



Fitness Matter: Fueling you gut microbes

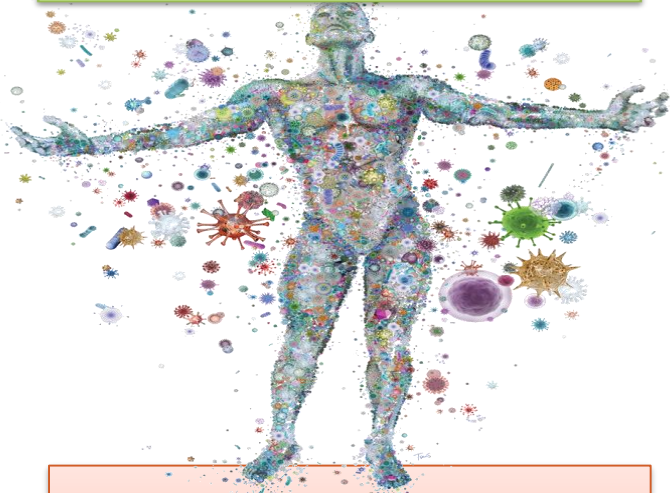
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Our Microbiome

Human

~10 trillion cells
~23 thousand genes



Microbiome

~100 **trillion** cells
~3 million genes

Bacteriome

Archaeome

Mycobiome

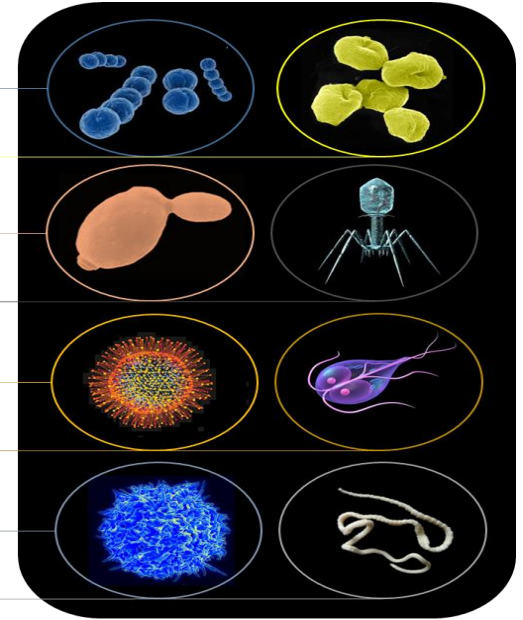
Phageome

Euvirome

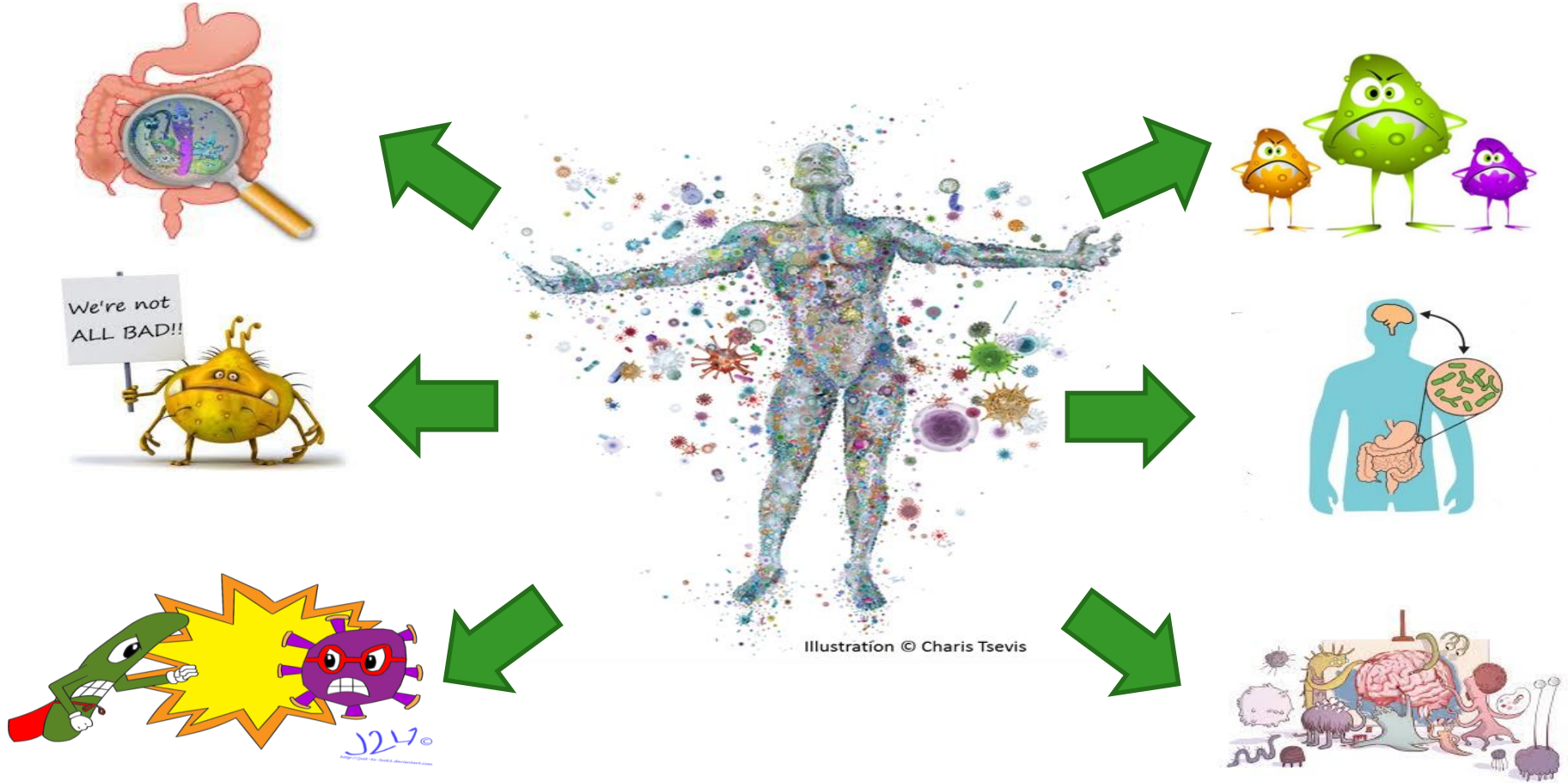
Protozome

Immunome

Helminthome



Why do we study the microbiome ?



How do we study the microbiome?

