## Sportomics, Metabolic Health, and Utilization of Functional Medicine Testing

Sportomics, Metabolic Health, and Utilization of Functional Medicine Testing

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## Sportomics, Metabolic Health, and Utilization of Functional Medicine Testing

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## Objectives for This Presentation

- At the conclusion of this program, participants will be able to:
- Define Sportomics and its application to Exercise Performance and Metabolic Health
- Identify and understand changes in metabolome and microbiome associated with exercise
- Apply metabolomic and microbiome Functional Medicine testing to inform recommendations for improved exercise performance and metabolic health


## OUTLINE

- Introduction to "SPORTOMICS"
- Exercise and Metabolome
- Exercise and Microbiome
- Sportomics and Exercise Performance
- Role of functional medicine testing
- Sportomics and Metabolic Health
- Role of functional medicine testing


## SPORTOMICS . . .

- Applies metabolomics to investigate the metabolic effects of physical exercise on individuals
- Works to advance knowledge in integrative physiology and the systems biology of movement with a goal to translate markers associated with metabolic challenges of training, or competition, to similar stresses of disease settings



## SPORTOMICS: CLINCIAL RELEVANCE

- EXERCISE = Medicine
- With a great "benefit to risk" profile
- Three major clinically relevant questions:
- What is "TIPPING POINT"?
- Where/When/How does exercise become less health promoting?
- Who are the "RESPONDERS" vs. "NONRESPONDERS"?
- How do you make a "NON-RESPONDER" a "RESPONDER"?
- EXERCISE = INVESTIGATIVE TOOL as a STRESSOR
to systems biology and integrative physiology



## SPORTOMICS

## MULTI-VARIATE

SYSTEMS BIOLOGY


## SPORTOMICS

- Must go beyond correlation/association studies
- Meta-omics and computational tools
- Potential for machine learning to predict exerciseinduce alteration and performance measures to distinguish responders and non responders
- by Zeevi et al



## SPORTOMICS: METABOLOME

- SUMMARY:
- Substantial heterogeneity of studies
- Volume was biggest driver of changes
- Generally short-lived response to acute exercise
- However, changes are apparent in consistent movers
- Multifactorial dose response relationship


## SPORTOMICS: METABOLOME

## - ACUTE EXERCISE:

- Responses resolved minutes to hours later
- Greater changes in less trained in response to marathon ${ }^{1}$
- INCREASE:
- Lactate, pyruvate, TCA intermediates, fatty acids, acyl-carnitines, ketone bodies ${ }^{2}$
- DECREASE:

- Bile acids ${ }^{2}$


## SPORTOMICS:

## METABOLOME \& PERFORMANCE

- CHRONIC EXERCISE:
- "Coherently healthier metabolic profile"
- Lower amino acids (especially isoleucine)
- Changes in several lipid metabolites
- Saturated $\rightarrow$ polyunsaturated profile
- Lower VLDL \& TG and Higher HDL
- Difference in intermediary metabolism, fuel substrate utilization, glucose transport, fatty acid oxidation, oxidative stress, steroid biosynthesis, insulin signaling
- Epigenomic, Transcriptomic, \& Proteomic studies are fewer but confirm gene changes


Glucose Lysophosphatidylethanolamines


## SPORTOMICS: METABOLOME



## SPORTOMICS:

## METABOLOME \&

 PERFORMANCE \& FUNCTIONAL TESTING- 38yo year-round athlete
- In winter with significant XC and downhill skiing routine



## SPORTOMICS: MICROBIOME

- 2014: $1^{\text {st }}$ report of exercise increasing gut microbial diversity in humans
- Low BMI control and athletes with high BMI significantly higher proportions of Akkermansia muciniphila
- Greater microbiome diversity
- HOWEVER, suggestion that exercise and protein intake were drivers of diversity



## SPORTOMICS: MICROBIOME

- Greater diversity compared to nonathletes ${ }^{1,2}$
- Greater growth of certain species such as Akkermansia muciniphila 1,3,4
- Relative increase in SCFA ${ }^{2,5}$
- Direct association between VO2max and $\mathrm{F} / \mathrm{B}$ ratio ${ }^{6}$




