



COST Action CA18131 ML4Microbiome

Statistical and Machine Learning techniques in human microbiome studies

Action Chair Marcus Claesson 4th Sept 2020





What is the Microbiome?

Microbiome definition:

The complete set of microbial genes & genomes in a given environment

Bacteriome, virome/phageome, mycobiome etc.

60-90% of Earth's biomass microbial

New technologies allows highthroughput analysis of microbial DNA from samples without culturing



Metagenomics &
Next Generation Sequencing:
DNA from 1000s species in parallel



Why is the Human Microbiome important?

The human microbiome

38 trillion bacterial cells (2-3kg) in a 70kg man

1.3:1 bacterial and human cells in/on body

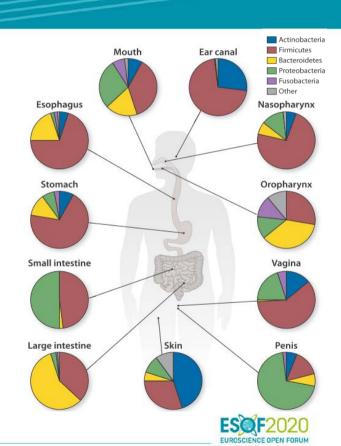
Digests food, synthesizes vitamins, prevents pathogens & trains the immune system

Microbiome importance for improving health

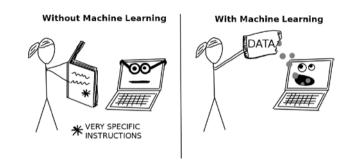
Dysregulation increasingly associated with health & disease => new treatments & diagnostics/prognostics

Optimise nutritional composition & uptake of food

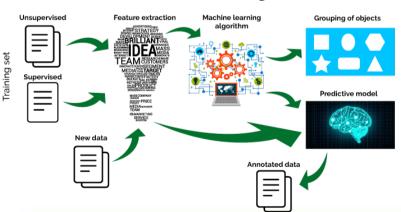
Optimise action, stability & safety of drugs



How can Machine Learning help?



Machine Learning



Microbiome datasets are huge, BUT so is their potential of improving human health!

Many challenges with these data, e.g.

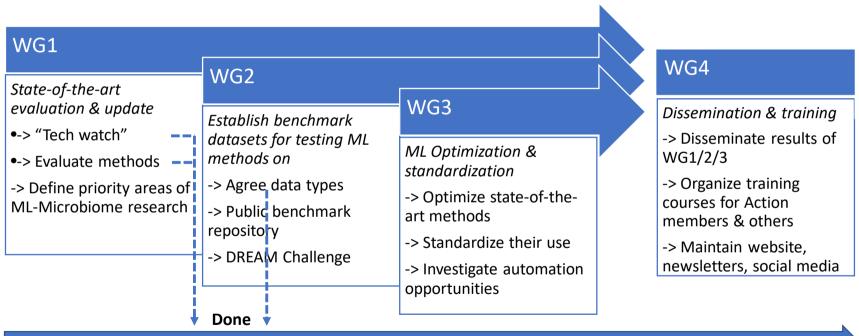
- Different data types
- Lots of noise & inter-individual variation
- # of microbiome samples << # of microbes
- ML models hard to generalise across studies



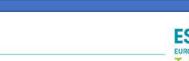
To first optimize and then standardize best practice of ML techniques for human microbiome research



How do we achieve this?



Start 2019

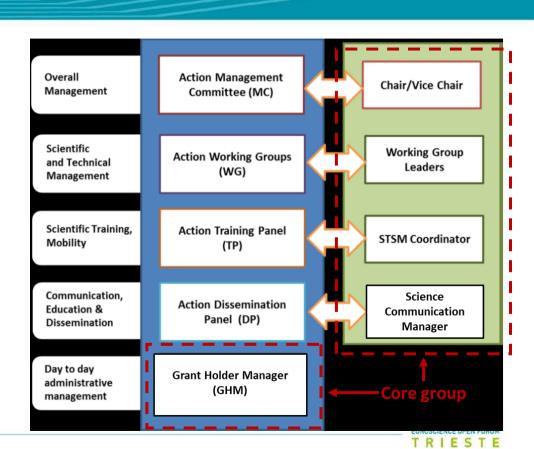


End

2022

Action size & structure

	<u>Start</u>	Now
Countries	24	34
Participants	57	113
ITC	54%	55%
Female	36%	39%
ECI	?	23%



Meetings, training schools & STSMs









+ 7 Short-Term Specific Missions



A huge thanks to

Grant Holder Manager Chloe Huseyin

Vice Chairs David Cabrero Lopez & Randi Betelsen

The Core Group

All MC Members & Substitutes

Karina Marcus & Olga Gorczyca