- Traditional



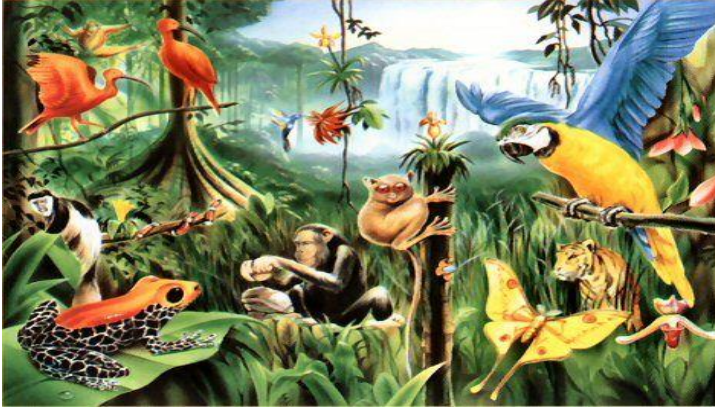
- Modern



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TCCGAGTTCCCTGGAACTGGGACGCA  
GGAATAACCGTGGTAATCTAGAGCTA  
ACGCCATAGAGGGTGAGAGCCCTGAG  
TTCCGAGTTCCCTGGAACTGGGACGCA  
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CGTCTGGTAGGACACCCAGCCCTG
```

What makes a healthy microbiome ?

Diversity is Key



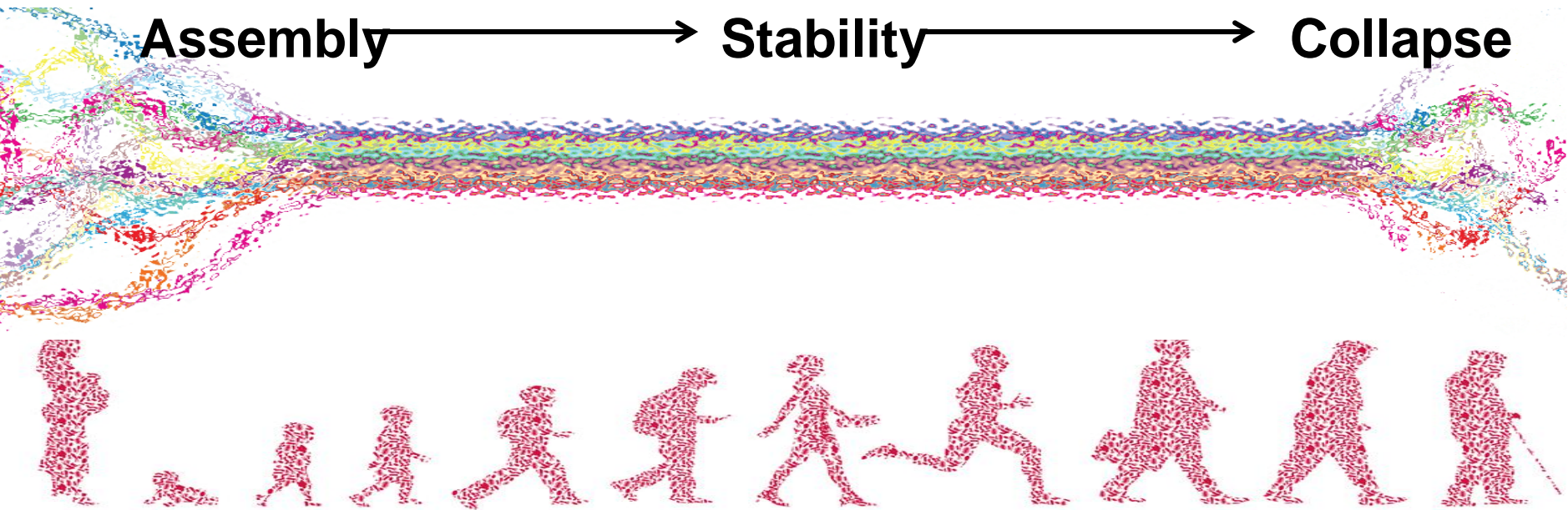
High Diversity



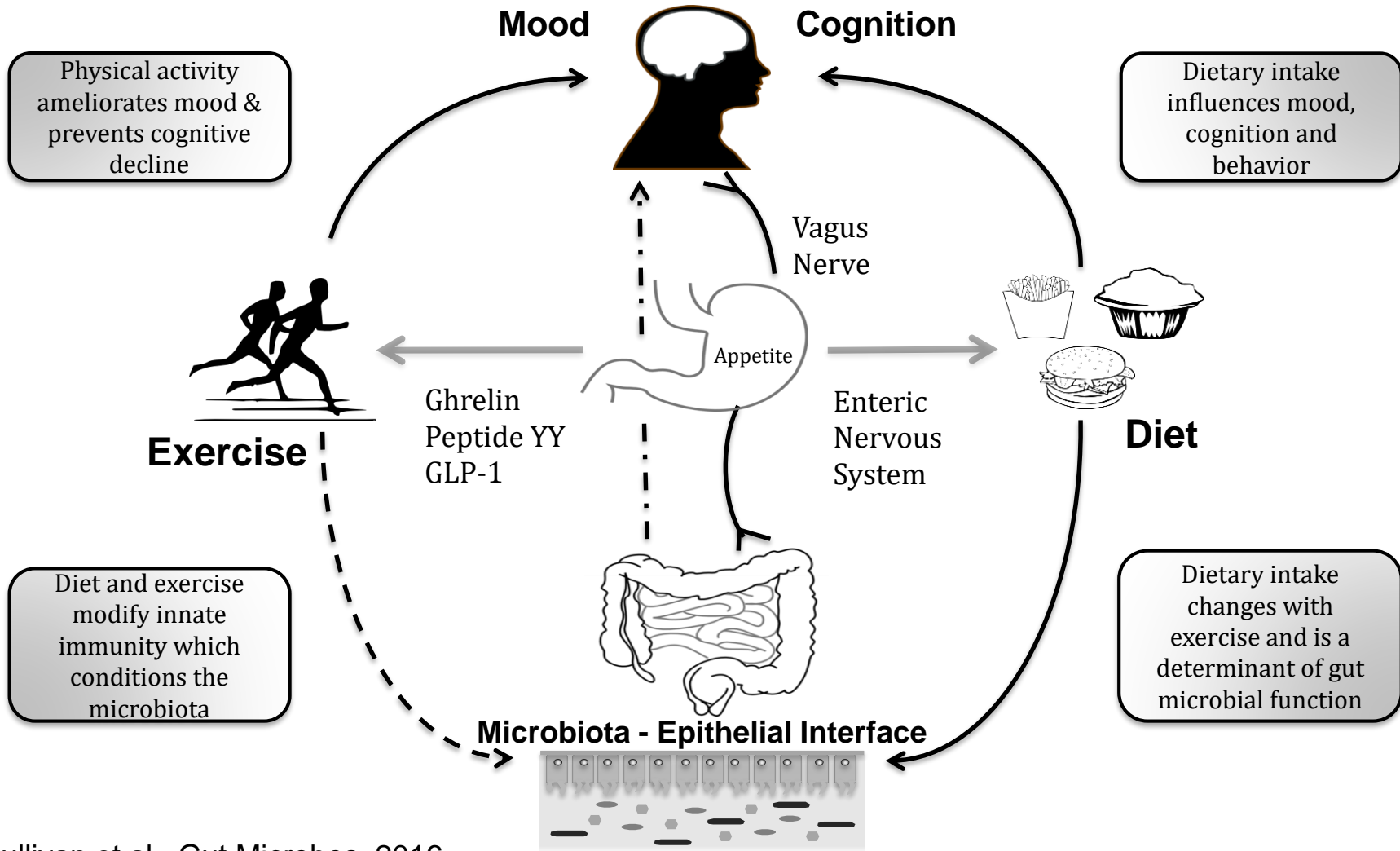
Low Diversity

Low total diversity within the gut microbiota is generally regarded as less desirable and has been observed in children