Human Teaching, Machine Learning –

developing a sharable syllabus from the recommendations generated through ML4Microbiome (EU-CA18131)











APC 4.11 Claesson Group 8th of June, 2023

Jamie.FitzGerald @ ucc.ie

background

Jamie FitzGerald (see left of screen)

microbiome / microbial ecology Research scientist, FHI







ML4Microbiome STSM: Utrecht University, the Netherlands

August 20th – August 27th, 2022







background

- Frequently focused on **discrimination** within a dataset
 - groups, classes, treatments...
- Diet intervention: ML identified persistent effect of probiotics in rare microbiome
 - not detected with multivariate approaches
- Roosmarijn Luiken, UU (PI: Aldert Zomer @UU, ML4Microbiome)
 - Microbiology, health, and disease
 - similar interest in exploring microbiome through ML





motivation

→ Both Roosmarijn & I interested in ML, but lack background in mathematics &/ machine learning

Somewhat typical:



Microbiome Analysis:

- largely self-taught
- low level of standardisation
- relatively specialised approaches

Machine learning:

- is a huge field in itself
- is very complex (especially for the selfdirected learner)
- often, no clear "best" method

motivation

Enter ML4Microbiome:

- Perspective/state of the art, appropriate methods, new and current applications, etc.
- In-person and online workshops
- Extensive online resources for training
- \rightarrow Suitable syllabus
- \rightarrow Expert guidance
- \rightarrow No need to re-invent any wheels



STSM proposal

1

2

3

condense the ML4Microbiome syllabus, for dissemination to microbiome researchers

create a set of learning exercises to allow training in ML-microbiome

focus on being:

- clear (for microbiome researchers)
- generalisable (for microbiome studies)

STSM in action

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3

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Conversations @ UU:

quickly refined the background, the scope, and the format

issue: "one-size-fits-all" approach not appropriate for a workshop scenario *(even within scope)*

- **solution**: run one method (RF, XGB, SVM); compare outputs to that of the other two methods:
 - highlight where certain approaches succeed, and why (as possible)

1 condensed syllabus

major methods of classification/prediction: Extreme Gradient Boosting, Support Vector Machines, and Random Forest

Connecting ML and the microbiome – appropriate hypothesis testing and study design, and relating this to the underlying classification approaches

data pre-processing - making different microbiome data robust for classification approaches

Understanding and communicating microbiome-ML outcomes (post-processing!)

2 learning exercises

- Identifying / generating hypotheses
- feature engineering and normalisation
- feature selection example
 - sample classification example
 - performance evaluation
 - communication (visualisation & interpretation)

candidate data:

- ABX treatment (UU dataset)
- environmental example (e.g. TARA Oceans)
- IBD data human health outcome

Method of analysis: print("In R we trust")

Method of delivery (independent of language): HTML; RMD v. IPYNB

Duration: 3x (2 hours theory / 4 hours practical)

- 1. ML approaches for the microbiome
- 2. Preparing microbiome for ML
- 3. Applying ML to microbiome
- 4. Evaluating and communicating outcomes

Zoning allows rearrangement / alternate modes of delivery

outcomes



scope : range of requirements feasible within short teaching timeframe

framework : foundational set of connected topics from ML4Microbiome's outputs, to allow informed implementation of ML

practical exercises : applications of framework, using publicly available data

mode : allows for rapid sharing/dissemination, as well as future improvements /

personal: better understanding of machine learning, better appreciation for complexity and role in microbiome analysis



Takeaway: Check COST.eu, apply for your own STSM!

Get funded

- \rightarrow to go somewhere you want
 - \rightarrow to talk to your peers
 - \rightarrow about something you're interested in
 - \rightarrow and then put it on your CV

Applying to EU COST Action:

- ✓ **<u>COST.eu</u>** straightforward portal. Many, many Actions, on all topics
- ✓ 2 straightforward documents asks you to understand what you are applying for, and to structure your application accordingly.
- ✓ COST staff / administration very helpful.
- \checkmark Excellent experience in starting applications for funding



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Recommendations:

- Do serious work, apply for serious funding
- Reporting and submitting claims very straightforward: use report template as a diary
- Getting the most out of your trip plan ahead, clear your schedule!

With many *thank* you's to:





Claesson group



Vision 1 group

Roosmarijn Luiken & Aldert Zomer

ML4Microbiome, esp. workshop authors:

Matti Ruuskanen Dimitrios Vlachakis Aleksandra Gruca

& Marcus Claesson

EU-COST admin: Aoife Tierney Tatjana Turukalo Anna Power



Utrecht University







MICROBIOME



