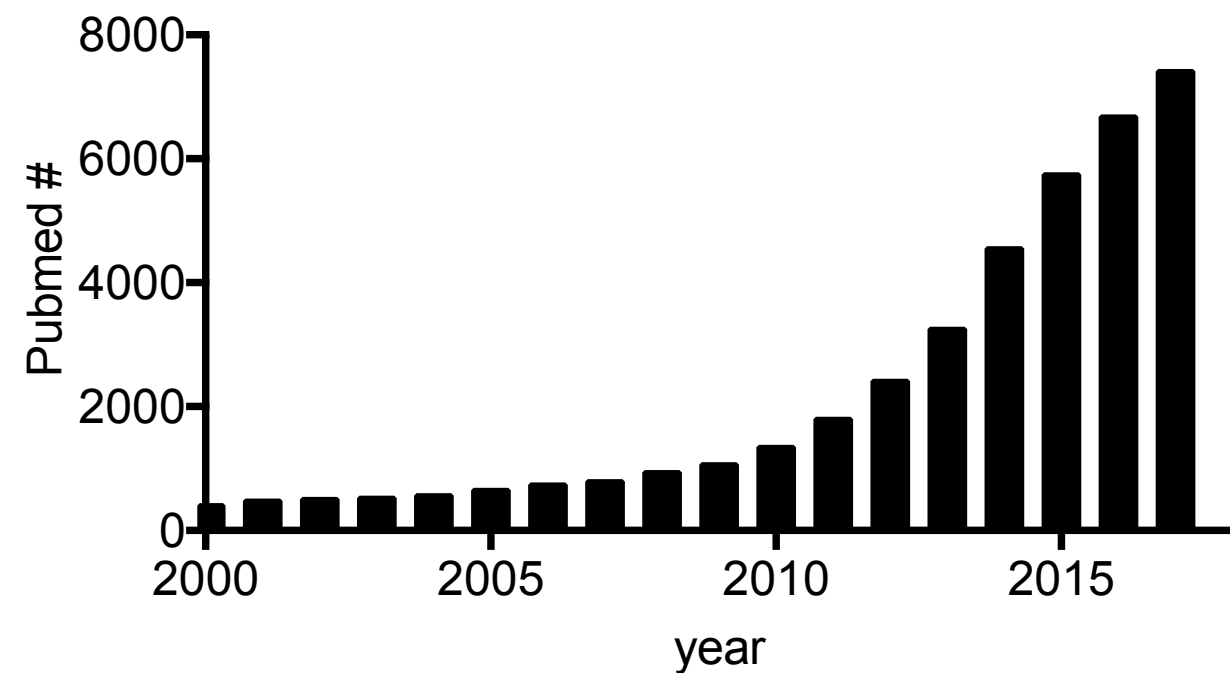
**Probiotics are “live micro-organisms that when administered in adequate amounts, confer a health benefit on the host”**

World Health Organization (WHO), Food and Agricultural Organization (FAO) 2015

“microbiota” or “microflora”

🔥 **HOT TOPIC** 🔥

## Microbiome in health and disease





## Targeting the Microbiome for Mental Health: Hype or Hope?

Jane A. Foster

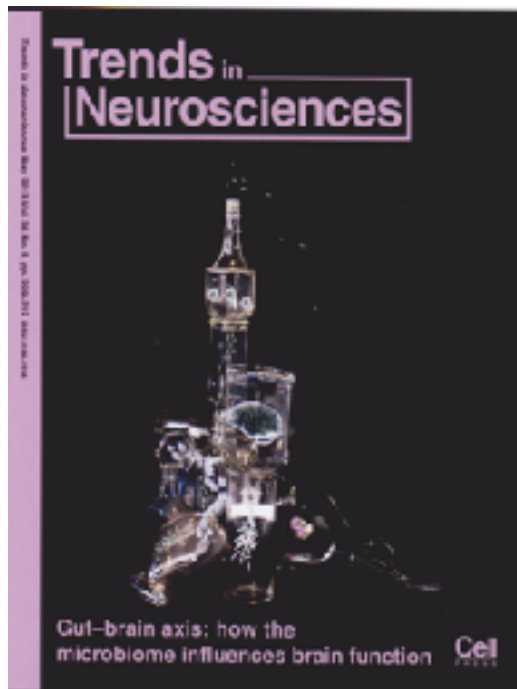
Key Questions for Today :

- What do we know about the microbiome?
- What is the evidence that microbes influence mood?
- How does diet influence the microbiome?
- What are the opportunities for therapeutic development and precision medicine approaches?

# Microbes - microbiome - microbiota



- All of the surfaces of your body are covered with microbes
- Microbes include bacteria, viruses, fungi, protozoa, and parasites
- "Microbiome" refers to all of the microbes and their related genetic material
- "Microbiota" refers to the microbes themselves
- Much of the research focused on gut microbiota - also referred to as commensal bacteria

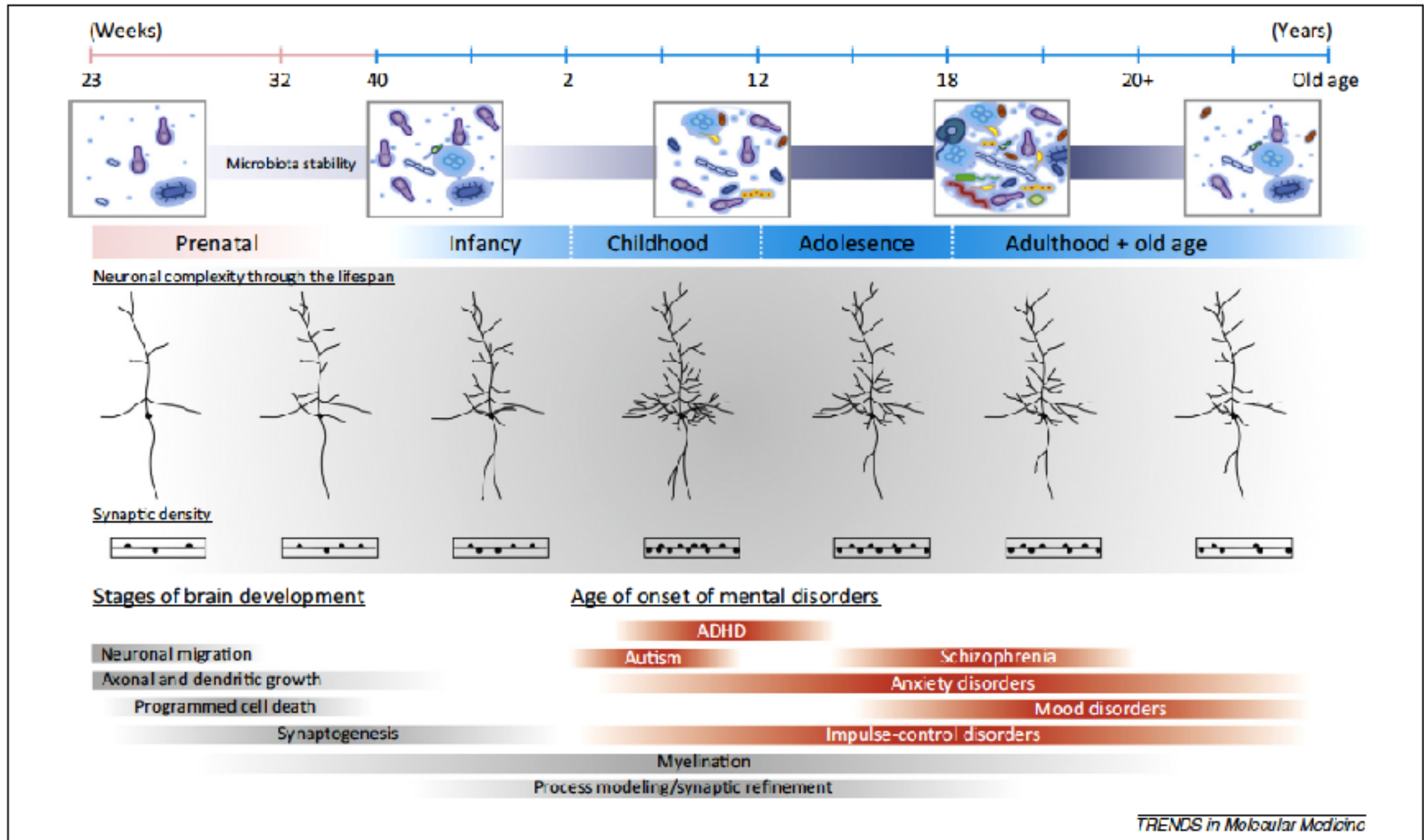


# What do we know about gut microbiota?

- The gastrointestinal tract of an adult human contains 100 trillion viable bacteria
- Exposure to microbes and colonization occurs primarily at birth and continues through development
- Microbiota are essential to pathogen defence, nutrient uptake, and metabolism
- Microbiota are essential to the development and function of the immune system
- Recent evidence shows that microbiota-brain communication is important to healthy brain development

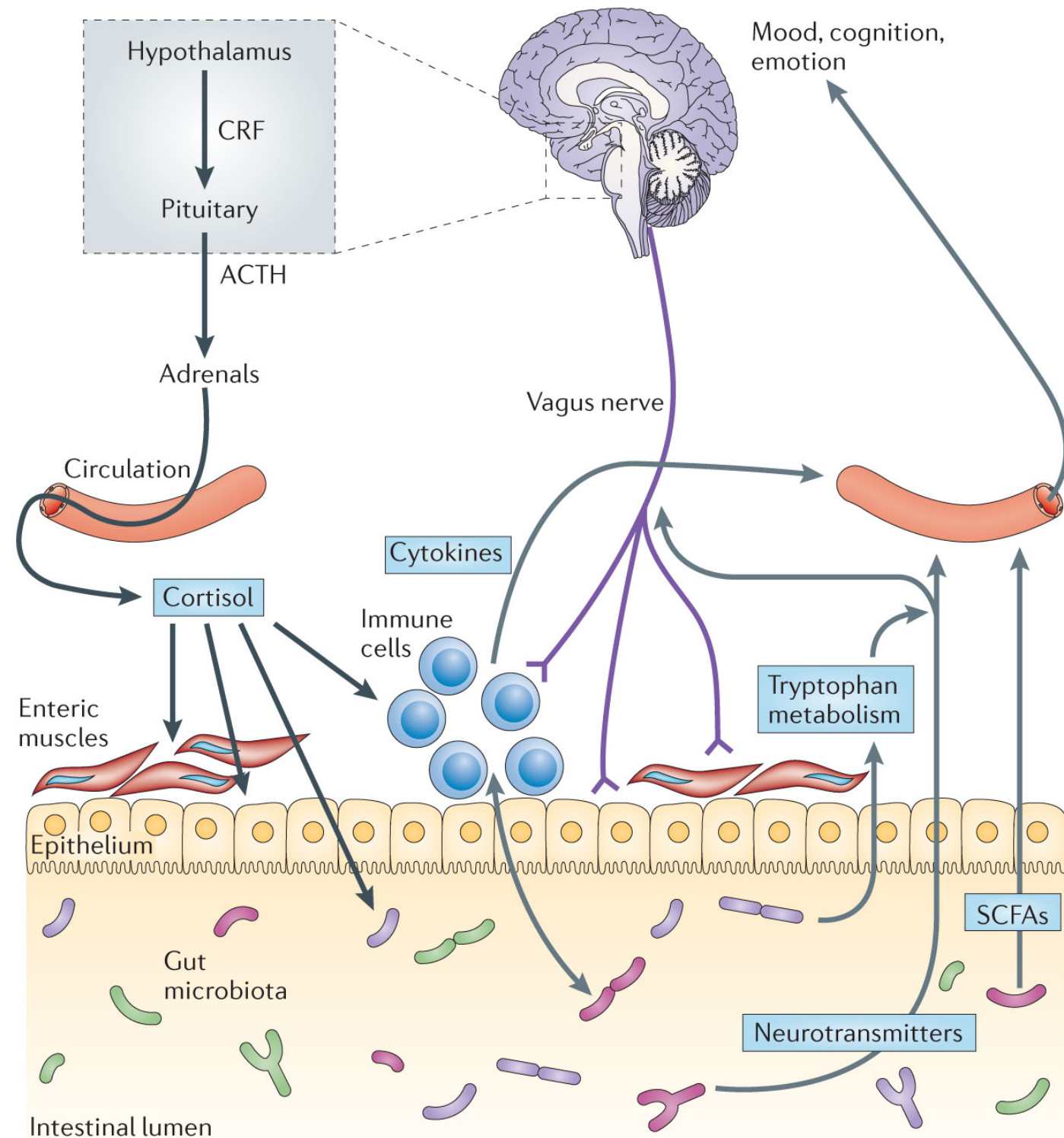


- In healthy infants, dynamic changes in microbiota composition and diversity over the first year of life - influenced by diet (breast vs bottle-fed) and mode of delivery (vaginal vs c-section)



# How do microbiota communicate with the brain?

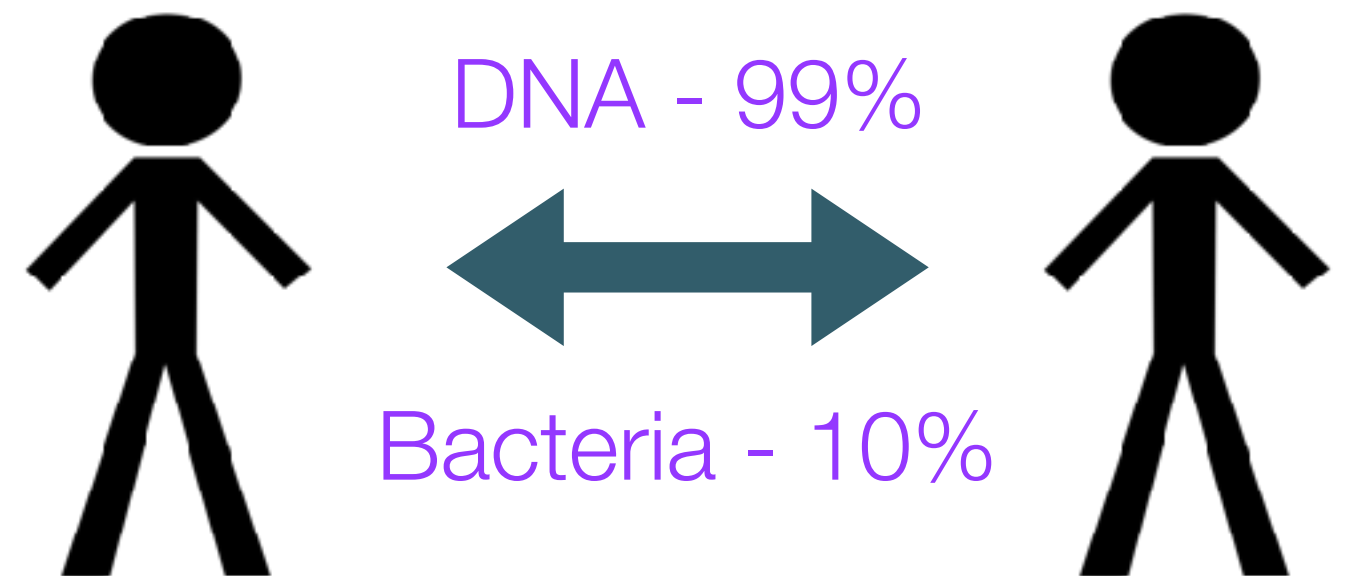
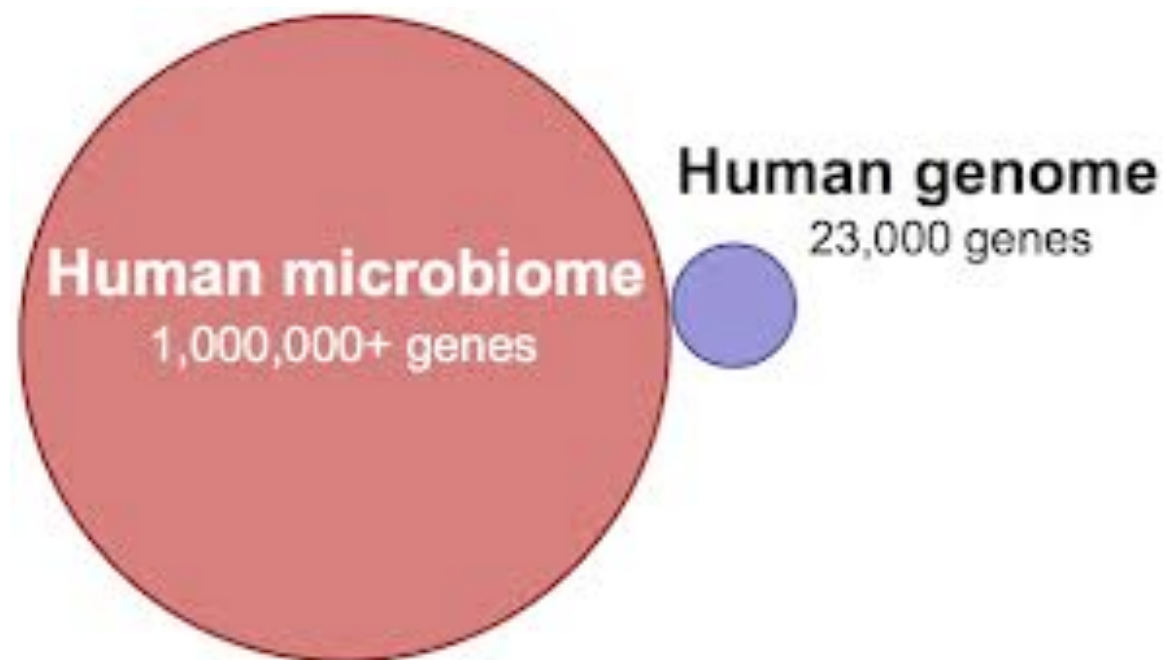
1. Neural
2. Humoral
3. Cellular
4. Metabolites
5. Neuroactive molecules