that are more susceptible to allergies as well as sufferers of IBD, IBS and *C. difficile* infection (among others)

Modulating our microbes



Opportunities for interventions/diagnostics for food, biotech & pharma



Fitness Matters



VS.

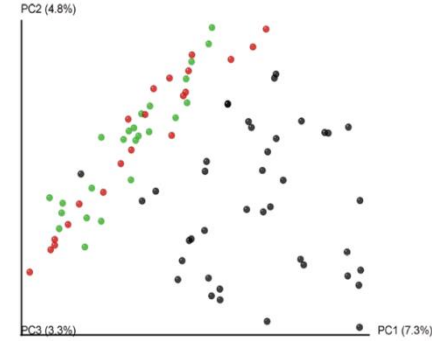


Figure 3 Unweighted UniFrac separates the athlete and control microbiota. Unweighted UniFrac principal coordinate analysis (PCoA) of fecal microbiota from 96 athletes (green and red) and 96 controls (black).

What was found?

- Athletes have more diverse gut microbes than controls at compositional, functional and metabolic levels
- This diversity was associated with protein intake and/or fitness levels
- Microbiome of athletes was primed for muscle repair, protein degradation and vitamin recovery



Can we train out gut?

8 weeks

Exercise (n=25)

Exercise + protein (whey; n=27)

Whey protein (n=22)



What was found?

- No change in the microbiome was noted
- It was concluded that microbiome diversity can be attributed to the level of physical fitness rather than frequency of exercise undertaken





Fitness Matters– V02 Max

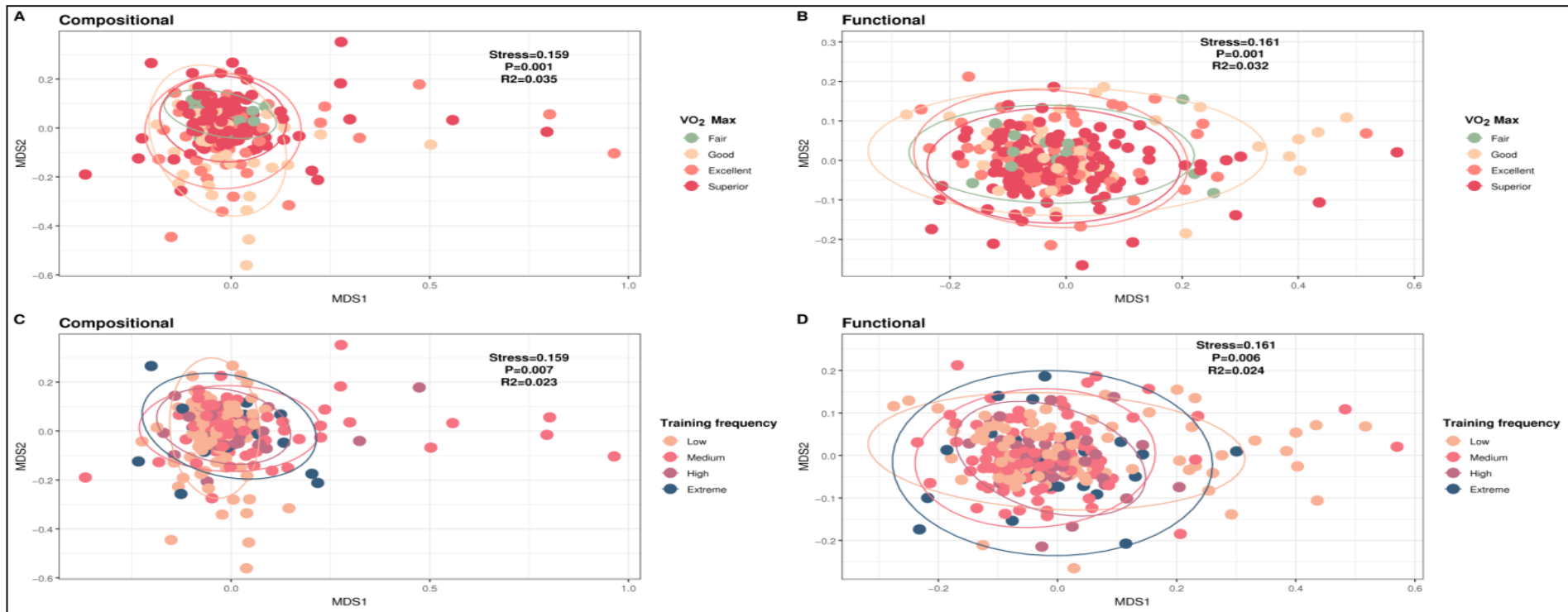
WHY?