## SPORTOMICS: MICROBIOME



## SPORTOMICS:

## MICROBIOME \& PERFORMANCE



Fermentable fiber \& microbes


Fermentable fiber \& microbes

## SPORTOMICS:

## MICROBIOME \& PERFORMANCE



Benefits for the athlete

- Production of bioactive metabolites (i.e.short chain fatty acids, neurotransmitters)
- Maintenance of intestinal barrier function
- Modulation of immune system
- Improved energy harvest and utilization
- Regulation of muscle metabolism


## EXERCISE \& PERFORMANCE

## Benefits on the gut microbiota

- Higher microbial richness
- Higher abundancy of beneficial Akkermansia, Veillonella, Prevotella
- Selection advantage for lactate-utilizing bacteria



## SPORTOMICS:

## MICROBIOME \& PERFORMANCE

GENOVA



## SPORTOMICS:

## MICROBIOME \& PERFORMANCE

- INCREASED Veillonella spp (esp. Veillonella atypica) post marathon
- Veillonella atypica gavage $\rightarrow$ increased exercise performance in mice



## SPORTOMICS:

## MICROBIOME \& PERFORMANCE

- IMPROVED EXERCISE PERFORMANCE in mice with Veillonella atypica gavage
- PROPOSED MECHANISM OF ACTION:
- LACTATE production and conversion into propionate



## SPORTOMICS:

## MICROBIOME \& PERFORMANCE

- Veillonella spp. = oral nitrate-reducing bacteria
- ADDITIONAL PROPOSED MECHANISM OF ACTION: Nitric oxide (NO) production by the oral microbiome

Dietary nitrate

$\longrightarrow$

eNOS

Increased mitochondrial efficiency


Increased muscle force

SPORTOMICS:
MICROBIOME \& PERFORMANCE \& FUNCTIONAL TESTING

| Gastrointestinal Microbiome (PCR) |  |  |  |
| :---: | :---: | :---: | :---: |
| Sommensal Bacteria (PCR) | Result $\mathrm{CFH} / \mathrm{g}$ stool |  | Reference Range CFU/g stool |
| Bacteroidetes Phylum |  |  |  |
| Bacteroides-Prevotella group | 2.4E8 | $\longmapsto \quad 1$ | 3.4E6-1.5E9 |
| Bacteroides vulgatus | 1.2E9 | \# , , , , | $<2.2 \mathrm{EP}$ |
| Barnesiella spp. | 3.6E7 | $\bigcirc \bigcirc$ | < $=1.6 \mathrm{E} 8$ |
| Odoribacter spp. | 7.157 | ¢ , , , - | <=8.0E7 |
| Prevotella spp. | 1.4E8 H | 1 1 1 | 1.4E5-1.6E7 |
| Firmicutes Phylum |  |  |  |
| Anaerotruncus colihominis | 3.4 EF H | \# 1 , 1 | $<=3.2 \mathrm{EF}$ |
| Butyrivibrio crossotus | 5.0 E 7 H | 1 , , | 5.5E3-5.9E5 |
| Clostridium spp. Lactate PRODUCER ${ }^{2.158}$ |  | $\longmapsto \quad 1$ | 1.7E8-1.5E10 |
| Coprococcus eutactus | 1.0E8 | ¢ 1 , ¢ , | $<=1.2 \mathrm{E} 8$ |
| Faecalibacterium prausnitzii | 7.5E8 | ¢ , 1 1 | 5.8E7-4.7E9 |
| Lactobacillus spp. | 1.6E8 | セ, 1 | 8.3E6-5.2E9 |
| Pseudoflavonifractorspp. | 3.0 E 8 H | ¢ 1 , | 4.2E5-1.3E8 |
| Roseburia spp. Lactate D | 7.6E7 L | - 1 , 1 | 1.3E8-1.2E10 |
| Ruminococcus spp. | 1.9 E 9 H | 1 1 1 | 9.5E7-1.6E9 |
| Veillonella spp. | 1.5 E 8 H | 1 1 , | 1.2E5-5.5E7 |
| Actinobacteria Phylum |  |  |  |
| Bifidobacterium spp. | 1.5E8 | $\because \bullet 1$ | < $=6.4 \mathrm{E} 9$ |
| Biffobacterium longum | 1.4E8 | ए , , 1 | $<=7.2 \mathrm{E} 8$ |
| Collinsella aerofaciens | 5.1 E 8 | - , , 1 | 1.4E7-1.9E9 |
| Proteobacteria Phylum |  |  |  |
| Desulfovibrio piger | $8.7 \mathrm{E7}$ H | $1 \quad 1 \quad 1$ | < 1.8 EE 7 |
| Escherichia coli | 1.3 E 8 H | - , , | 9.0E4-4.6E7 |
| Oxalobacter formigenes | 5.0 E 7 H | $\bigcirc \quad 1 \quad \longrightarrow$ | <=1.5E7 |
| Euryarchaeota Phylum |  |  |  |
| Methanobrevibacter smithii | 1.4 EB H | $\xrightarrow{+1}$ | < $=8.6$ E7 7 |
| Fusobacteria Phylum |  |  |  |
| Fusobacterium spp. | 2.3 E 7 H | 1 1 , + | < $=2.4 \mathrm{E} 5$ |
| Verrucomicrobia Phylum |  |  |  |
| Akkermansia muciniphila | 3.1 E7 | $\bigcirc 1+$ | $>=1.2 \mathrm{E} 6$ |
| Firmicutes/Bacteroidetes Ratio |  |  |  |
| Firmicutes/Bacteroidetes (F/B Ratio) | 11 L | -1 1 1 | 12-620 |