# Are We Really Vastly Outnumbered? Revisiting the Ratio of Bacterial to Host Cells in Humans

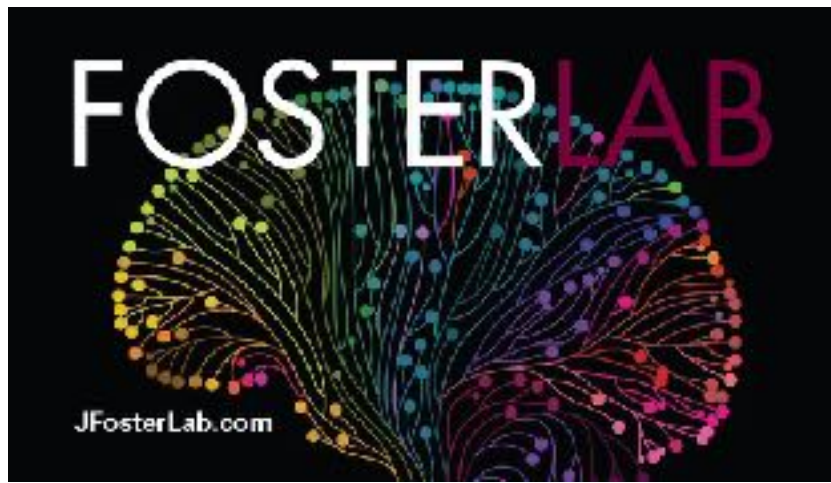


Ron Sender,<sup>1</sup> Shai Fuchs,<sup>2,3,\*</sup> and Ron Milo<sup>1,\*</sup>

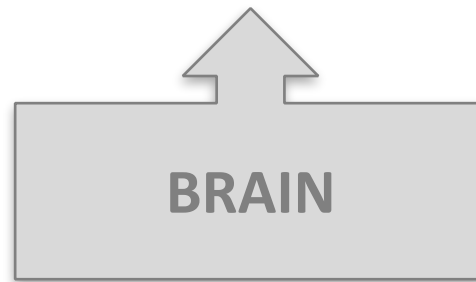
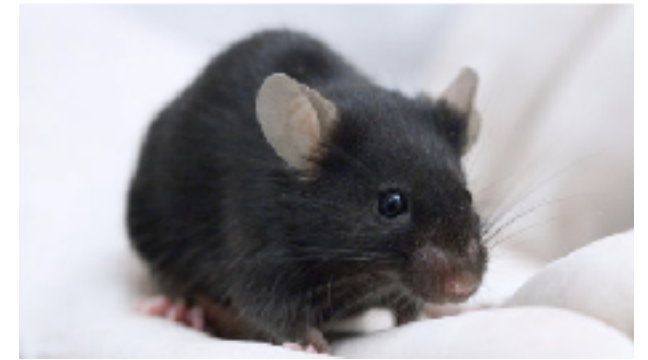


in healthy adults...

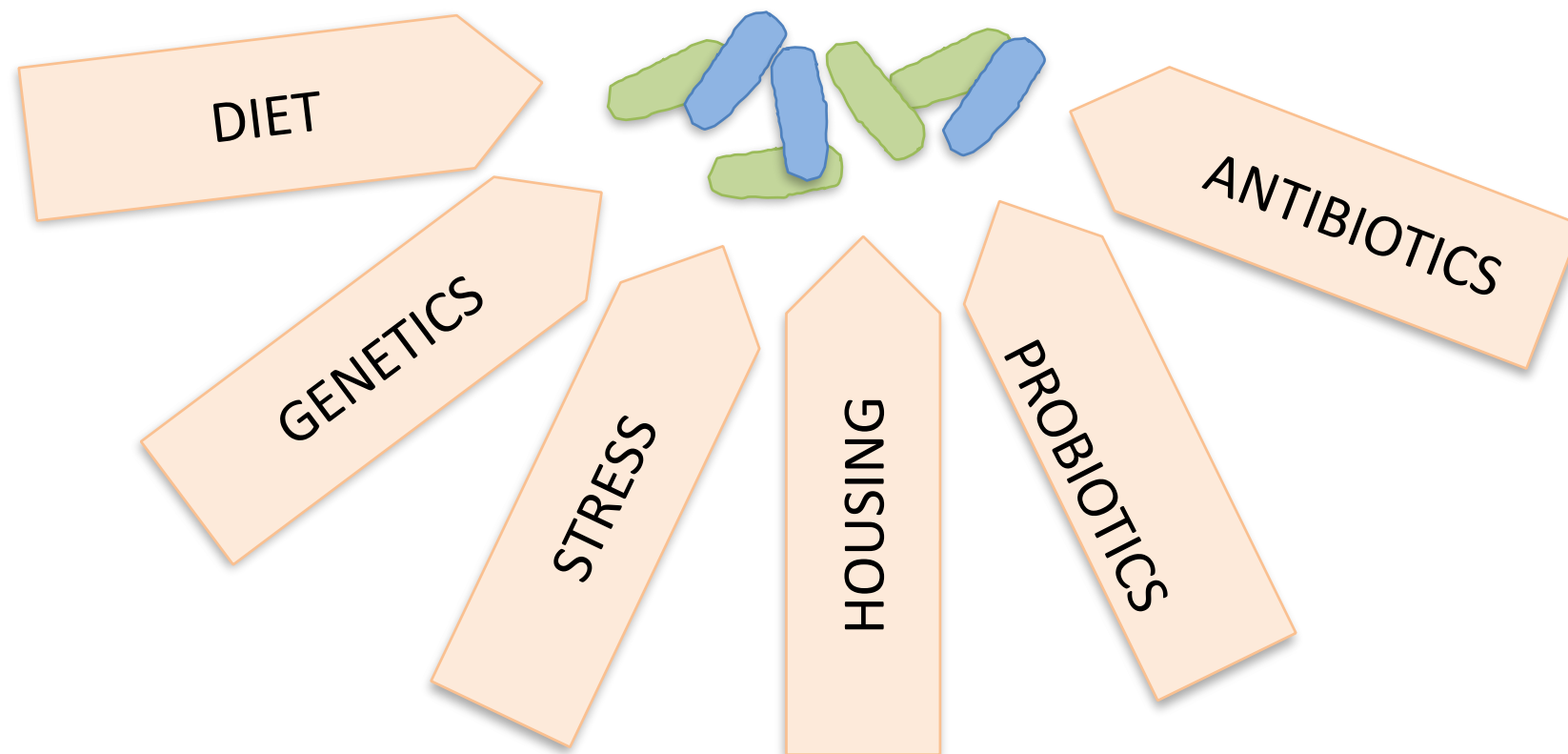




BEHAVIOUR



MICROBIOTA



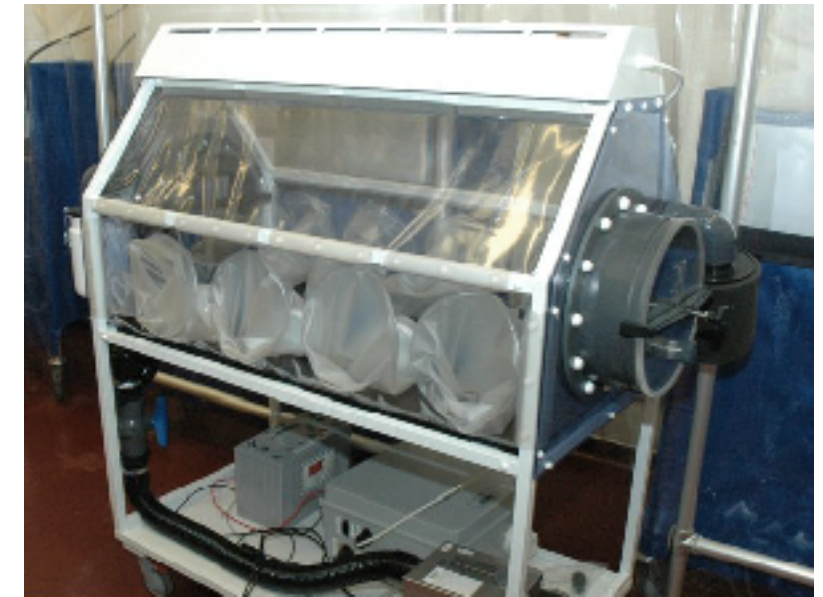
# The starting point...

## Postnatal microbial colonization programs the hypothalamic-pituitary-adrenal system for stress response in mice

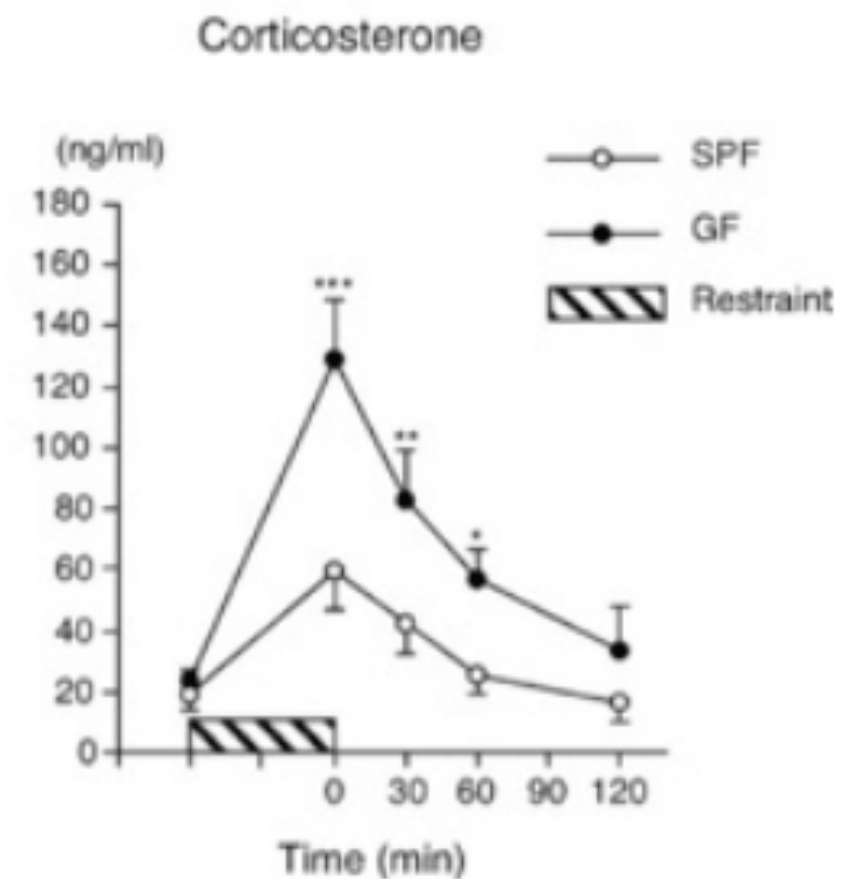
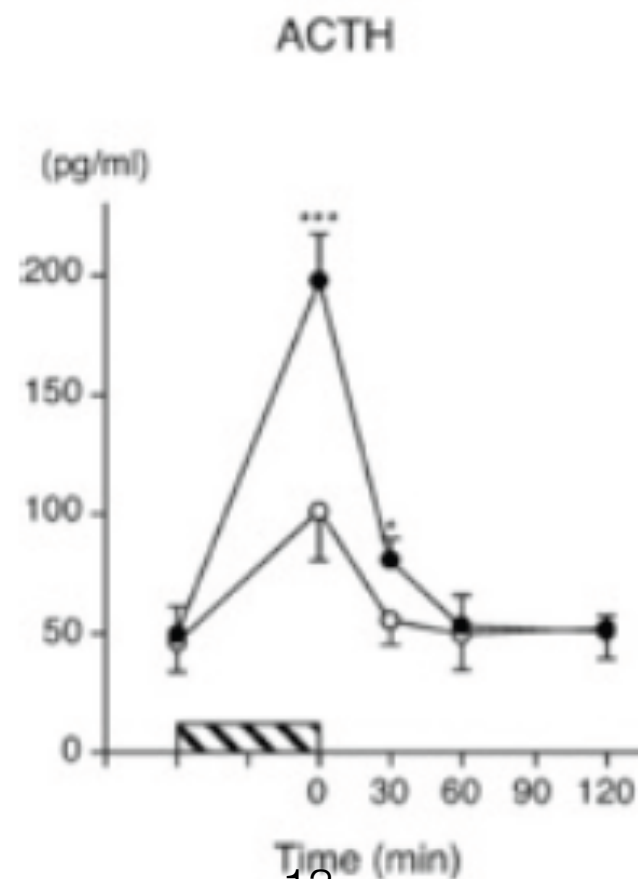
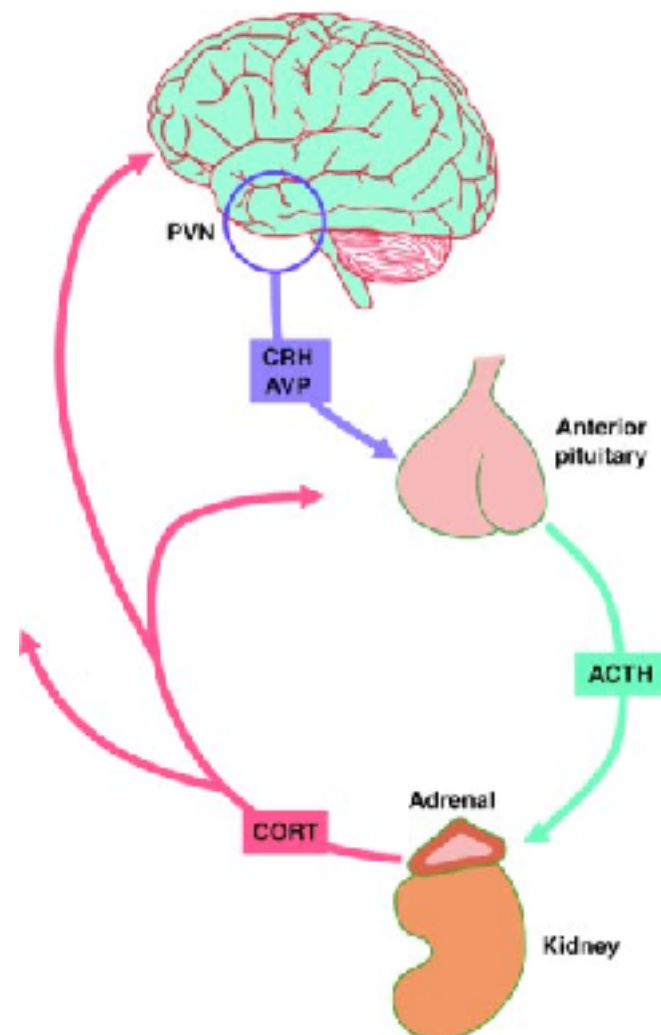
Sudo et al 2004 J Physiol 558: 263-275

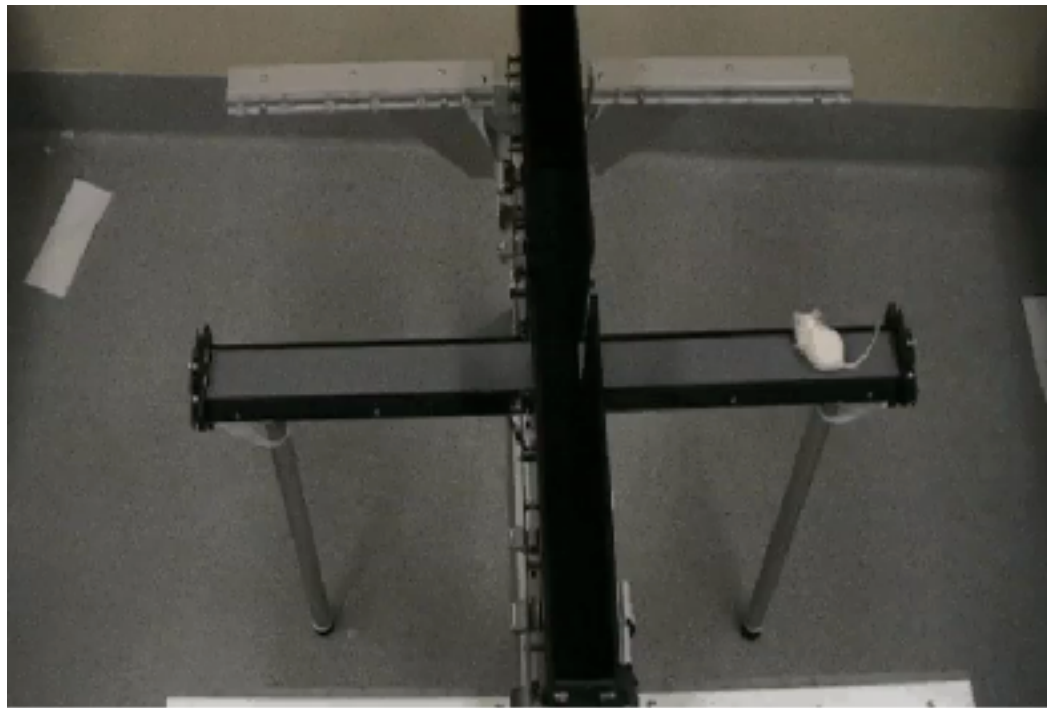
Nobuyuki Sudo<sup>1,2</sup>, Yoichi Chida<sup>1</sup>, Yuji Aiba<sup>3,4</sup>, Junko Sonoda<sup>1</sup>, Naomi Oyama<sup>1</sup>, Xiao-Nian Yu<sup>1</sup>, Chiharu Kubo<sup>1</sup> and Yasuhiro Koga<sup>3</sup>