To investigate potential factors influencing gut microbiome composition and function in an active cohort.

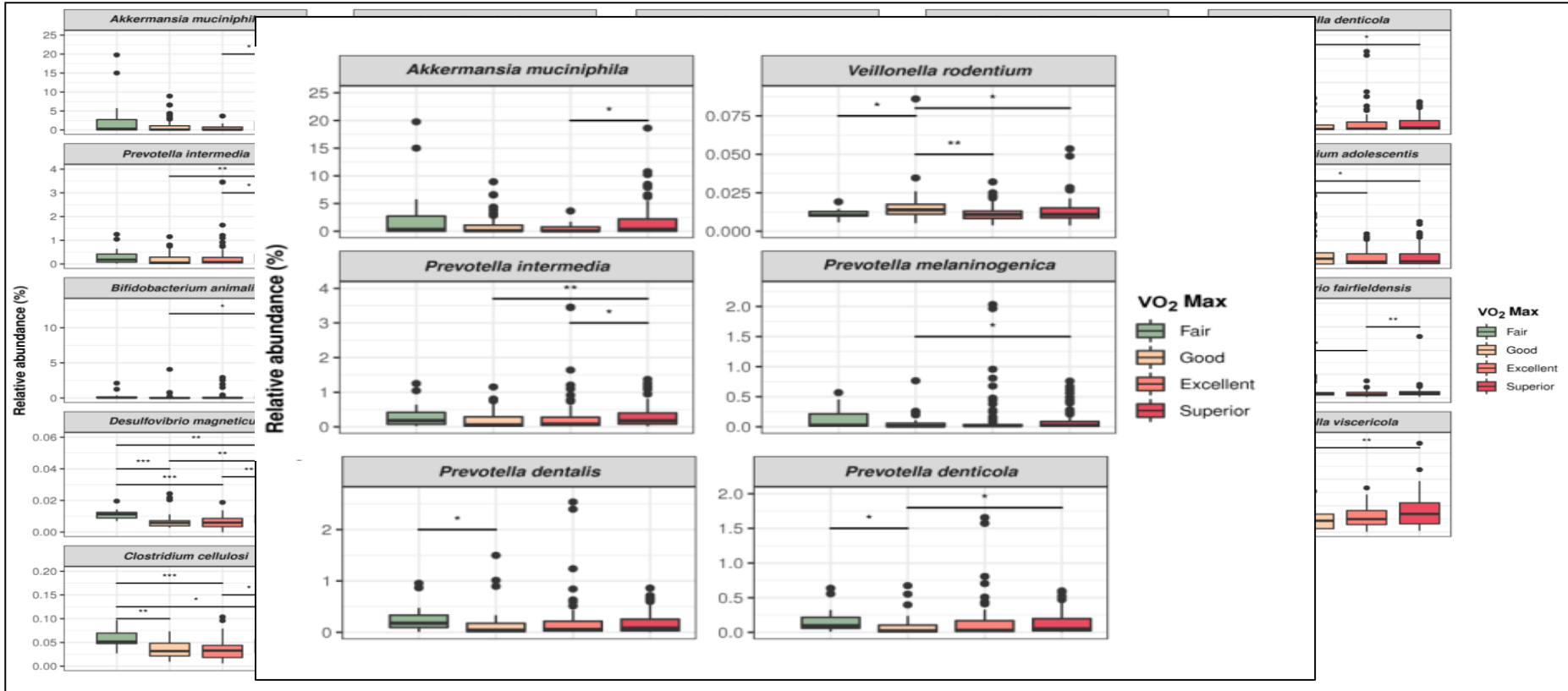
Total participants	62
Total samples	238
Age	34 (± 9) years
Sex	
Male	39
Female	23
VO2 max	46.8 (± 6.6)
BMI (kg/m²)	25.6 (± 3.1)
Height (cm)	174 (± 9)



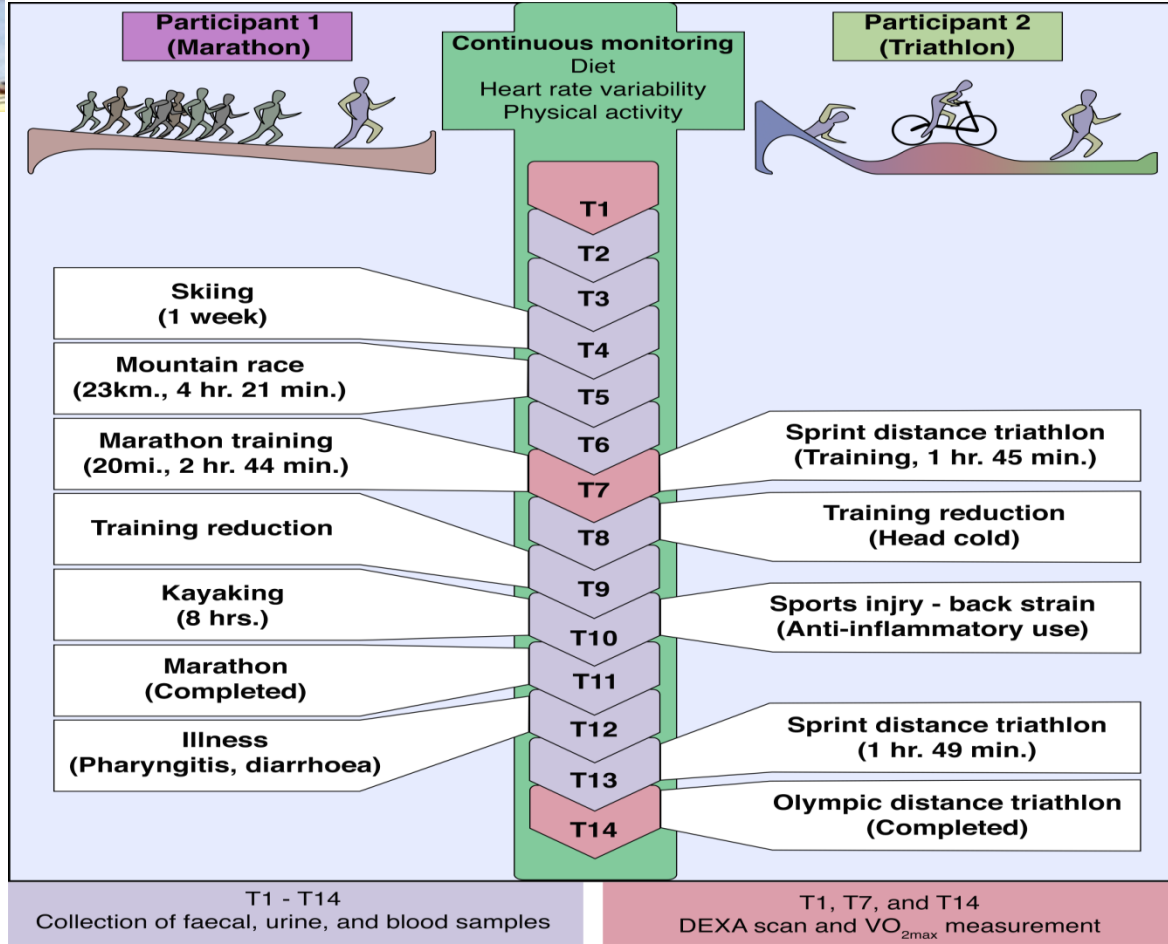
Separation Based on VO₂ Max and Training load



Significantly Different Species Based on VO_2 max



A STEP TOWARDS PERSONALIZED MEDICINE FOR ATHLETES

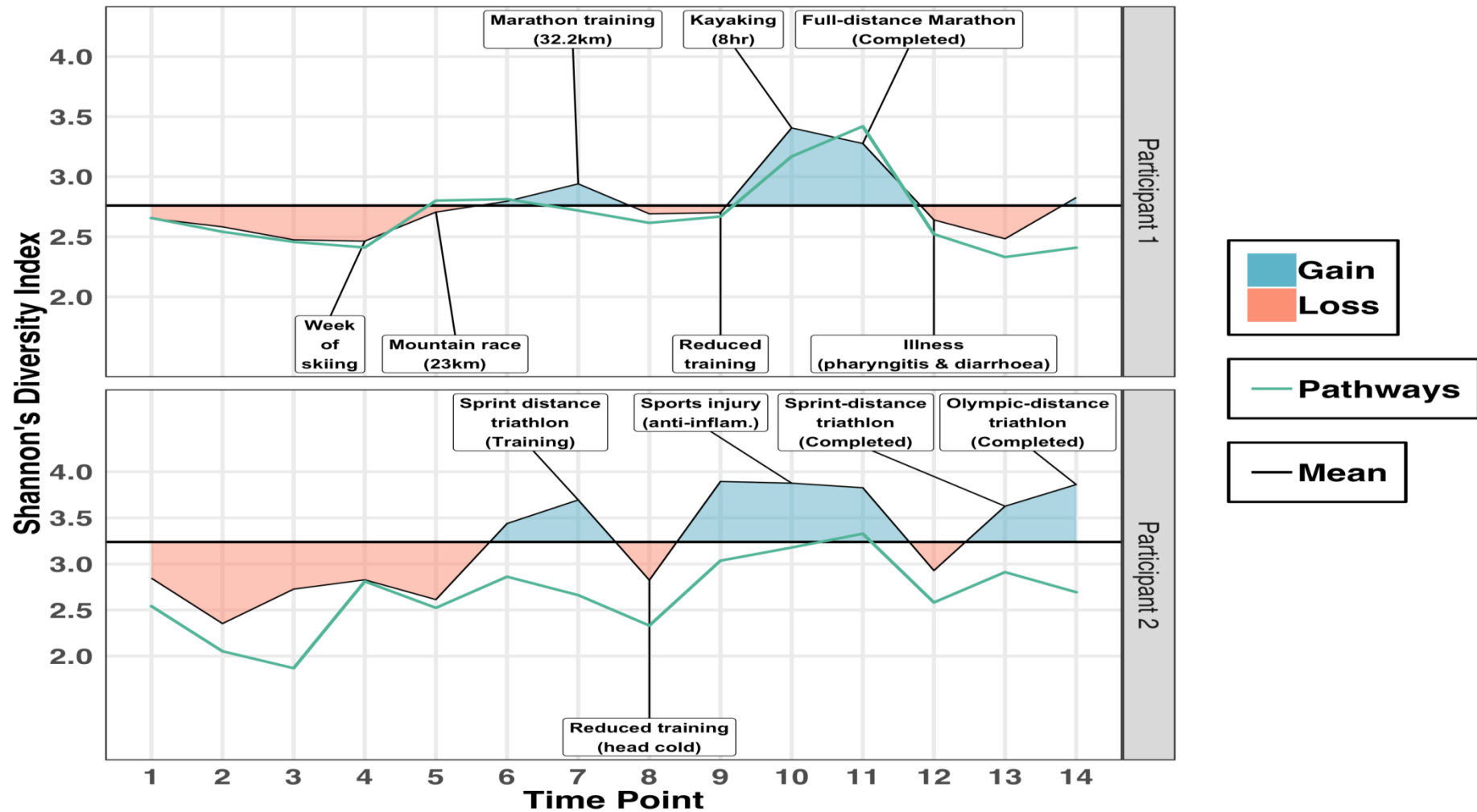


Improved body composition

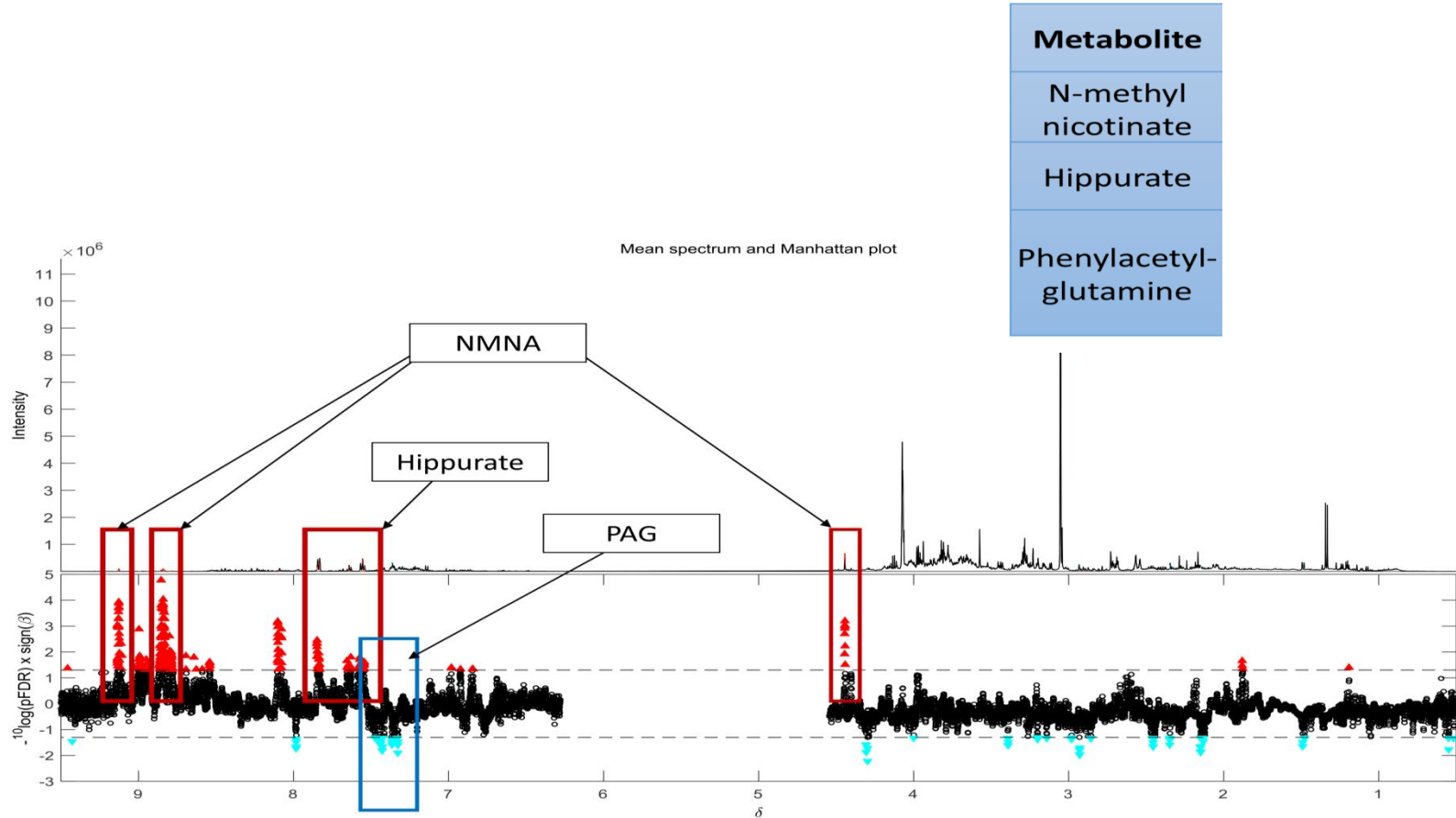
Improved cardiorespiratory
fitness

Patient characteristics	Values					