## SPORTOMICS:

## MICROBIOME \& PERFORMANCE \& FUNCTIONAL TESTING

- 40yo with chronic fatigue syndrome, pain amplification syndrome and exercise intolerance - previously failed graded exercise program


| Gastrointestinal Microbiome (PCR)** |  |  |  |
| :---: | :---: | :---: | :---: |
| Commensal Bacteria (PCR) | Result cFU/g stool | 1 st QUINTILE DISTRIBUTION <br> 2nd 3rd 4 th 5th | Reference Range CFU/g stool |
| Bacteroidetes Phylum |  |  |  |
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| Barresiella spp. | 1.2 ET | ¢ , 1 , , | < $=1.6 \mathrm{E} 8$ |
| Odoribacter spp. | 7.2E6 | ए, 1, 1 | < $=8.0 \mathrm{E} 7$ |
| Prevotella spp. | 4.8E6 | $\longmapsto$ 1 , 1 1 | 1.4E5-1.6E7 |
| Firmicutes Phylum |  |  |  |
| Anaerotruncus colihominis | 9.4 E 5 | ■ - , 1 | $\ll 3.2 \mathrm{E} 7$ |
| Butyrivibrio crossotus | <DL L | -1 1 1 | 5.5E3-5.9E5 |
| Clostridium spp. | 6.8E8 | $\longmapsto \quad 1$ | 1.7E8-1.5E10 |
| Coprococcus eutactus | 3.8E5 | ए, 1 1 | < $=1.2 \mathrm{E} 8$ |
| Faecalibacterium prausnitzii | 6.0 E7 | $\longmapsto \quad 1$ | 5.8E7-4.7E9 |
| Lactobacillus spp. | 1.6E8 | -1 1 1 1 | 8.3E6-5.2E9 |
| Pseudoflavonifactorspp. | 2.3 E7 | ¢ , , , , | 4.2E5-1.3E8 |
| Roseburia spp. | 3.3 E7 L | -11 1-1 | 1.3E8-1.2E10 |
| Ruminococcus spp. | 5.2 E 7 L | -11 1 1 | 9.5E7-1.6E9 |
| Veillonella spp. | <DL L | -11 1 1 | 1.2E5-5.5E7 |
| Actinobacteria Phylum |  |  |  |
| Bifidobacterium spp. | 2.1 E7 |  | $<=6.4 \mathrm{E} 9$ |
| Bifidobacterium longum | <DL | $\longmapsto \quad 1$ | < $=7.2 \mathrm{E} 8$ |
| Collinsella aerofaciens | 1.7E8 | 1, 1, 1 | 1.4E7-1.9E9 |
| Proteobacteria Phylum |  |  |  |
| Desulfovibrio piger | 8.6 E 5 | 1 1 - | < $=1.8 \mathrm{E} 7$ |
| Escherichia coli | <DL L | - 1 1 1 | 9.0E4-4.6E7 |
| Oxalobacter formigenes | 1.9 E 6 | $\longmapsto \quad 1$ | <=1.5E7 |
| Euryarchasota Phylum |  |  |  |
| Methanobrevibacter smithii | 6.4E6 |  | < $=8.6 \mathrm{EE} 7$ |
| Fusobacteria Phylum |  |  |  |
| Fusobacterium spp. | 1.1E4 | $\bigcirc \quad 1$ | < $=2.4$ E 5 |
| Verrucomicrobia Phylum |  |  |  |
| Akkermansia muciniphila | 1.4 E 7 |  | > $=1.2 \mathrm{E} 6$ |
| Firmicutes/Bacteroidetes Ratio |  |  |  |
| Firmicutes/Bacteroidetes (F/B Ratio) | 7 L | 1111 | 12-620 |