<sup>1</sup>Department of Psychosomatic Medicine and <sup>2</sup>Department of Health Care Administration & Management, Graduate School of Medical Sciences, Kyushu University, Fukuoka, Japan, <sup>3</sup>Department of Infectious Diseases, Tokai University School of Medicine, Isehara, Kanagawa, Japan and <sup>4</sup>Wakama Pharmaceutical Co. Ltd, Ohi-machi, Kanagawa, Japan



**Germ-free mice showed exaggerated stress response**





## Reduced anxiety-like behavior and central neurochemical change in germ-free mice

K. M. NEUFELD,<sup>a,\*</sup>† N. KANG,<sup>a,‡</sup> J. BIENENSTOCK,<sup>a,§</sup> & J. A. POSTER,<sup>a,‡</sup>

PNAS

## Normal gut microbiota modulates brain development and behavior

Rochellys Diaz Heijtz<sup>a,b,1</sup>, Shugui Wang<sup>c</sup>, Farhana Anuar<sup>d</sup>, Yu Qian<sup>a,b</sup>, Britta Björkholm<sup>d</sup>, Annika Samuelsson<sup>d</sup>, Martin L. Hibberd<sup>c</sup>, Hans Forsberg<sup>b,e</sup>, and Sven Pettersson<sup>c,d,1</sup>

Molecular Psychiatry (2012), 1–8

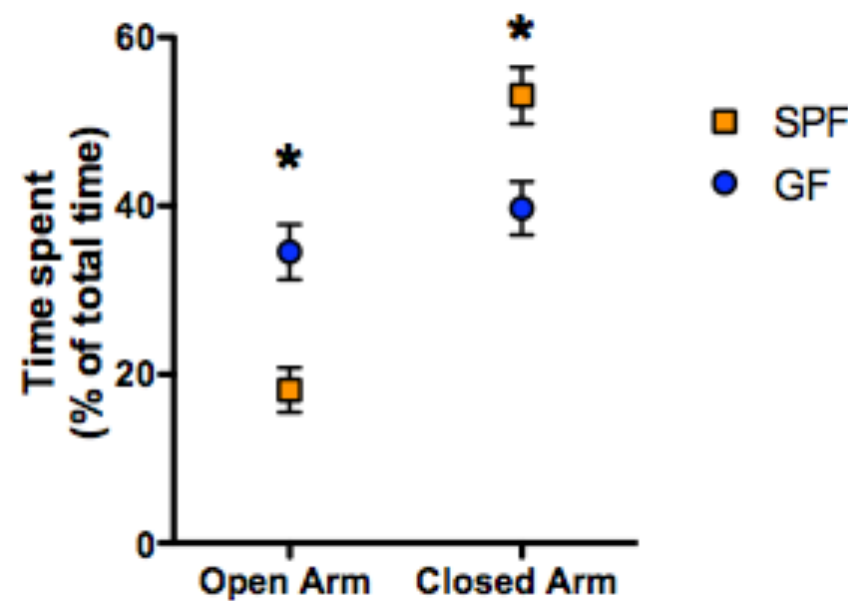
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www.nature.com/mp

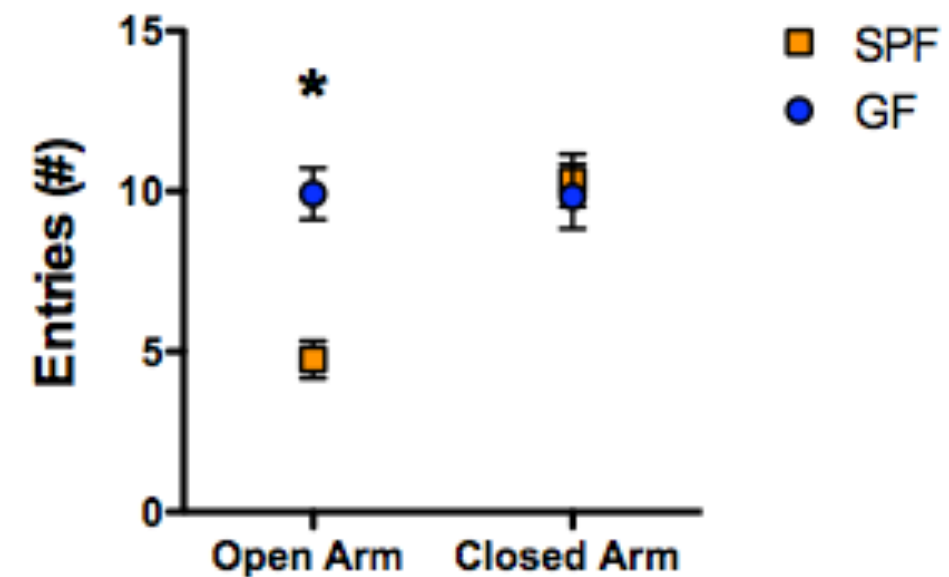
### ORIGINAL ARTICLE

## The microbiome-gut-brain axis during early life regulates the hippocampal serotonergic system in a sex-dependent manner

G. Clarke<sup>1,2</sup>, S. Grenham<sup>1</sup>, P. Scully<sup>1</sup>, P. Fitzgerald<sup>1</sup>, R.D. Moloney<sup>1</sup>, F. Shanahan<sup>1,3</sup>, T.G. Dinan<sup>1,2</sup> and J.F. Cryan<sup>1,4</sup>



GF mice spent more time in the open arms of the EPM



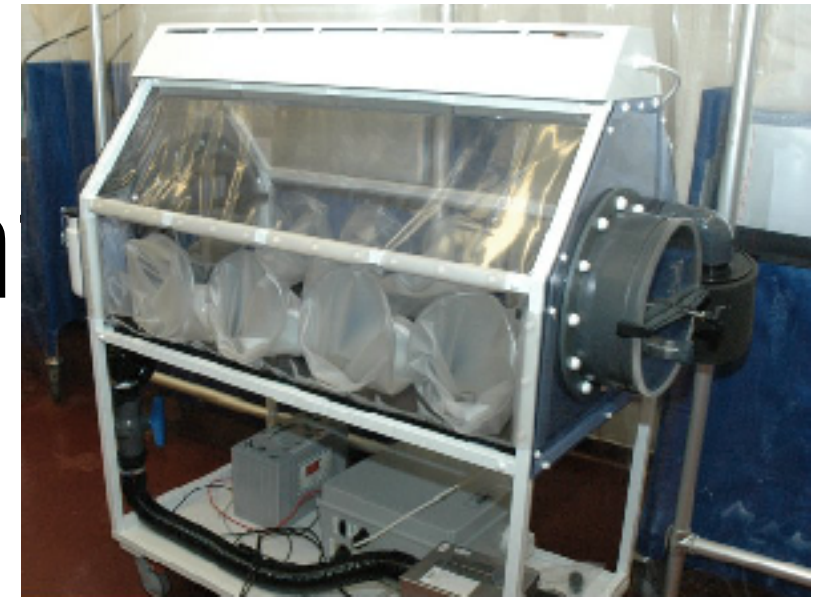
GF mice showed increased open arm entries



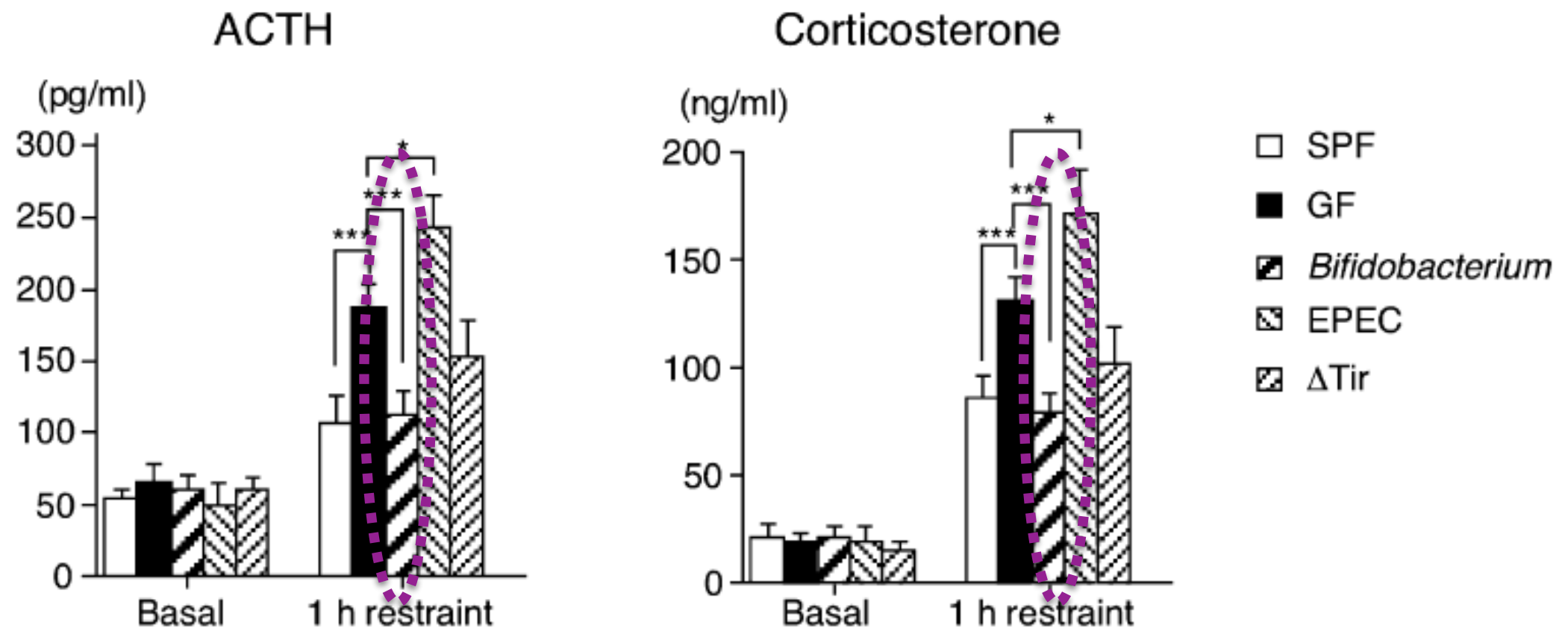
# Postnatal microbial colonization programs the hypothalamic–pituitary–adrenal system for stress response in mice

Nobuyuki Sudo<sup>1,2</sup>, Yoichi Chida<sup>1</sup>, Yuji Aiba<sup>3,4</sup>, Junko Sonoda<sup>1</sup>, Naomi Oyama<sup>1</sup>, Xiao-Nian Yu<sup>1</sup>, Chiharu Kubo<sup>1</sup> and Yasuhiro Koga<sup>3</sup>

<sup>1</sup>Department of Psychosomatic Medicine and <sup>2</sup>Department of Health Care Administration & Management, Graduate School of Medical Science, Kyushu University, Fukuoka, Japan, <sup>3</sup>Department of Infectious Diseases, Tokai University School of Medicine, Isehara, Kanagawa, Japan and <sup>4</sup>Wakamu Pharmaceutical Co. Ltd, Ohi-machi, Kanagawa, Japan