	Participant 1 (Marathoner)			Participant 2 (Triathlete)		
	T ₀	T ₁₄	Δ	T ₀	T ₁₄	Δ
Age (years)	30	—	—	33	—	—
Height (cm)	181	—	—	182	—	—
Weight (kg)	93.8	89.2	-4.6	104.9	103.4	-1.5
BMI (kg/m ²)	28.6	27.2	-1.4	31.7	31.2	-0.5
Waist:Hip ratio	0.92	0.92	0.0	0.95	.91	-0.04
Body fat (%)	25.6	21.7	-3.9	34.7	34.5	-0.2
Fat mass (kg)	23.9	19.4	-4.6	36.3	35.7	-0.6
Fat mass (trunk) (kg)	14.8	11.7	-3.1	20.9	20.4	-0.5
Lean tissue mass (kg)	65.6	65.9	0.2	64.97	64.2	-0.7
Estimated VO _{2max} (mls/kg/min)	41.1	46.6	5.5	33.6	38	4.4
Max HR (bpm)	183	179	-4	196	179	-17
Resting HR (bpm)	69	50	-19	58	72	-2
Systolic BP (mmHg)	122	116	-6	128	127	-1
Diastolic BP (mmHg)	77	75	-2	87	72	-15
Weekly PA (IPAQ, METS)	891.5	—	—	646.5	—	—
Weekly PA (IPAQ, kCals)	1,393.7	—	—	1,130.3	—	—

Diversity of Bacteria Species



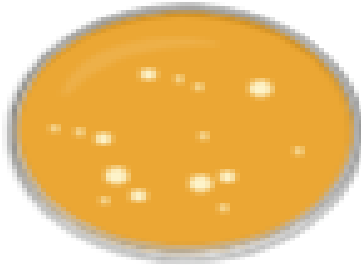
Metabolomics





Probiotics from the athlete gut?

Why athletes?



Why do athletes need probiotics?



DEPRESSION/ANXIETY



INFECTION



GASTROENTERITIS/ GUT
PERMEABILITY DYSFUNCTION



INFLAMMATION



ENERGY METABOLISM



OXIDATIVE STRESS

How we are isolating potential novel probiotics?

- Microbiological culture methods
- Biobank of elite athlete faecal samples