## SPORTOMICS: METABOLIC HEALTH

- LOW cardiorespiratory fitness independent predictor of cardiometabolic disease and mortality ${ }^{1}$
- Exercise capacity more powerful predictor of mortality from cardiometabolic disease than other established risk factors ${ }^{2}$
- VO2max is highly correlated with skeletal muscle mitochondrial capacity ${ }^{3}$


## SPORTOMICS: METABOLIC HEALTH

- METABOLIC FLEXIBILITY
- Ability to appropriately adjust substrate oxidation relative to substrate availability
- LACTATE SHUTTLE:
- Lactate = gluconeogenic precursor
- Increased with increased energy expenditure or reduction in energy from aerobic oxidation



## SPORTOMICS: METABOLIC HEALTH

## - LACTATE

- Glucose as specific fuel and lactate as universal fuel ${ }^{1}$
- Lactate as a substrate for generating fuel
- Radiolabeled lactate $\rightarrow$ labeling of TCA intermediates greater than radiolabeled glucose ${ }^{2}$


## Lactate-producing cell




Glucose

## Lactate-consuming cell



## SPORTOMICS: METABOLIC HEALTH

- Metabolic Disease
- Obesity
- Insulin Resistance
- With OBESITY/INCREASED VAT:
- Skeletal muscle tissue minimally increases glucose oxidation with insulin and preferentially partitions it with net lactate release ${ }^{1}$
- "VISCIOUS CORI CYCLE" 2
- Due to impaired pyruvate oxidation



## SPORTOMICS: METABOLIC HEALTH

- Metabolic Disease
- Obesity
- Insulin Resistance



## SPORTOMICS:

## METABOLIC HEALTH \& FUNCTIONAL TESTING

- EVIDENCE of "Vicious Cori Cycle" and impaired metabolic health
- ELEVATED Lactic Acid
(especially relative to Pyruvic Acid)
- LOW Citric acid
- Often seen in severe obesity
- LOWER TCA intermediates ${ }^{1}$
- Labeled by some as "hypometabolic state" or "mitochondria retraction"
- +/- HIGHER Malic Acid/Succinic Acid


## SPORTOMICS: METABOLIC HEALTH

## - EXERCISE \& DIABETES PREVENTION

- Overweight, prediabetic treatment-naïve males
- Identified responders to exercise intervention (improved glucose metabolism \& insulin resistance)
- Increased capacity for increased SCFA synthesis
- Increased capacity for BCAA catabolism
- FMT of responders improved mice response to exercise
- Through machine learning then predicted glycemic response to exercise in additional 30 subjects


Saccharolysis




## SPORTOMICS:

## METABOLIC HEALTH \& FUNCTIONAL TESTING

- FUNCTIONAL MEDICINE TESTING INFORMED RECOMMENDATIONS for metabolic health \& insulin resistance
- IF EXERCISING \& METABOLIC DISEASE
- POTENTIAL NON-RESPONDER WITH ..
- Elevated BCAAs and AAAs; Elevated glutamate and Decreased GABA; Elevated cysteine \& methionine
- PROBABLE RESPONDER WITH ...
- Increased Firmicutes \& SCFAs; Increased lactate utilizers // (absence of non-responder characteristics)
- IF NOT EXERCISING \& AT RISK FOR METABOLIC DISEASE
- POTENTIAL RESPONDER WITH ..
- Elevated Bacteroides // Elevated GABA
- IF NON-RESPONDER OR POTENTIAL NON-RESPONDER
- Work to elevate SCFAs (fiber, resistant starches, fermented foods/probiotics, butyrate)
- Decrease exercise volume/intensity


# THANK YOU FOR YOUR TIME AND ATTENTION 

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