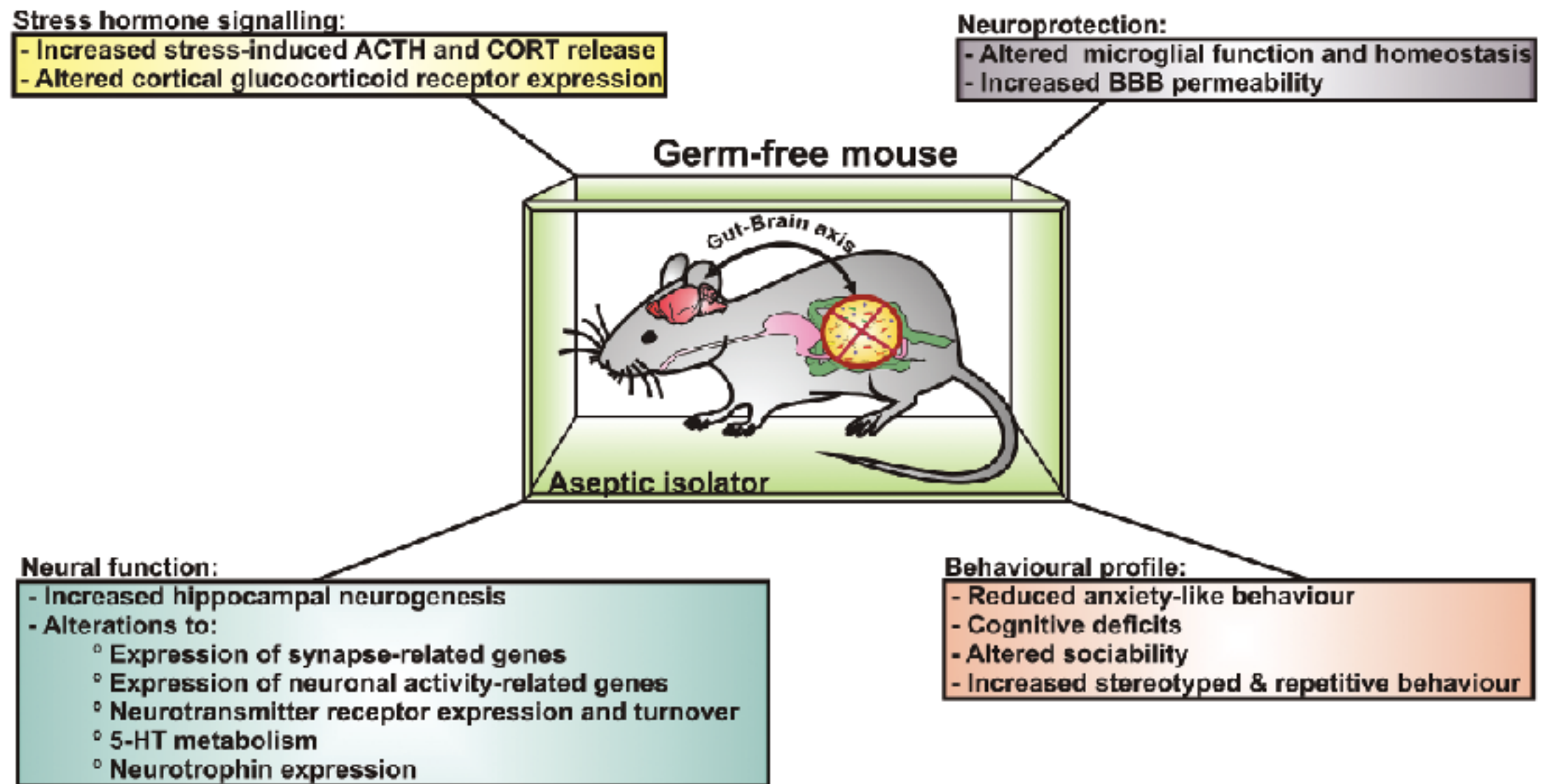


Colonization with *Bifidobacterium* normalized stress response



# Lessons learned from germ free mice

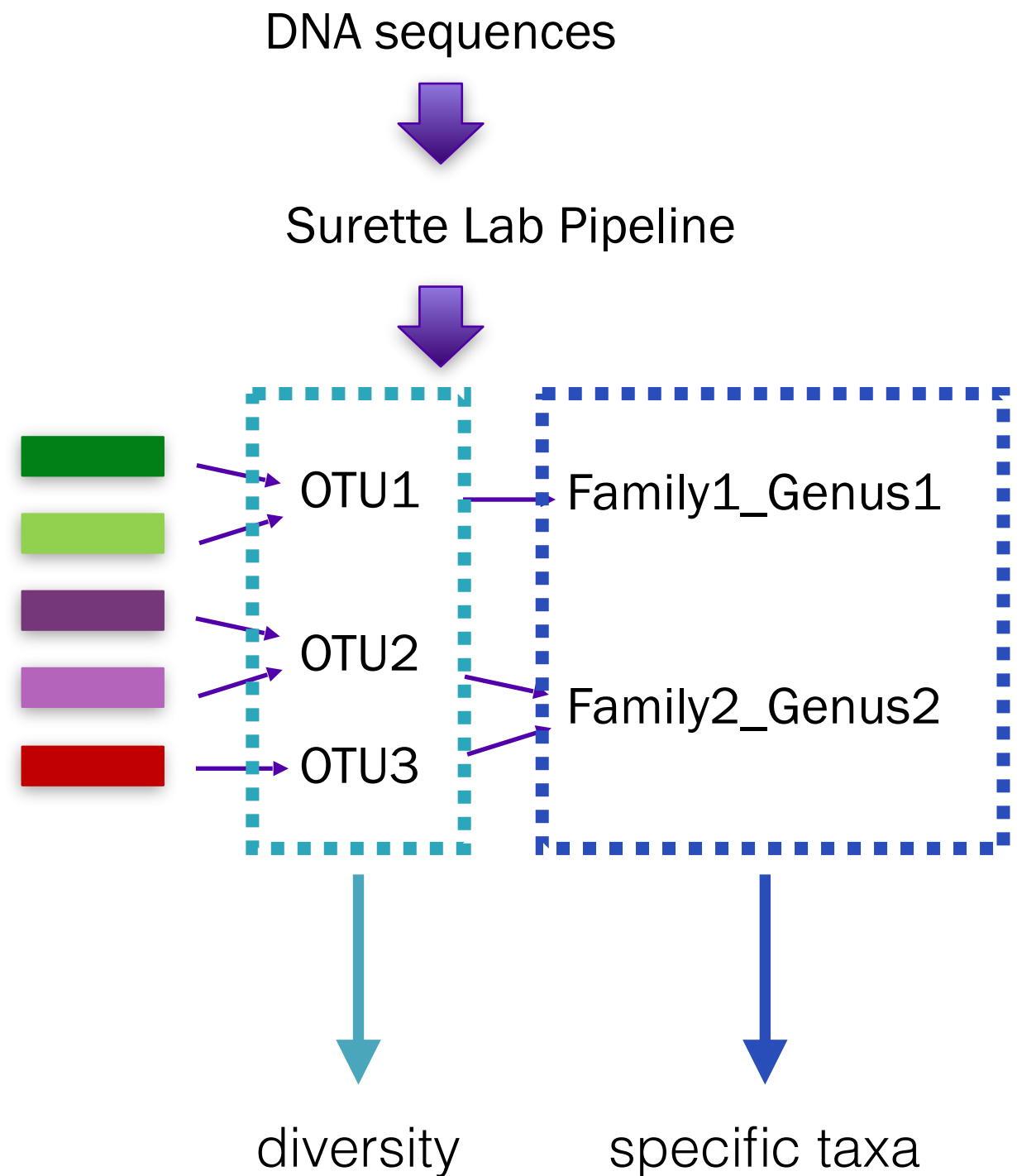
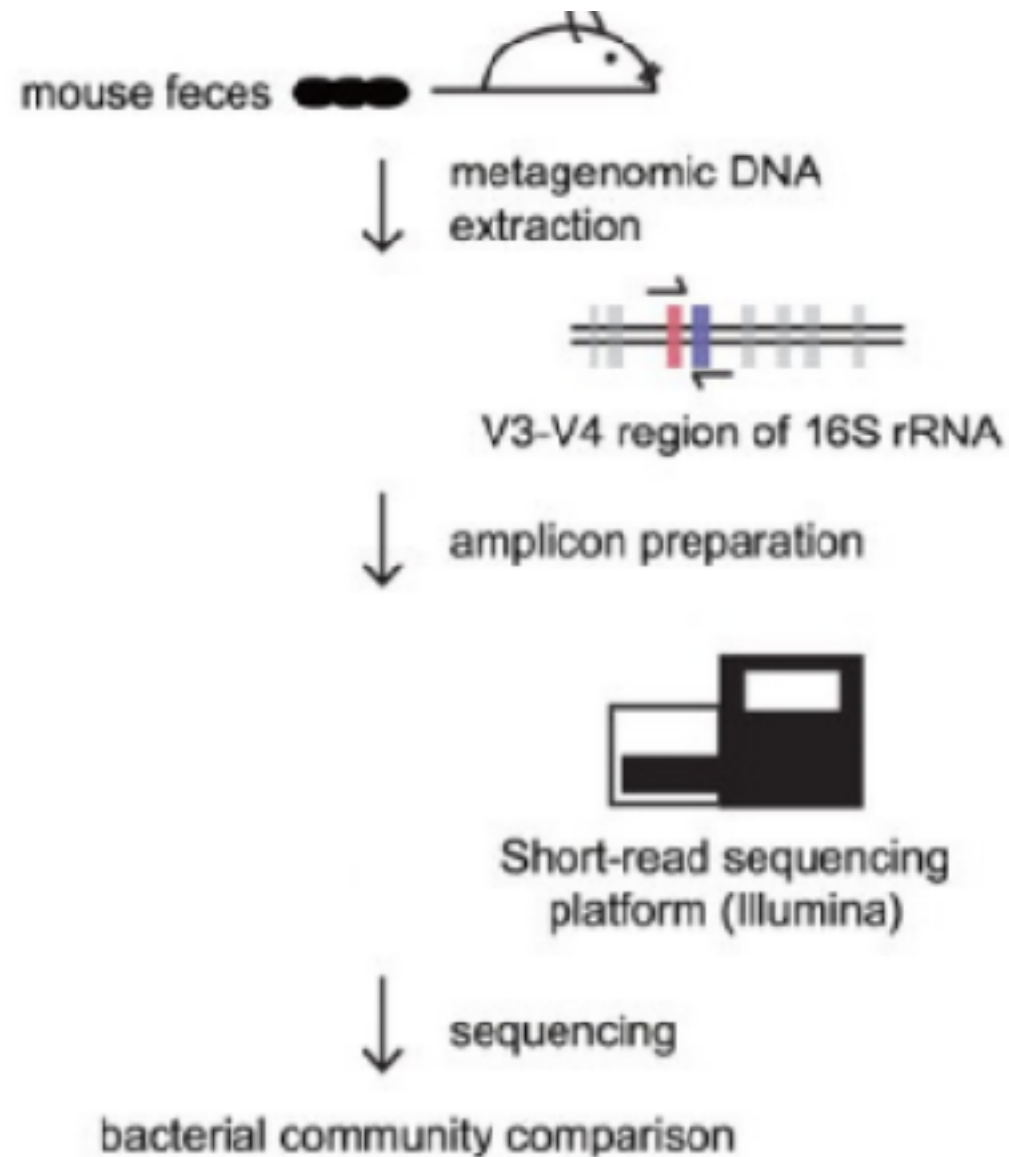
- microbiota influence behaviour, stress circuitry, stress responsively, and brain structure



IMPORTANCE OF ENTERIC NERVOUS SYSTEM  
West et al, 2016; McVey Neufeld et al, 2013

Luczynski et al 2016; Int J Neuropsychopharm

# 16s rRNA gene sequencing shows strain differences in bacterial composition





# What is the evidence that microbes influence mood in people?

- Several studies have shown that the gut-brain axis may play an important role in brain development, behaviour, and mood
- In particular, studies have examined the impact of probiotics on stress systems in healthy individuals
- Several studies have show benefit of probiotic consumption in healthy individuals
  - improved mood (Benton et al 2007)
  - influenced brain activity in emotional centers in healthy individuals (Tillisch et al 2013)
  - showed a beneficial effect on anxiety and depressive measures and reduced stress hormone levels (Messaoudi et al 2011)
  - showed reduction in cognitive reactivity to sad mood, specifically ruminative thoughts (Steenbergen et al 2015)

# Benefits of probiotics on mood

Table 1. Benefits of probiotics on mood

| Benefit   | Bacterial Taxa   | Population               | Dose (CFU)  | Ref |
|---|--|--------------------------|---|-----|
| Improved Mood   | <i>Lactobacillus casei</i>   | Healthy Individuals      | 6 X10 <sup>9</sup> /day                                     | [1] |
| Reduced anxiety and depressive measures                         | <i>Lactobacillus helveticus</i> R0052<br><i>Bifidobacterium longum</i> R0175   | Healthy Individuals      | 3 X10 <sup>9</sup> /day                                     | [2] |
| Reduced anxiety measures  | <i>Lactobacillus casei</i>   | Chronic fatigue syndrome | 2.4 X10 <sup>10</sup> /day                                  | [3] |
| Reduced stress hormone levels                                   | <i>Lactobacillus helveticus</i> R0052<br><i>Bifidobacterium longum</i> R0175   | Healthy Individuals      | 3 X10 <sup>9</sup> /day                                     | [2] |
| Reduced engagement of brain network to emotion recognition task | <i>Streptococcus thermophilis</i> (CNCM I-1630)<br><i>Lactobacillus bulgaricus</i> (CNCM I-1632 and I-1519)<br><i>Lactococcus lactis</i> supsp <i>lactis</i> (CNCM I-1631)<br><i>Bifidobacterium animalis</i> subsp <i>lactis</i> (CNCM I-2494)                      | Healthy Individuals      | 1.2 X10 <sup>9</sup> /day<br><br>1.25 X10 <sup>9</sup> /day | [4] |
| Reduction in cognitive reactivity to sad mood                   | <i>Bifidobacterium bifidum</i> W23<br><i>Bifidobacterium lactis</i> W52<br><i>Lactobacillus acidophilus</i> W37<br><i>Lactobacillus brevis</i> W63<br><i>Lactobacillus casei</i> W56<br><i>Lactobacillus salivarius</i> W24<br><i>Lactococcus lactis</i> W19 and W58 | Healthy Individuals      | 5 X10 <sup>9</sup> /day                                     | [5] |
| Reduced depression scores                                       | <i>Lactobacillus casei</i><br><i>Lactobacillus acidophilus</i><br><i>Bifidobacterium bifidum</i>   | Major depression         | 2 X X10 <sup>9</sup> /day                                   | [6] |
| Reduced serum insulin   | <i>Lactobacillus casei</i><br><i>Lactobacillus acidophilus</i><br><i>Bifidobacterium bifidum</i>   | Major depression         | 2 X X10 <sup>9</sup> /day                                   | [6] |
| Reduced inflammation  | <i>Lactobacillus casei</i><br><i>Lactobacillus acidophilus</i><br><i>Bifidobacterium bifidum</i>   | Major depression         | 2 X X10 <sup>9</sup> /day                                   | [6] |

CFU – colony forming units; R – Probio'Stick: batch no. 6533308; Institut Rosell-Lallemand, Blagnac, France; CNCM – French National Collection of Cultures of Microorganisms, Paris, France; W – Ecologic Barrier, Winclove probiotics, The Netherlands

# Benefits of probiotics on mood

1. Benton, D., C. Williams, and A. Brown, *Impact of consuming a milk drink containing a probiotic on mood and cognition*. Eur J Clin Nutr, 2007. **61**(3): p. 355-61.
2. Messaoudi, M., et al., *Beneficial psychological effects of a probiotic formulation (Lactobacillus helveticus R0052 and Bifidobacterium longum R0175) in healthy human volunteers*. Gut Microbes, 2011. **2**(4): p. 256-61.
3. Rao, A.V., et al., *A randomized, double-blind, placebo-controlled pilot study of a probiotic in emotional symptoms of chronic fatigue syndrome*. Gut Pathog, 2009. **1**(1): p. 6.
4. Tillisch, K., et al., *Consumption of fermented milk product with probiotic modulates brain activity*. Gastroenterology, 2013. **144**(7): p. 1394-401, 1401 e1-4.
5. Steenbergen, L., et al., *A randomized controlled trial to test the effect of multispecies probiotics on cognitive reactivity to sad mood*. Brain Behav Immun, 2015.
6. Akkasheh, G., et al., *Clinical and metabolic response to probiotic administration in patients with major depressive disorder: A randomized, double-blind, placebo-controlled trial*. Nutrition, 2016. **32**(3): p. 315-20.



# How to pick a probiotic?



## ***Probiotics: A Consumer Guide for Making Smart Choices***

*Developed by the International Scientific Association for Probiotics and Prebiotics ([www.ISAPPScience.org](http://www.ISAPPScience.org))*

**Probiotics** are live microorganisms that, when administered in adequate amounts, confer a health benefit on the host. They are present in numerous products, most commonly in foods and supplements.



## Clinical Guide to Probiotic Products Available in Canada

Indications, Dosage Forms and Clinical Evidence to Date - 2018 Edition

A microscopic view of various bacteria, including rod-shaped and spherical forms, in shades of blue.

**PREBIOTICS**

A microscopic view of various bacteria, including rod-shaped and spherical forms, in shades of blue.

**FERMENTED FOODS**

# Gut bacteria and depression

# Correlation between the human fecal microbiota and depression

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*Neurogastroenterol Motil* (2014) 26, 1155–1162

Neurogastroenterol Motil (2014) 26, 1155–1162

## A.11.11.11 Altered fecal microbiota composition in patients with major depressive disorder



Haiyin Jiang<sup>a,1</sup>, Zongxin Ling<sup>a,1</sup>, Yonghua Zhang<sup>b,1</sup>, Hongjin Mao<sup>c</sup>, Zhanping Ma<sup>d</sup>, Yan Yin<sup>c</sup>, Weihong Wang<sup>e</sup>, Wenxin Tang<sup>c</sup>, Zhonglin Tan<sup>c</sup>, Jianfei Shi<sup>c</sup>, Lanjuan Li<sup>a,2</sup>, Bing Ruan<sup>a,\*</sup>

# Possible association of *Bifidobacterium* and *Lactobacillus* in the gut microbiota of patients with major depressive disorder

## Transferring the blues: Depression-associated gut microbiota induces neurobehavioural changes in the rat

Journal of Psychiatric Research 82 (2016) 109–118

John R. Kelly<sup>a, b</sup>, Yuliya Borre<sup>a</sup>, Ciaran O' Brien<sup>a, c</sup>, Elaine Patterson<sup>a, c</sup>, Sahar El Aidy<sup>a, d</sup>,  
Jennifer Deane<sup>c</sup>, Paul J. Kennedy<sup>a, c</sup>, Alan E. Hoban<sup>a</sup>, Lucinda Scott<sup>l</sup>, Gerard Clarke<sup>a, b</sup>, John F. Cryan<sup>a, b</sup>

Prevotella and Klebsiella proportions in fecal microbial communities are potential characteristic parameters for patients with major depressive disorder

Journal of Affective Disorders 207 (2017) 300–304

Ping Lin<sup>a,1</sup>, Bingyu Ding<sup>b,c,1</sup>, Chunyan Feng<sup>d</sup>, Shuwei Yin<sup>b</sup>, Ting Zhang<sup>b</sup>, Xin Qi<sup>b</sup>, Huiying Lv<sup>b</sup>,  
Xiaokui Guo<sup>c</sup>, Ke Dong<sup>c</sup>, Yongzhang Zhu<sup>c,1</sup>, Qingtian Li<sup>b,\*,1</sup>

# Gut bacteria and depression

Table 1. Bacterial taxa differences observed in individuals with major depressive disorder

| Experimental Design       |                           |                              |                                       |                       |
|---------------------------|---------------------------|------------------------------|---------------------------------------|-----------------------|
| Reference                 | MDD Sample (n)            | Comparison Group (n)         | OTU Picking                           | Taxon Assignment      |
| Naseribafrouei et al 2014 | mild to severe MDD (37)   | neurological outpatient (18) | Closed Reference, UClust modified     | RDP database          |
| Jiang et al 2015          | mild to moderate MDD (29) | healthy volunteers (30)      | Mothur ver1.25.0, custom Perl scripts | RDP database          |
| Kelly et al 2016          | MDD (34)                  | healthy volunteers (33)      | USEARCH v7                            | BLAST, Silva v.111    |
| Zheng et al 2016          | MDD (58)                  | healthy volunteers (63)      | Roche software                        | RDP database          |
| Lin et al 2017            | MDD (10)                  | healthy volunteers (10)      | Mothur v.1.30                         | Silva v.119 in mothur |



| Differences in Relative Abundance  |                        |                      |                               |  |
|--|------------------------|----------------------|-------------------------------|--|
| Phyla  | Order                  | Class                | Family                        | Genus  |
| <i>Naseribfrouei et al 2014 - method</i>   |                        |                      |                               |  |
| Bacteroidetes (up)   |                        | Bacteroidales (down) | Lacnospiraceae (down)         | <i>Alistipes</i> (up)<br><i>Oscillibacter</i> (up) |
| <i>Jiang et al 2015 - Mothur metastats</i>   |                        |                      |                               |  |
| Bacteroidetes (up)   |                        |                      | Acidaminococcaceae (up)       | <i>Alistipes</i> (up)                              |
| Proteobacteria (up)  |                        |                      | Enterobacteriaceae (up)       | <i>Blautia</i> (up)                                |
| Firmicutes (down)  |                        |                      | Fusobacteriaceae (up)         | <i>Clostridium XIX</i> (up)                        |
|  |                        |                      | Porphyromonadaceae (up)       | <i>Lachnospiraceae</i> (up)                        |
|  |                        |                      | Rikenellaceae (up)            | <i>Megamonas</i> (up)                              |
|  |                        |                      | Bacteroidaceae (down)         | <i>Parabacteroides</i> (up)                        |
|  |                        |                      | Erysipelotrichaceae (down)    | <i>Parasutterella</i> (up)                         |
|  |                        |                      | Lacnospiraceae (down)         | <i>Phascolarctobacterium</i> (up)                  |
|  |                        |                      | Prevotellaceae (down)         | <i>Oscillibacter</i> (up)                          |
|  |                        |                      | Ruminococcaceae (down)        | <i>Roseburia</i> (up)                              |
|  |                        |                      | Veillonellaceae (down)        | <i>Bacteroides</i> (down)                          |
|  |                        |                      |                               | <i>Dialister</i> (down)                            |
|  |                        |                      |                               | <i>Faecalibacterium</i> (down)                     |
|  |                        |                      |                               | <i>Prevotella</i> (down)                           |
|  |                        |                      |                               | <i>Ruminococcus</i> (down)                         |
| <i>Jiang et al 2015 - LefSe LDA; alpha level = 0.05, effect size threshold = 2</i> |                        |                      |                               |  |
|  | Enterobacteriales (up) |                      | Polphyromonadaceae (up)       | <i>Alistipes</i> (up)                              |
|  |                        |                      | Enterobacteriaceae (up)       | <i>Parabacteroides</i> (up)                        |
|  |                        |                      | Rikenellaceae (up)            | <i>Butyrivibrio</i> (up)                           |
|  |                        |                      | Erysipelotrichaceae (up)      | <i>Flavonifractor</i> (up)                         |
|  |                        |                      | Peptostreptococcaceae (down)  | <i>Haemophilus</i> (down)                          |
|  |                        |                      | Pasteurellaceae (down)        | <i>Dialister</i> (down)                            |
|  |                        |                      | Ruminococcaceae (down)        | <i>Faecalibacterium</i> (down)                     |
|  |                        |                      |                               | <i>Escherichia shigella</i> (down)                 |
|  |                        |                      |                               | <i>Ruminococcus</i> (down)                         |
| <i>Kelly et al 2016 - Mann-Whitney U test, FDR adjusted 10%</i>                    |                        |                      |                               |  |
|  |                        |                      | Prevotellaceae (down)         | <i>Prevotella</i> (down)                           |
|  |                        |                      | Thermoanaerobacteriaceae (up) | <i>Dialister</i> (down)                            |
|  |                        |                      |                               | <i>Eggerthella</i> (up)                            |
|  |                        |                      |                               | <i>Holdemania</i> (up)                             |
|  |                        |                      |                               | <i>Gelria</i> (up)                                 |
|  |                        |                      |                               | <i>Turicibacter</i> (up)                           |
|  |                        |                      |                               | <i>Paraprevotella</i> (up)                         |
|  |                        |                      |                               | <i>Anaerofilum</i> (up)                            |

|  |  |  |
|--|--|--|
| <i>Lin et al 2017 - Student's t-test (Phyla) and Wilcoxon's Sign Rank Test (Genus)</i> |  |  |
| Bacteroidetes (down)   |  | <i>Prevotella</i>                            |
| Firmicutes (up)  |  | <i>Klebsiella</i>                            |
|  |  | <i>Streptococcus</i>                         |
|  |  | <i>Clostridium XIX</i>                       |
| <i>Zheng et al 2016 - Random Forest Classifier</i>                                     |  |  |
|  | Actinomycineae (up)                      | <i>Parvimonas</i> (up)                       |
|  | Coriobacterineae (up)                    | <i>Anerostipes</i> (up)                      |
|  | Lactobacillaceae (up)                    | <i>Blautia</i> (up)                          |
|  | Streptococcaceae (up)                    | <i>Dorea</i> (up)                            |
|  | Clostridiales incertae sedis XI (up)     | <i>Lachnospiraceae incertae sedis</i> (up)   |
|  | Eubacteriaceae (up)                      | <i>Clostridium IV</i> (up)                   |
|  | Lachnospiraceae (up)                     | <i>Alistipes</i> (down)                      |
|  | Ruminococcaceae (up)                     | <i>Coproccus</i> (down)                      |
|  | Erysipelotrichaceae in certae sedis (up) | <i>Clostridium XIVa</i> (down)               |
|  | Bacteroidaceae (down)                    | <i>Phascolarctobacterium</i> (down)          |
|  | Rikenellaceae (down)                     | <i>Megamonas</i> (down)                      |
|  | Lachnospiraceae (down)                   | <i>Lachnospiraceae incertae sedis</i> (down) |
|  | Acidaminococcaceae (down)                | <i>Roseburia</i> (down)                      |
|  | Vellonellaceae (down)                    | <i>Faecalibacterium</i> (down)               |
|  | Sutterellaceae (down)                    |  |