- Testing potential isolates for probiotic traits



Fitness and gut health

Influencing factors

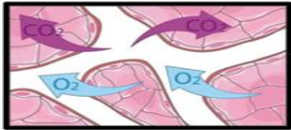
Type of sport



Travel



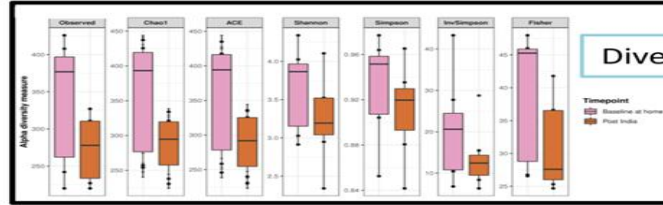
VO₂ max



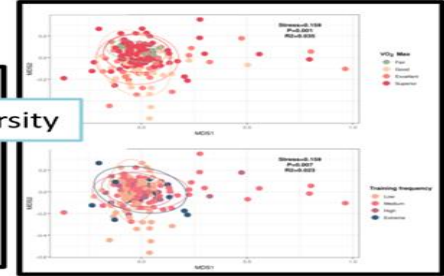
Training frequency



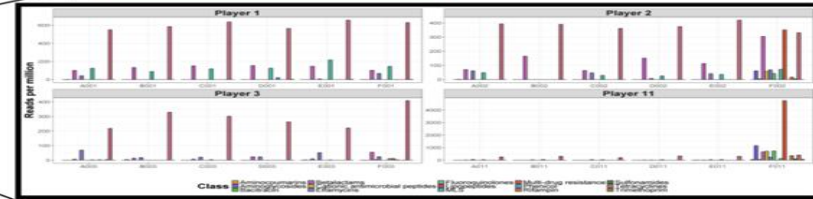
Gut microbiome aspects impacted



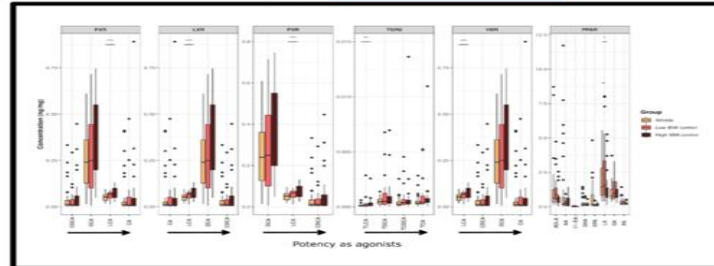
Diversity



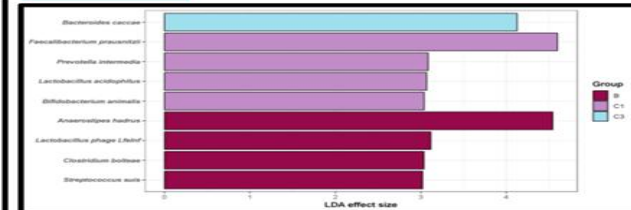
Antibiotic resistance



Metabolites



Species



Acknowledgements



All athletes both elite and non



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