# Gut bacteria and depression

Nutrition 32 (2016) 315–320

Applied nutritional investigation

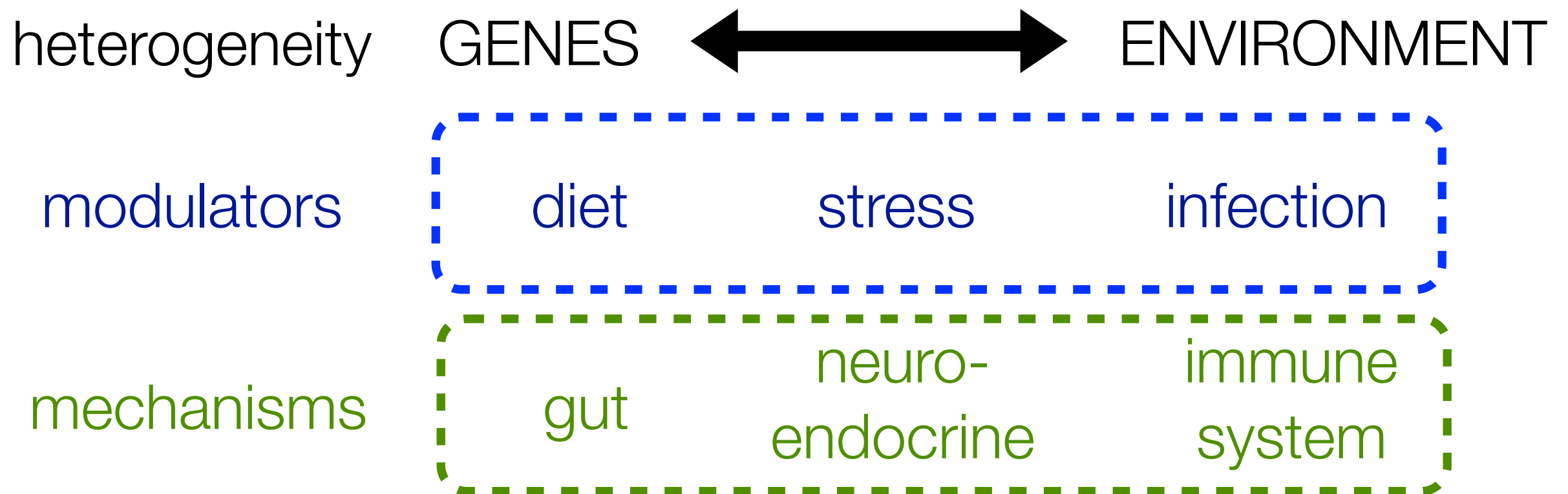
Clinical and metabolic response to probiotic administration in patients with major depressive disorder: A randomized, double-blind, placebo-controlled trial

Ghodarz Akkasheh M.D.<sup>a</sup>, Zahra Kashani-Poor M.D.<sup>a</sup>,  
Maryam Tajabadi-Ebrahimi Ph.D.<sup>b</sup>, Parvaneh Jafari Ph.D.<sup>c</sup>, Hossein Akbari Ph.D.<sup>d</sup>,  
Mohsen Taghizadeh Ph.D.<sup>e</sup>, Mohammad Reza Memarzadeh Ph.D.<sup>f</sup>,  
Zatollah Asemi Ph.D.<sup>e,\*</sup>, Ahmad Esmailzadeh Ph.D.<sup>g,h,i</sup>

- Probiotic supplementation for 8 week compared to placebo - *Lactobacillus casei*, *L. acidophilus*, *Bifidobacterium bifidum*
- Probiotic supplement was associated with
  - reduced depression scores (Beck Depression Score)
  - reduced serum insulin
  - reduce inflammatory marker - C-reactive protein



# Can peripheral measures such as microbiota explain heterogeneity in health and disease?



# Many factors are important to gut health and the composition of gut bacteria?

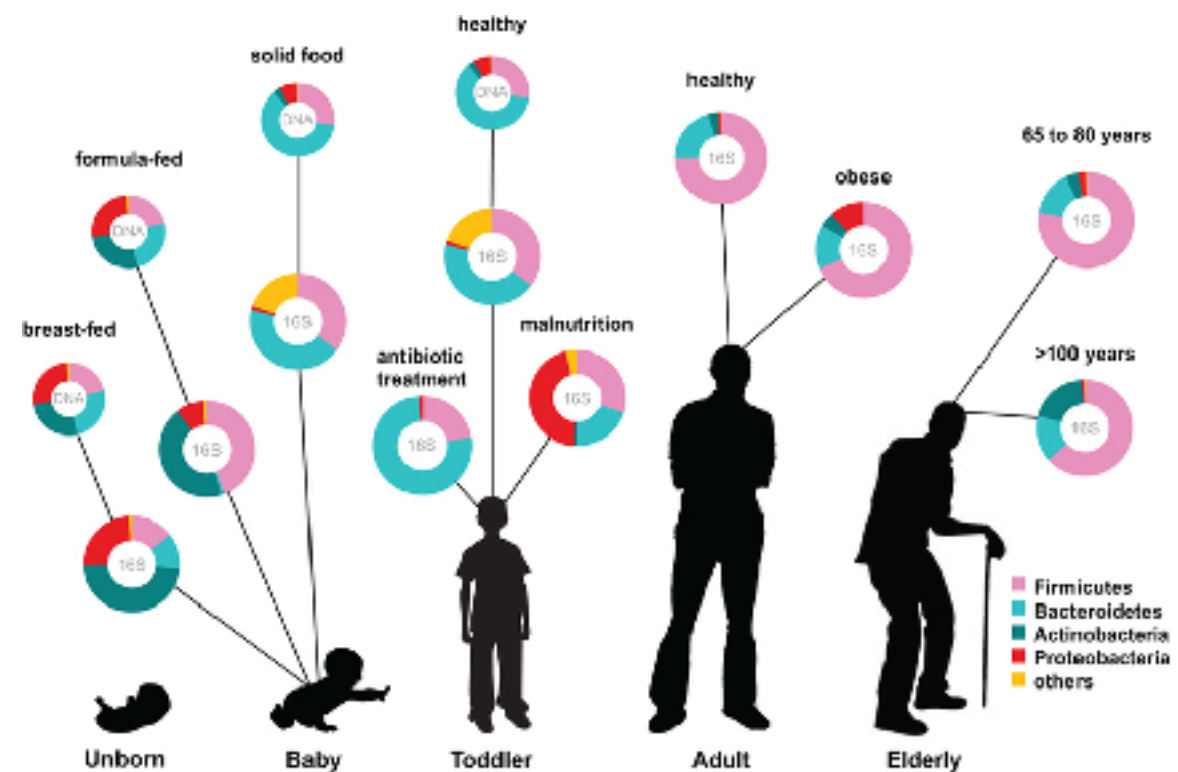
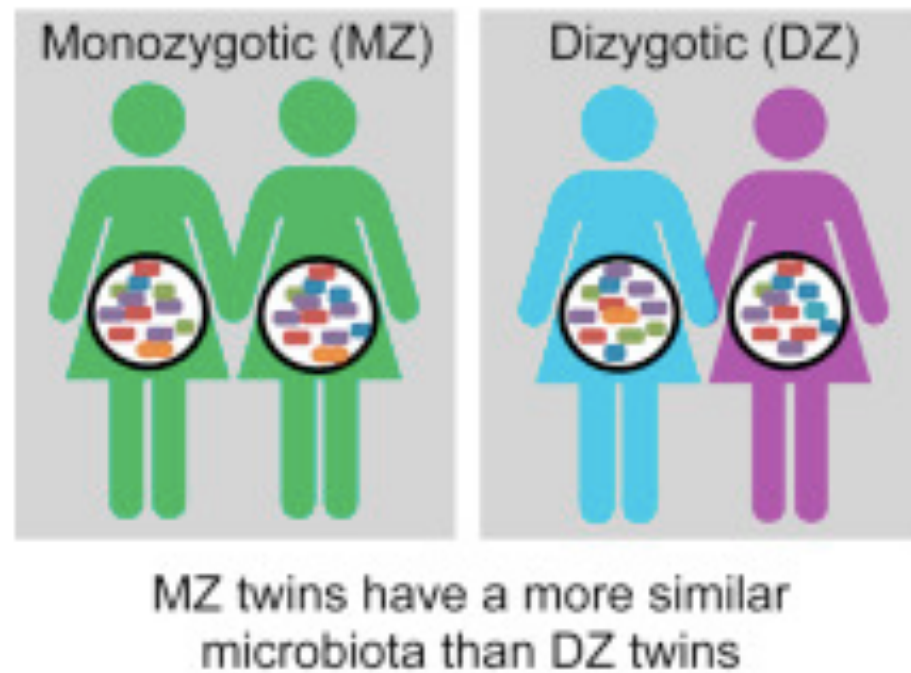
Genetics

Diet

Exercise

Drugs

Stress



Goodrich et al 2015 Cell 159, 789-799

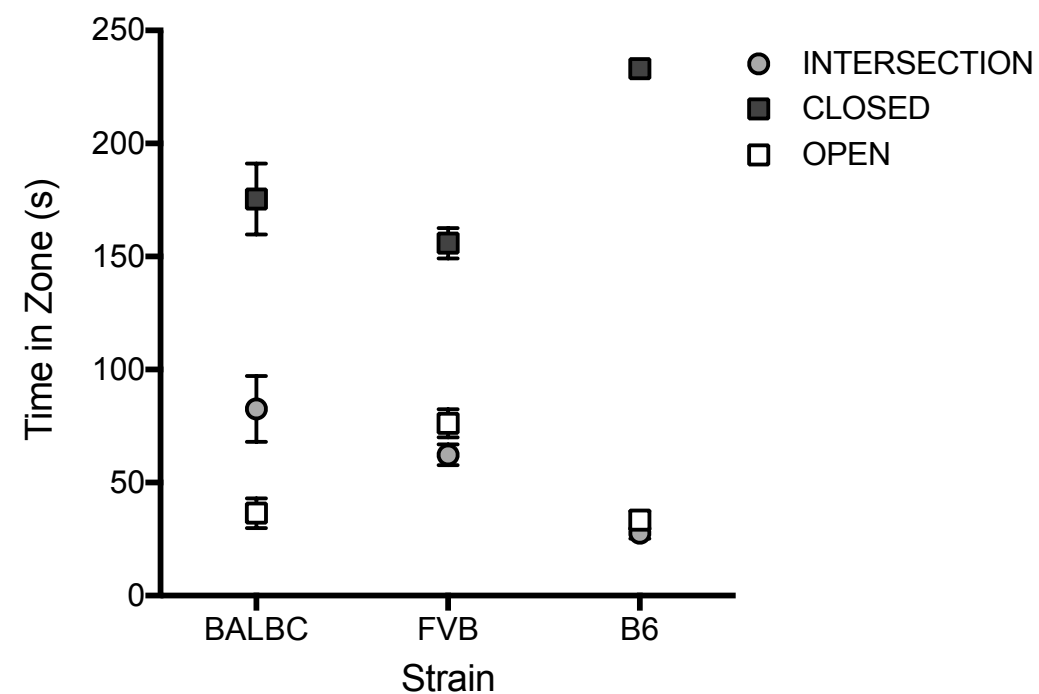
Age

Sex

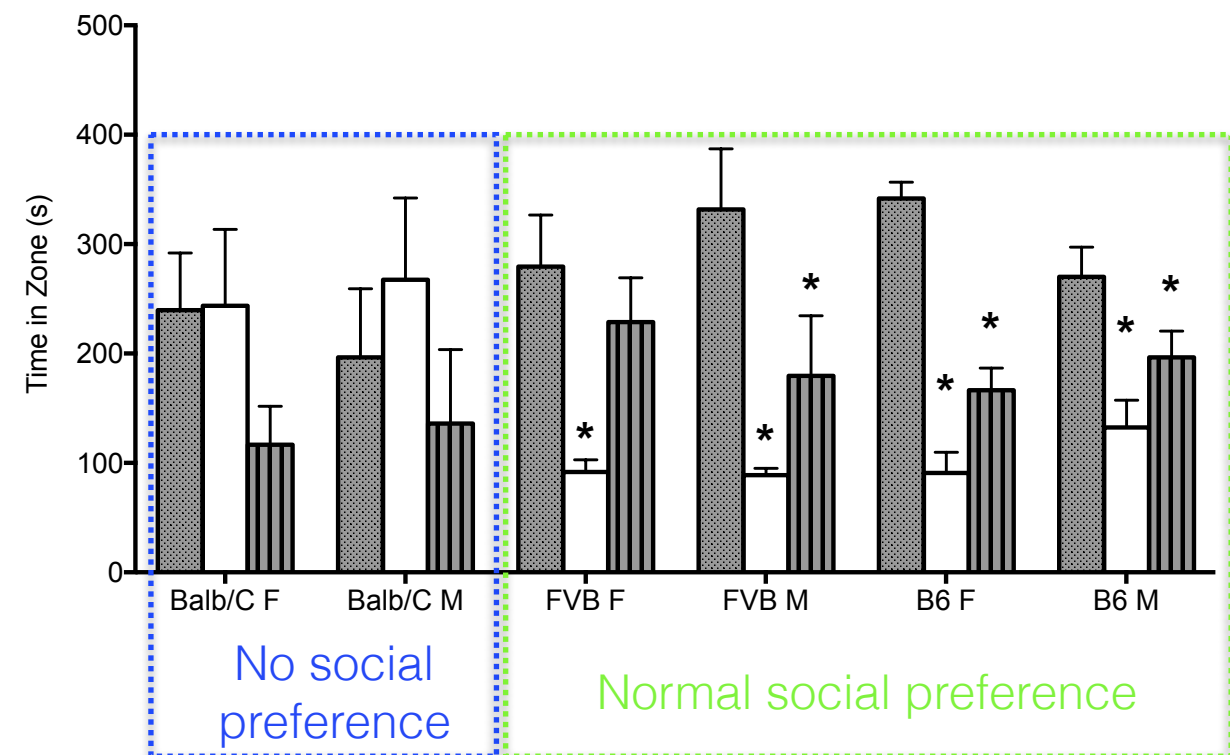
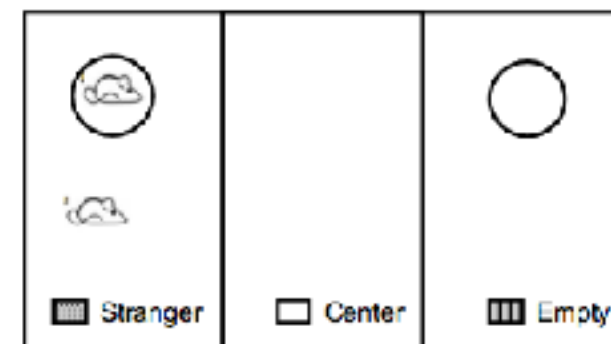
# Host Genetics influences Behaviour



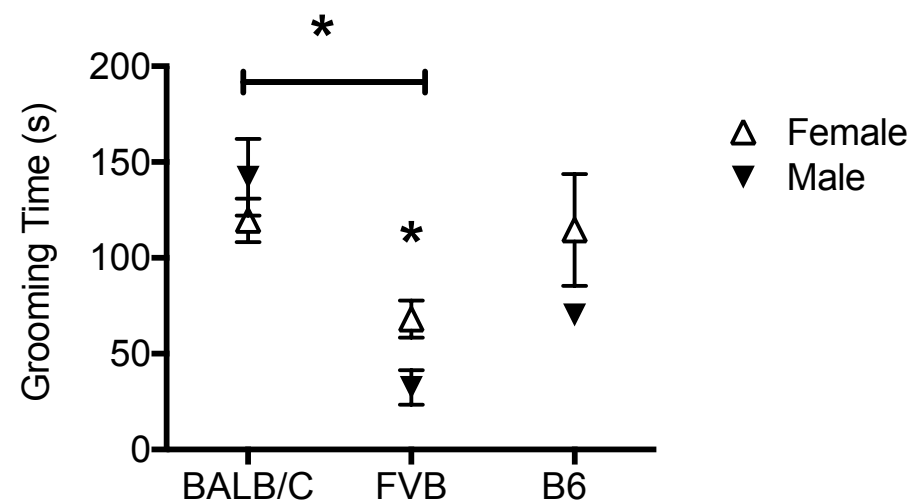
## Elevated Plus Maze behaviour



## Social behaviour

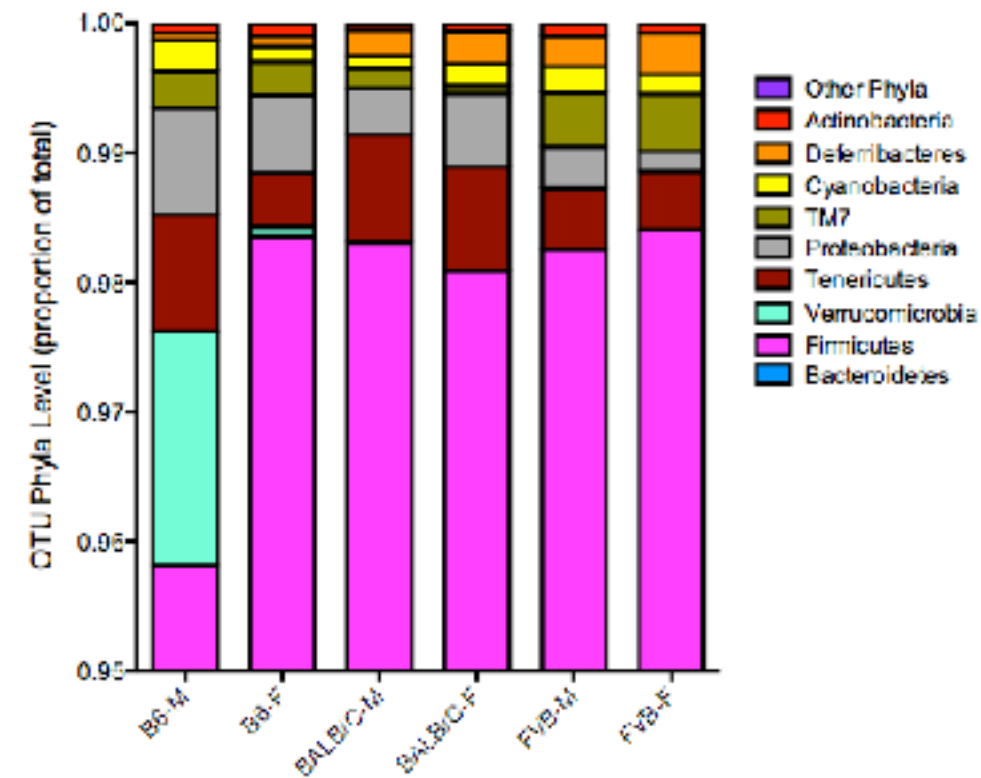
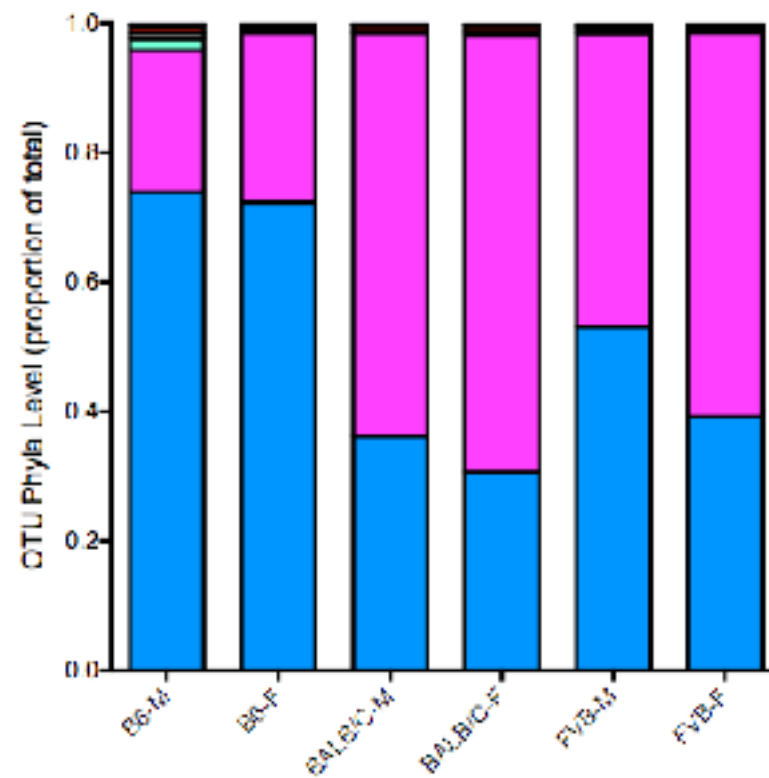
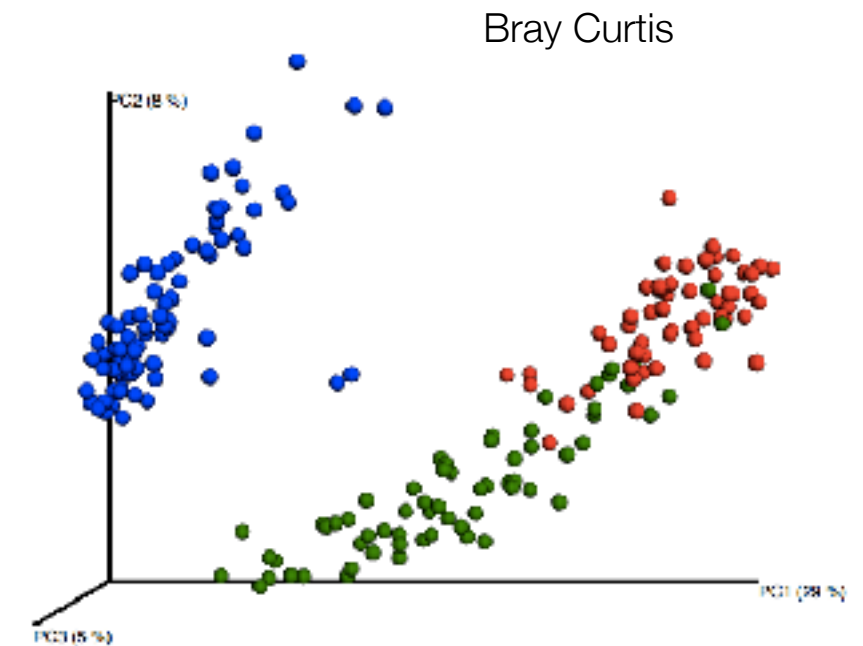
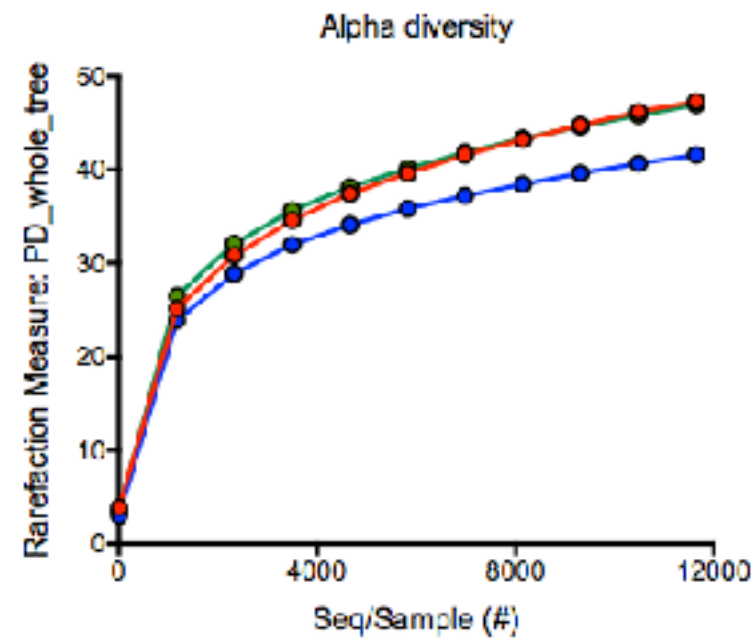


## Self-grooming behaviour



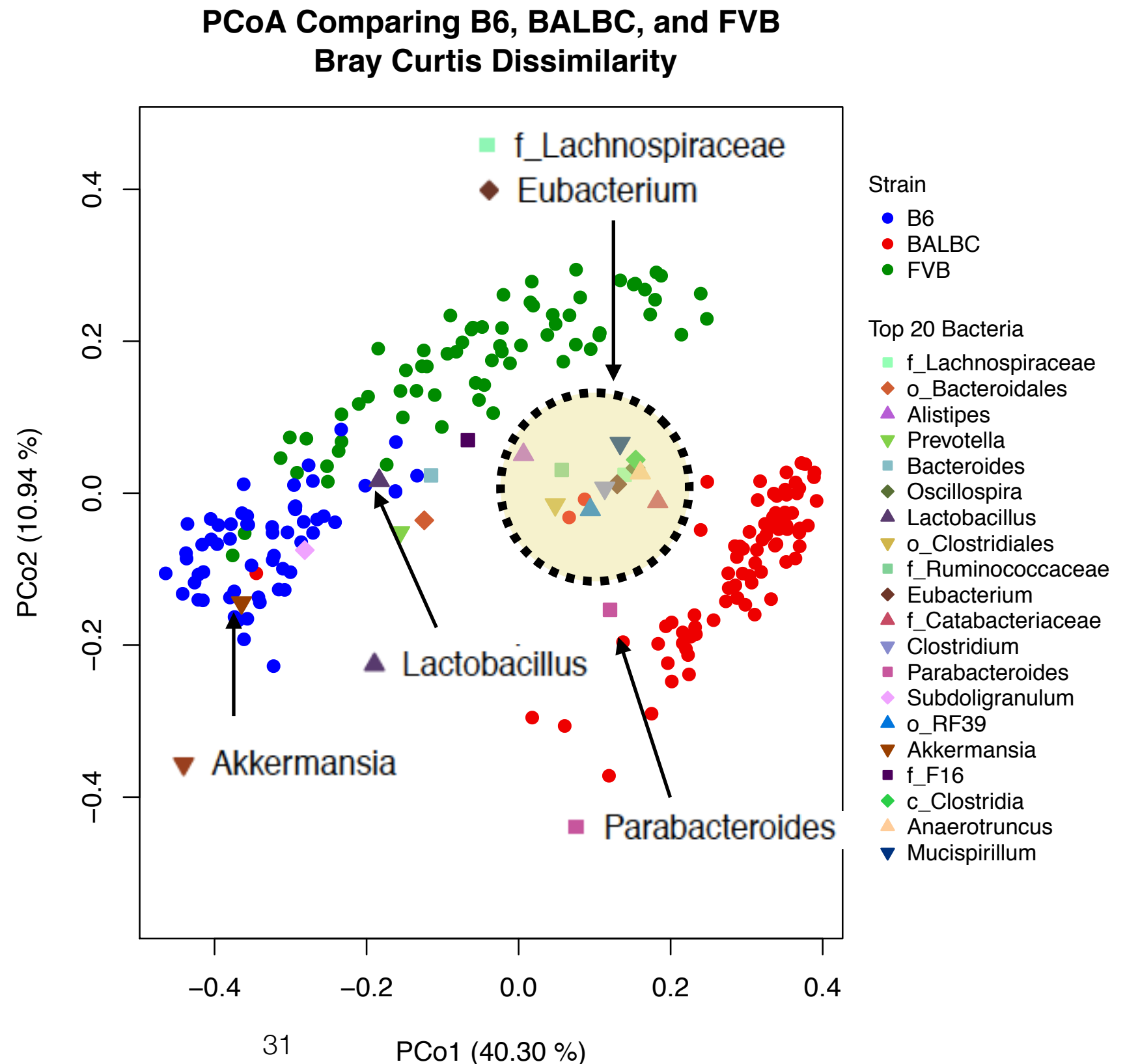


# 16s rRNA analysis of bacterial composition



# Are specific taxa associated with different strains of mice?

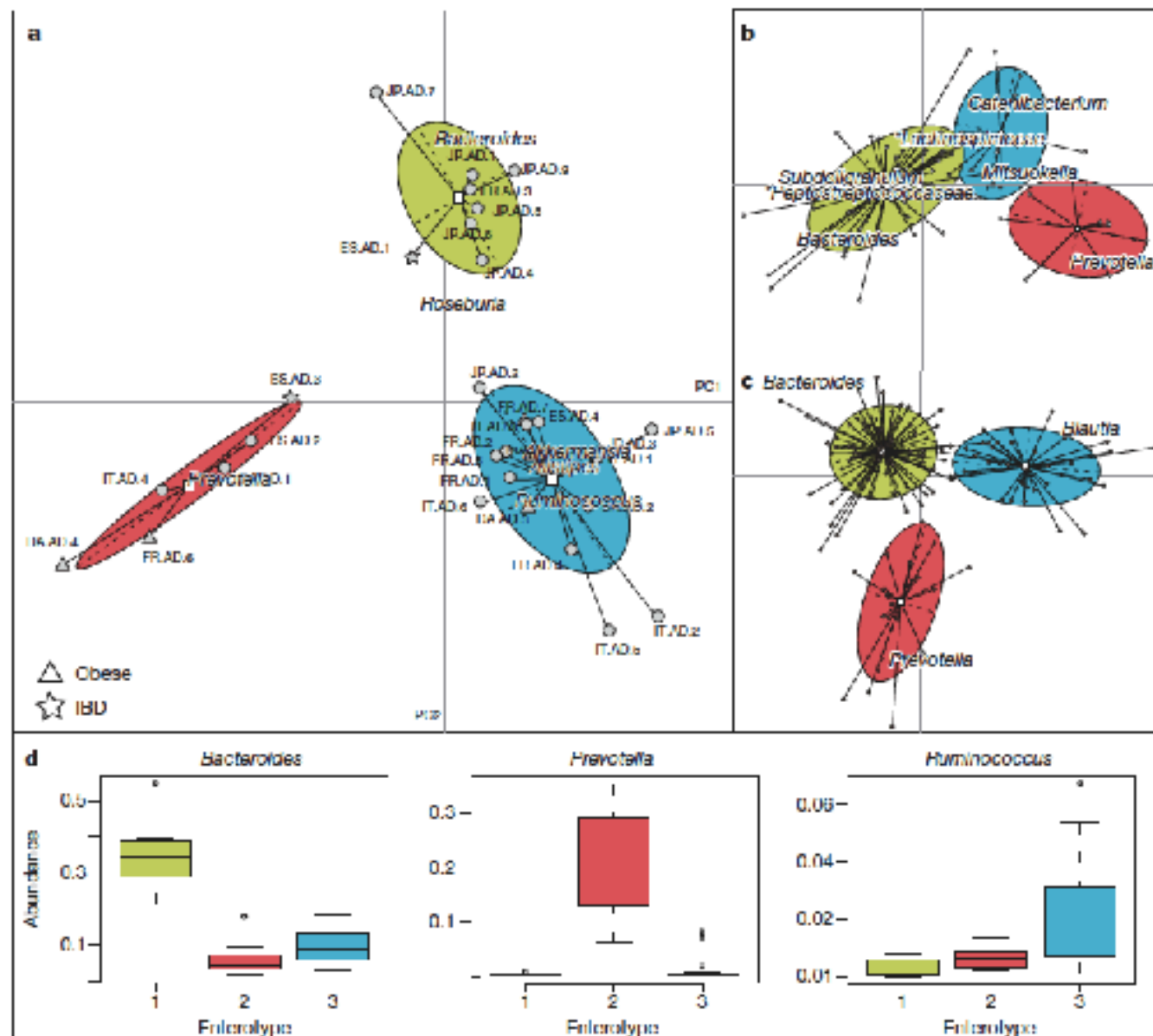
PCo Analysis  
with  
count-weighted ave  
for top 20 taxa





# Enterotypes of the human gut microbiome

Manimozhiyan Arumugam<sup>1,4</sup>, Jeroen Raes<sup>1,2,4</sup>, Eric Pelletier<sup>3,4,5</sup>, Denis Le Paslier<sup>3,4,5</sup>, Takuji Yamada<sup>1</sup>, Daniel R. Mende<sup>1</sup>, Gabriel R. Fernandes<sup>1,6</sup>, Julien Tap<sup>1,7</sup>, Thomas Bruls<sup>3,4,5</sup>, Jean Michel Batto<sup>7</sup>, Marcelo Bertalan<sup>8</sup>, Natalia Borruecl<sup>9</sup>, Francesc Casellas<sup>9</sup>, Leyden Fernandez<sup>10</sup>, Laurent Cautier<sup>8</sup>, Torben Hansen<sup>11,12</sup>, Masahira Hattori<sup>13</sup>, Tetsuya Hayashi<sup>14</sup>, Michiel Kleerebezem<sup>15</sup>, Ken Kurokawa<sup>16</sup>, Marion Leclerc<sup>7</sup>, Florence Levenez<sup>7</sup>, Chaysavanh Manichanh<sup>9</sup>, H. Bjørn Nielsen<sup>8</sup>, Trine Nielsen<sup>11</sup>, Nicolas Pons<sup>7</sup>, Julie Poulain<sup>3</sup>, Junjie Qin<sup>17</sup>, Thomas Sicheritz-Ponten<sup>8,18</sup>, Sebastian Tims<sup>13</sup>, David Torrents<sup>10,19</sup>, Edgardo Ugarte<sup>3</sup>, Erwin G. Zoetendaal<sup>15</sup>, Jun Wang<sup>17,20</sup>, Francisco Guarner<sup>9</sup>, Oluf Pedersen<sup>11,21,22,23</sup>, Willem M. de Vos<sup>15,24</sup>, Søren Brunak<sup>8</sup>, Joel Doré<sup>7</sup>, MetaHIT Consortium†, Jean Weissenbach<sup>3,4,5</sup>, S. Dusko Ehrlich<sup>7</sup> & Peer Bork<sup>1,25</sup>



## Key points to consider

- Inter-individual differences in healthy human gut microbiota can be reduced by clustering individuals into subgroups, referred to as enterotypes, based on enrichment of specific taxa at the genus level
- Several tools available to examine the microbiome (composition, function, and active gene expression)



# Human Microbiome Project



- Human microbiome project - Phase I HMP, Phase II iHMP
- Phase I HMP examined diversity and composition of the human microbiome in healthy individuals
- Phase 2 iHMP examines the role of the microbiome in human health and disease - longitudinal (3 y) studies on pregnancy, gut disease onset (IBD), and respiratory viral infection and onset of type 2 diabetes
- information and publications to data at <https://www.hmpdacc.org/hmp/>

# Do microbes influence brain function and behaviour in people?

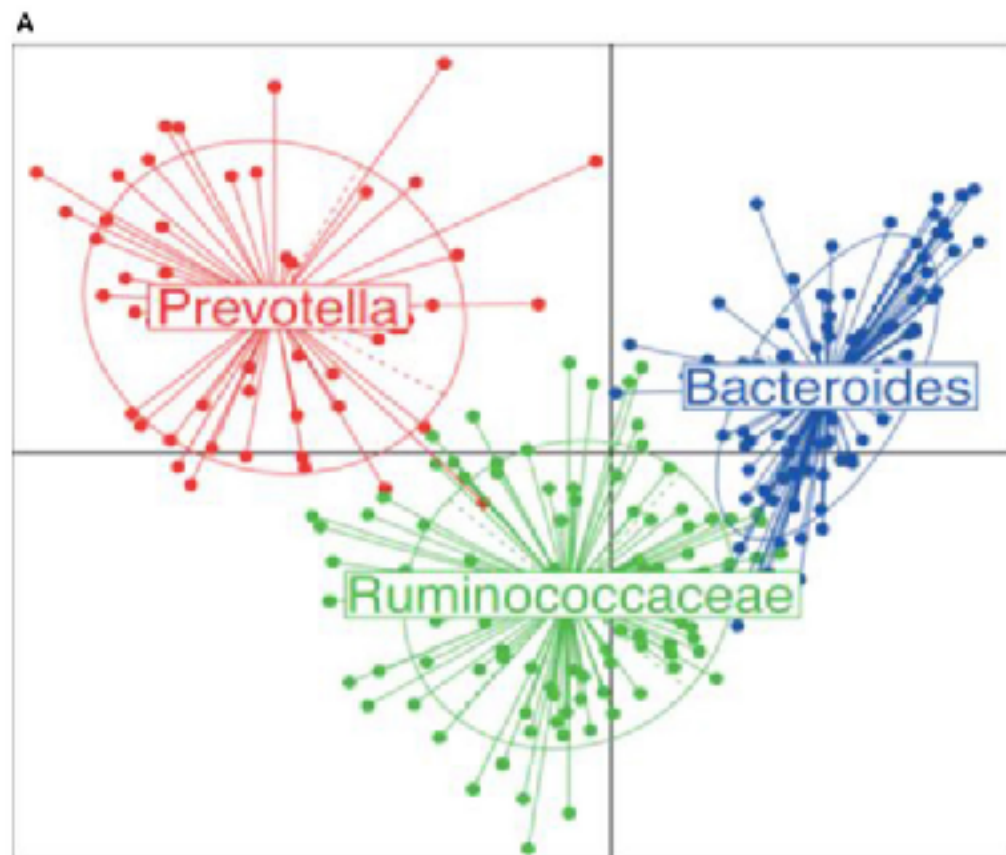
## Brain Structure and Response to Emotional Stimuli as Related to Gut Microbial Profiles in Healthy Women

Kirsten Tillisch, MD, Emeran A. Mayer, MD, PhD, Arpana Gupta, PhD, Zafar Gill, BSc, Rémi Brazeilles, MSc, Boris Le Nevé, PhD, Johan E.T. van Hylckama Vlieg, PhD, Denis Guyonnet, PhD, Muriel Derrien, PhD, and Jennifer S. Labus, PhD

- In healthy women, identified two enterotypes/clusters, a *Bacteroides* cluster and a *Prevotella* cluster
- Using functional MRI, structural MRI and diffusion tensor imaging, the investigators identified association between these 2 groups and emotional response, white matter connectivity, and brain volume

## Enterotype May Drive the Dietary-Associated Cardiometabolic Risk Factors

Ana C. F. de Moraes<sup>1</sup>, Gabriel R. Fernandes<sup>2</sup>, Isis T. da Silva<sup>1</sup>, Bianca Almeida-Pititto<sup>3</sup>, Everton P. Gomes<sup>4</sup>, Alexandre da Costa Pereira<sup>4</sup> and Sandra R. G. Ferreira<sup>1\*</sup>



- Identified 3 enter-types in Brazilian cohort
- Significantly more strict vegetarians in the Prevotella cluster and associated lower LDL-cholesterol in this group
- Within each cluster specific bacterial taxa were associated with cardiometabolic measures - some in a positive manner some as a risk factor



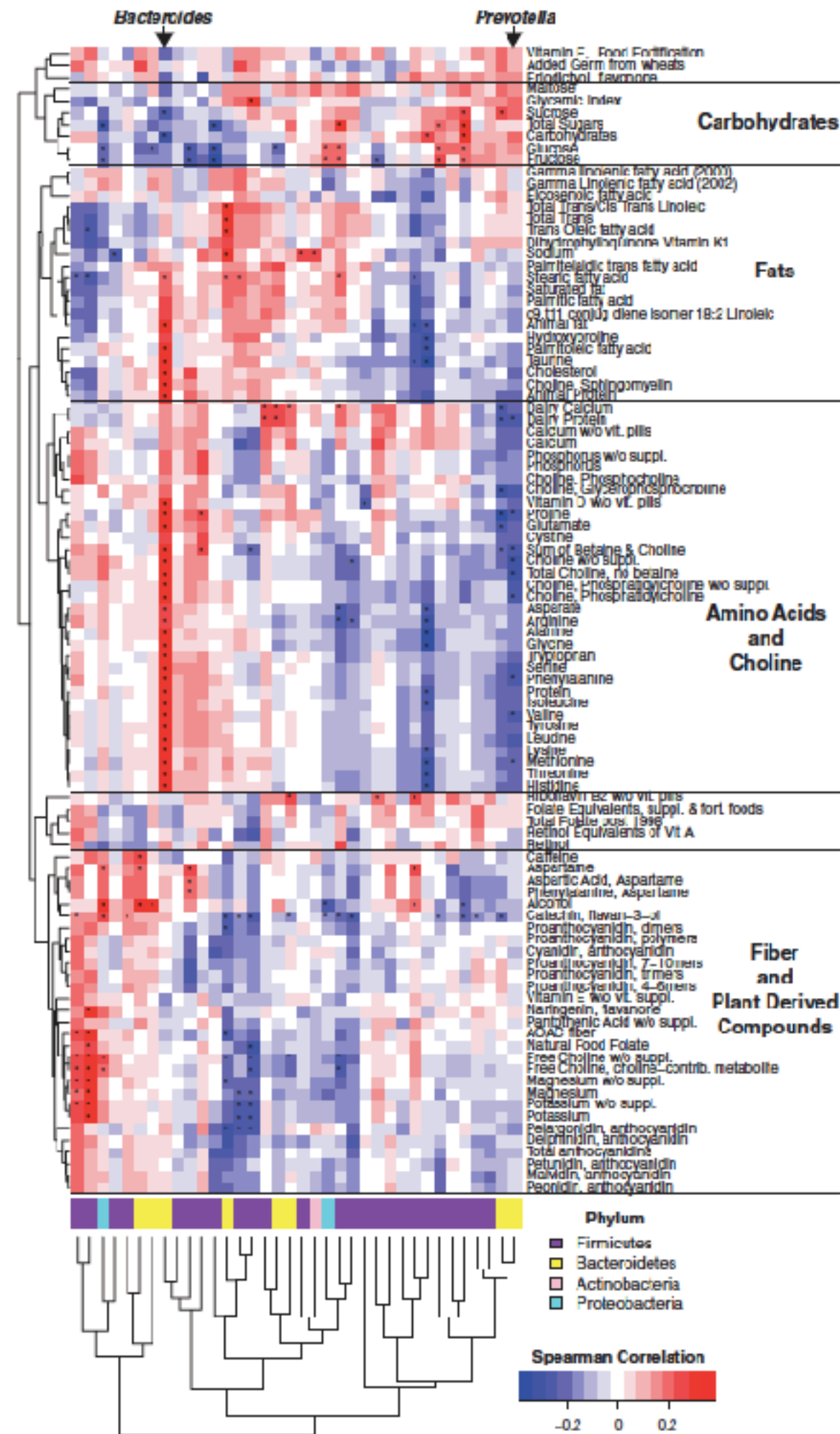
# Diet influences microbiota composition

## Linking Long-Term Dietary Patterns with Gut Microbial Enterotypes

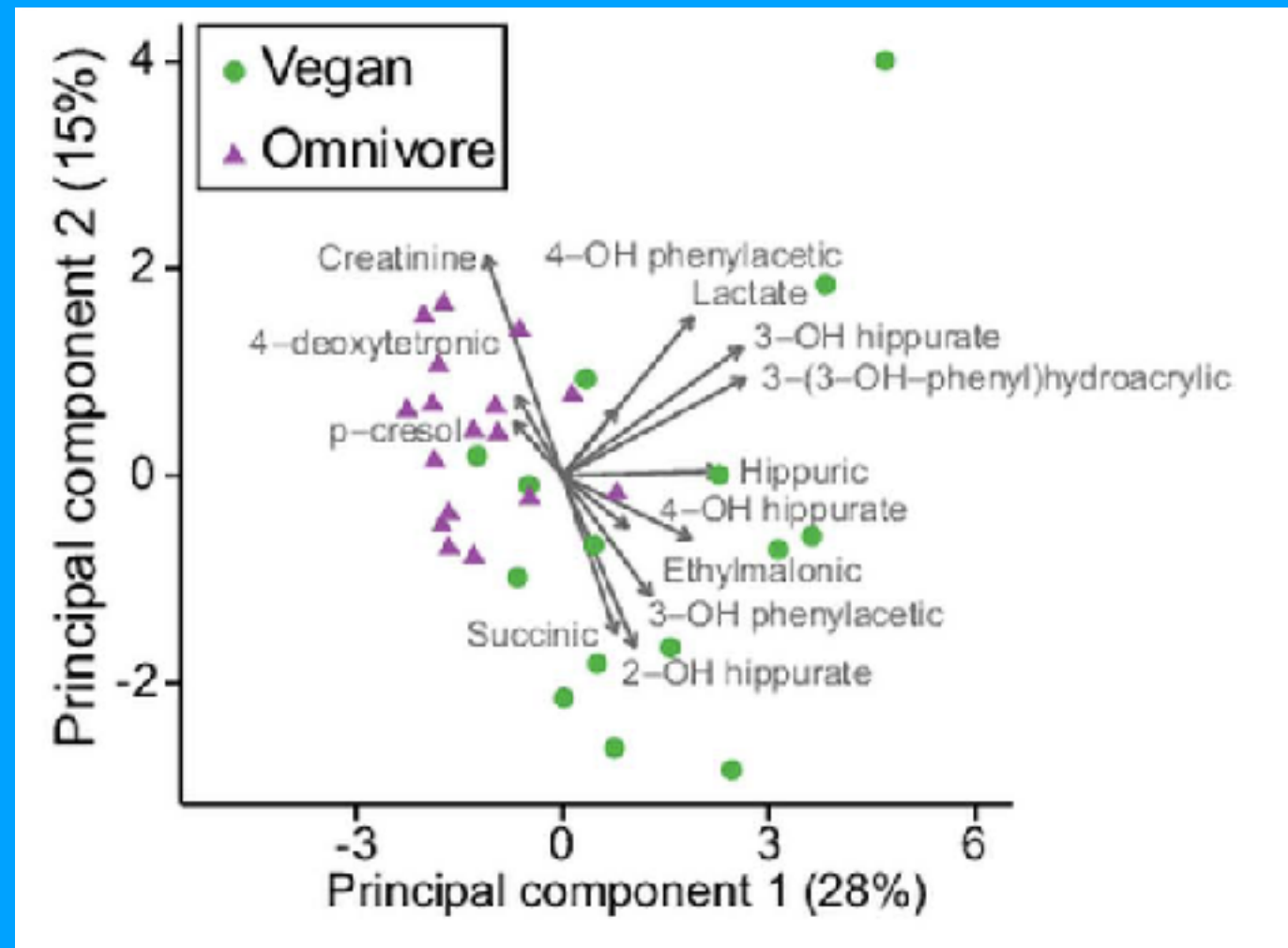
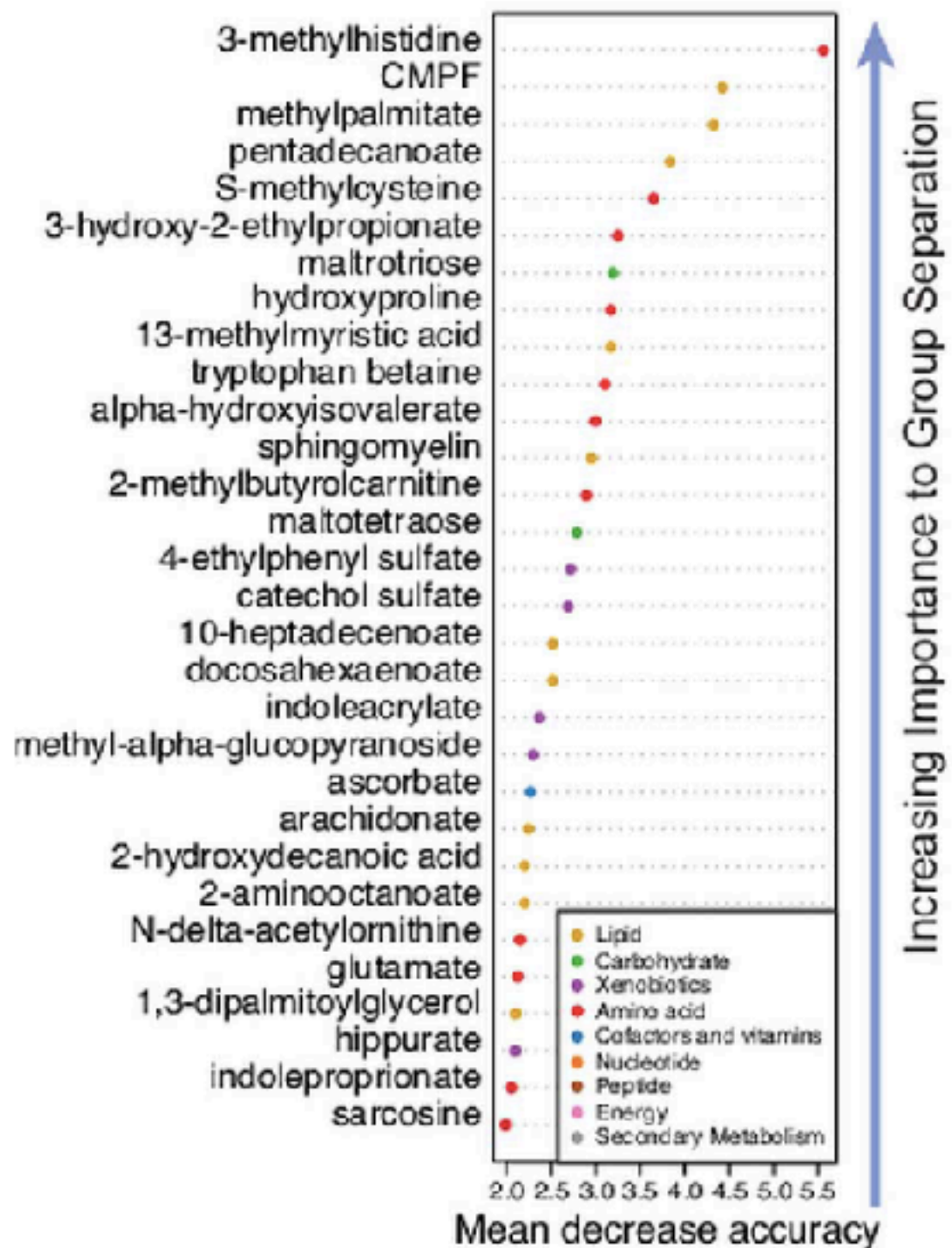
Gary D. Wu,<sup>1\*</sup> Jun Chen,<sup>2,3</sup> Christian Hoffmann,<sup>4,5</sup> Kyle Bittinger,<sup>4</sup> Ying-Yu Chen,<sup>1</sup> Sue A. Keilbaugh,<sup>1</sup> Meenakshi Bewtra,<sup>1,2</sup> Dan Knights,<sup>6</sup> William A. Walters,<sup>7</sup> Rob Knight,<sup>8,9</sup> Rohini Sinha,<sup>4</sup> Erin Gilroy,<sup>2</sup> Kernika Gupta,<sup>10</sup> Robert Baldassano,<sup>10</sup> Lisa Nessel,<sup>2</sup> Hongzhe Li,<sup>2,3</sup> Frederic D. Bushman,<sup>4\*</sup> James D. Lewis<sup>1,2,3\*</sup>

SCIENCE VOL 334 7 OCTOBER 2011

- 2 main clusters identified
- Bacteriodes cluster associated with diet higher in protein and animal fat; Prevotella cluster associated with diet higher in carbohydrates



# Diet influences metabolites

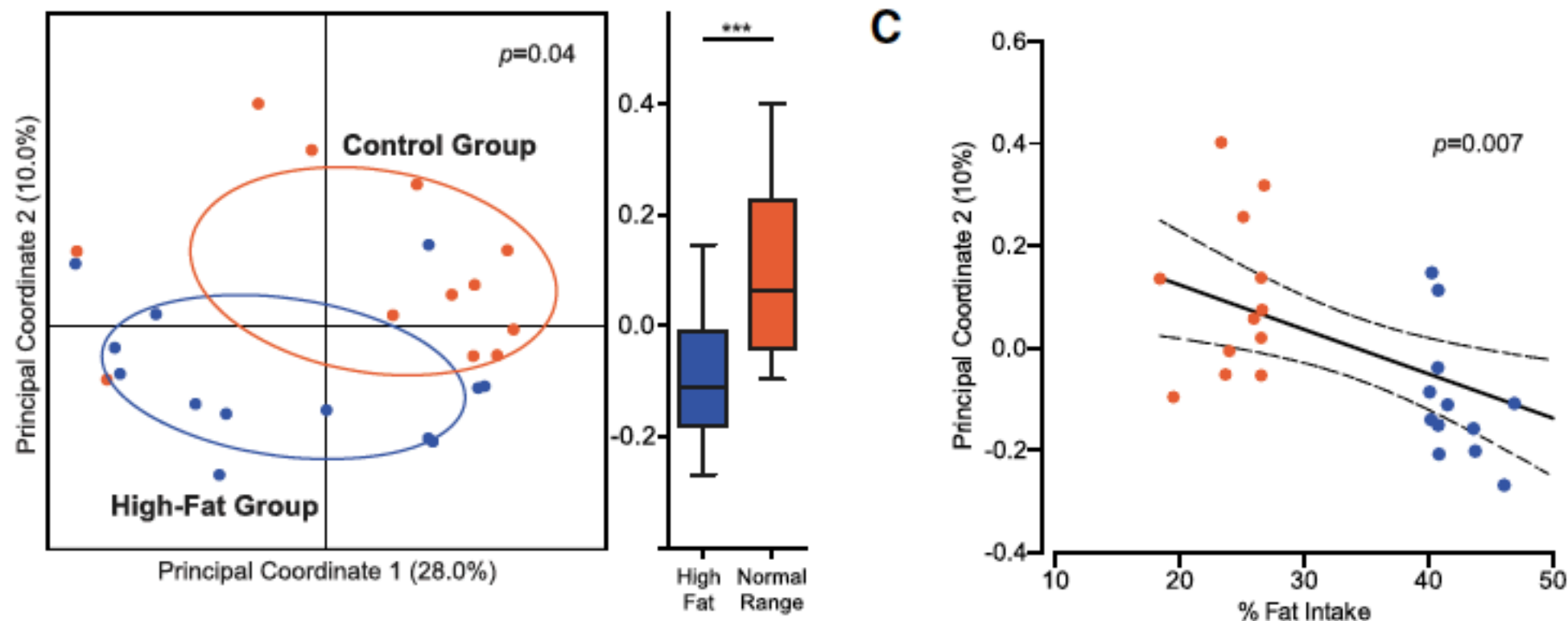


# Maternal diet influences infant's microbiome

Chu et al. *Genome Medicine* (2016) 8:77

## The early infant gut microbiome varies in association with a maternal high-fat diet

Derrick M. Chu<sup>1,2,3</sup>, Kathleen M. Antony<sup>1</sup>, Jun Ma<sup>1</sup>, Amanda L. Prince<sup>1</sup>, Lori Showalter<sup>1</sup>, Michelle Moller<sup>1</sup> and Kjersti M. Aagaard<sup>1,2,3,4,5\*</sup>





# Many factors interact to influence the microbiome

RESEARCH ARTICLE

## Sex, Body Mass Index, and Dietary Fiber Intake Influence the Human Gut Microbiome

Christine Dominianni<sup>1</sup>, Rashmi Sinha<sup>3</sup>, James J. Goedert<sup>3</sup>, Zhiheng Pei<sup>2,4,5,6</sup>, Liying Yang<sup>6</sup>, Richard B. Hayes<sup>1,2</sup>, Jiyoung Ahn<sup>1,2\*</sup>

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PLOS ONE | DOI:10.1371/journal.pone.0124599 April 15, 2015

## Extensive impact of non-antibiotic drugs on human gut bacteria

Lisa Maier<sup>1\*</sup>, Mihaela Pruteanu<sup>1†\*</sup>, Michael Kuhn<sup>2\*</sup>, Georg Zeller<sup>2</sup>, Anja Telzerow<sup>1</sup>, Exene Erin Anderson<sup>1</sup>, Ana Rita Brochado<sup>1</sup>, Keith Conrad Fernandez<sup>1</sup>, Hitomi Dose<sup>3</sup>, Hirotada Mori<sup>3</sup>, Kiran Raosaheb Patil<sup>2</sup>, Peer Bork<sup>2,4,5,6</sup> & Athanasios Ilyas<sup>1,2</sup>

# Many factors are important to gut health and the composition of gut bacteria

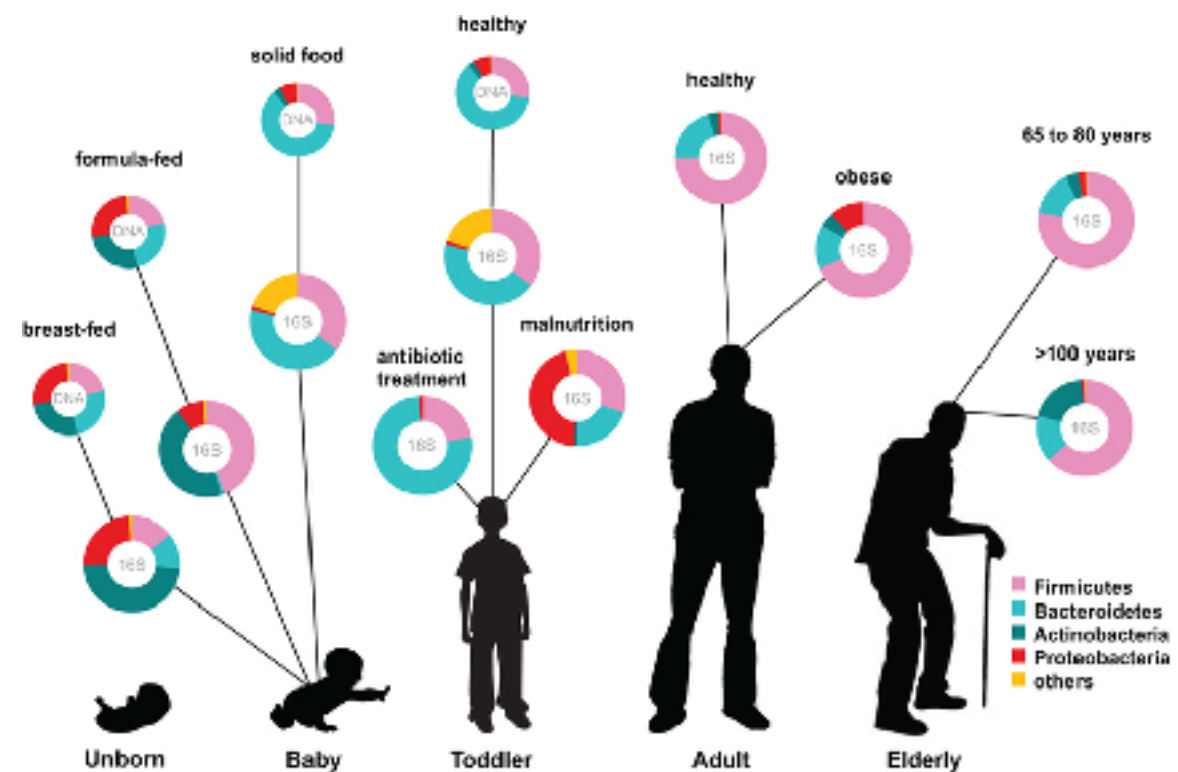
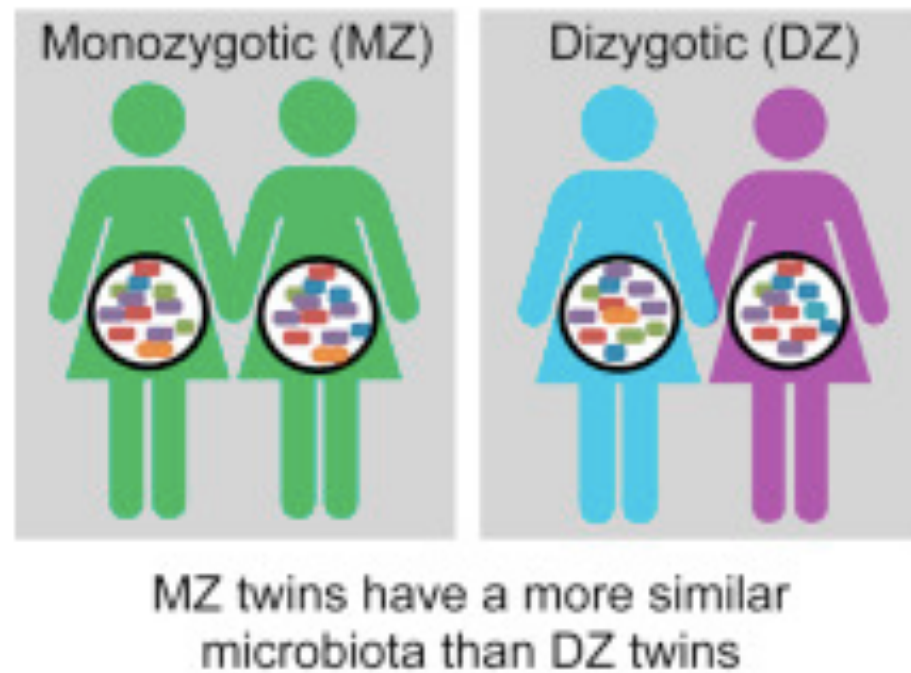
Genetics

Diet

Exercise

Drugs

Stress

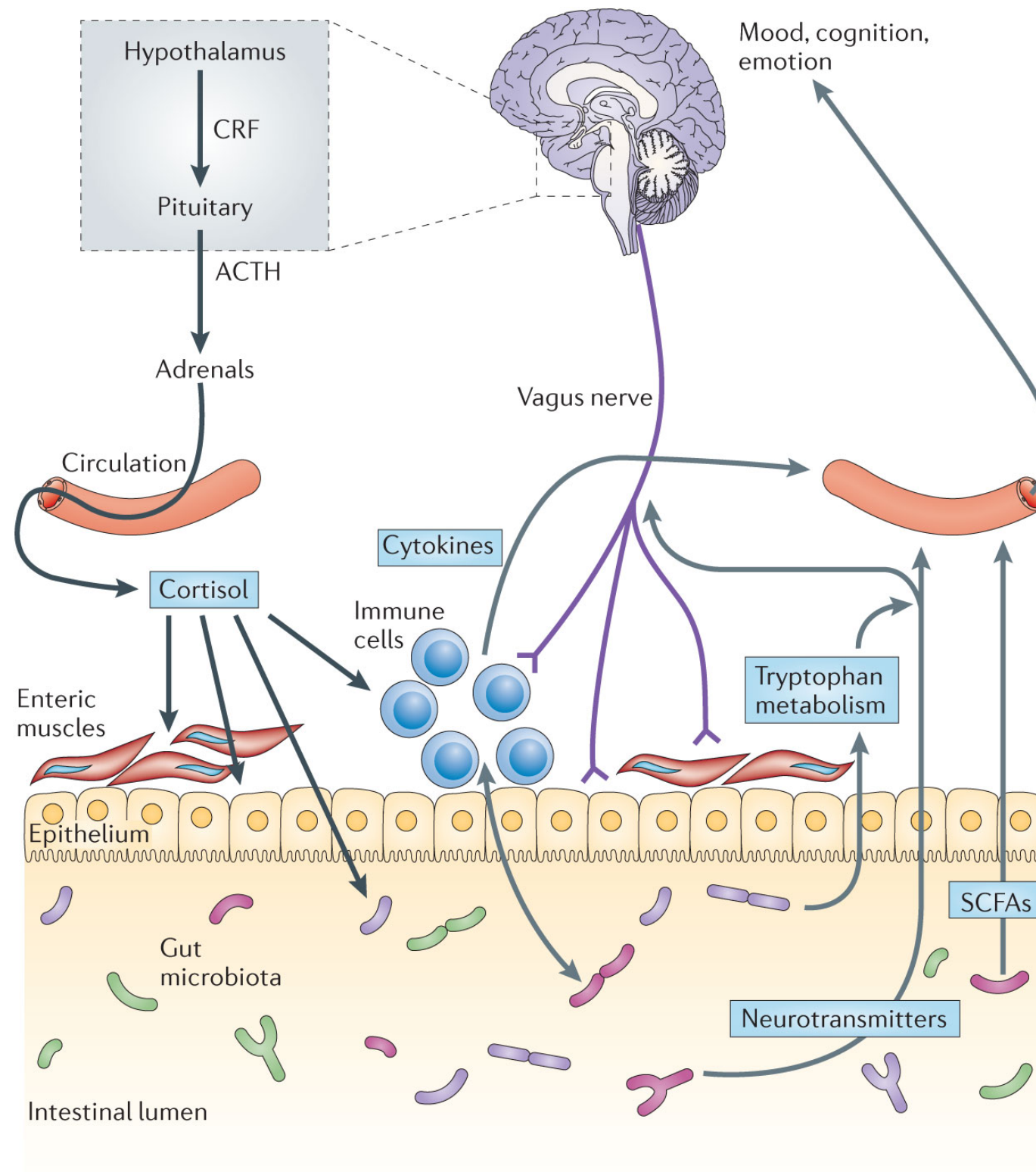


Goodrich et al 2015 Cell 159, 789-799

Age

Sex

# There are many ways that diet can influence microbiota-brain communication





# What is the benefit of looking at microbes and mental health?

- Identification of biomarkers that will help understand individual biological differences and help subgroup clinical populations to predict best treatment
- Identify individuals at risk for early intervention
- Provide novel targets for drug development
- Expansion and development of diet, prebiotic, probiotic, or other intervention strategies for psychiatric illness

# Research Funding Sources



Canada Foundation  
for Innovation

Fondation canadienne  
pour l'innovation

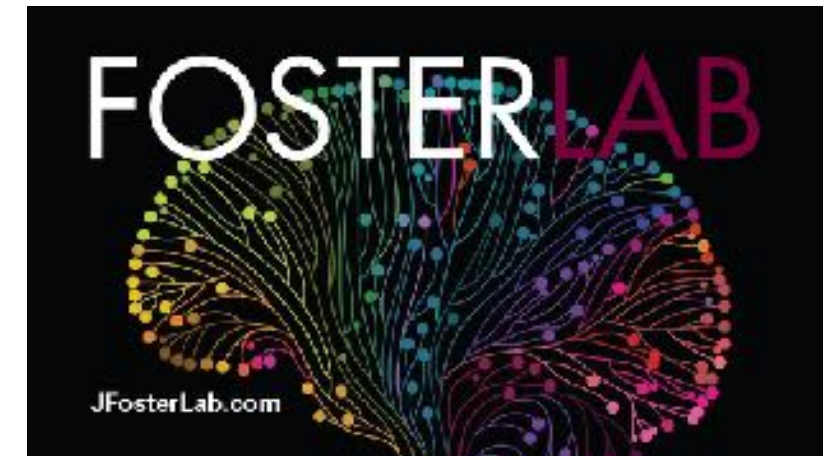


ONTARIO  
BRAIN  
INSTITUTE



Brain Canada





## **Collaborators**

Jason Lerch  
 Jacob Ellegood  
 Mike Surette  
 Aadil Bharwani  
 OBI POND research group  
 CAN-BIND research group

## **Former Lab Members**

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 Kelly Rilett  
 Karen-Anne McVey-Neufeld  
 Linda Zhou  
 Robyn MacKenzie  
 James St. Pierre  
 Daiana Pogacean

## **Current Lab Members**

Shawna Thompson  
 Rachael Horne  
 Cassandra Francesca  
 Bryce Kwiecien-Delaney  
 Owen Luo  
 Alexandra Moyssakos  
 Emily Long-Huckle



# Trends in **Neurosciences**



**Gut–brain axis: how the  
microbiome influences brain function**

**Cell**  
PRESS

Using clear bottles as a metaphor for the physical inner workings of the human body, BrainGut Axis depicts the re-conception of the brain as part of an interconnected system, rather than the more traditional view of the brain as superior to, or in control of all other processes. There is movement throughout this system of various shapes and colors meant to insinuate components at work within the body such as microbiota and hormones, as well as to symbolize moods, impulses and the more abstract processes happening within us. In addition, these “inner-workings” leak outside of the system, reminding us that things are not fully contained, but are continuously moving beyond us, and wandering into us, as we maneuver our way through the world.

Thanks to:

Dr. Renee Jackson (PhD Art Education)  
Dr. Suzanne McCallaugh (PhD Philosophy)  
Dr. Dominic Letarte (PhD Computer Science)