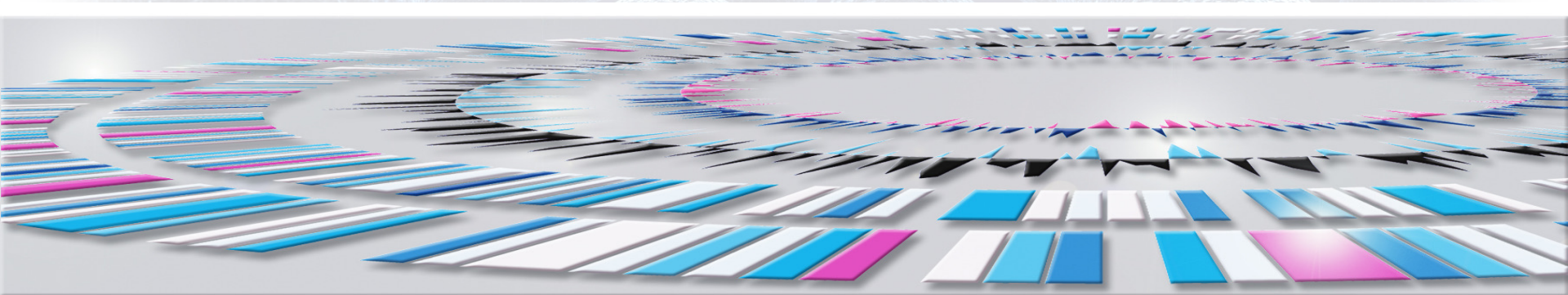



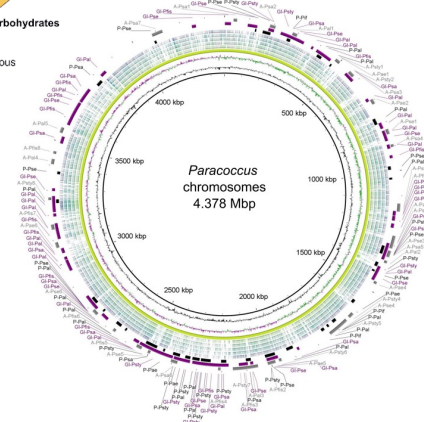
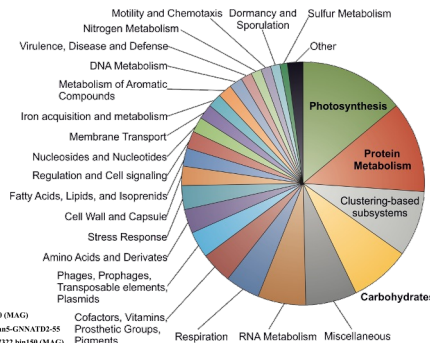
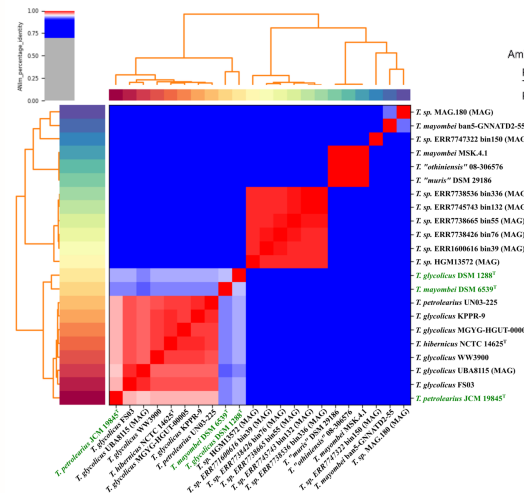
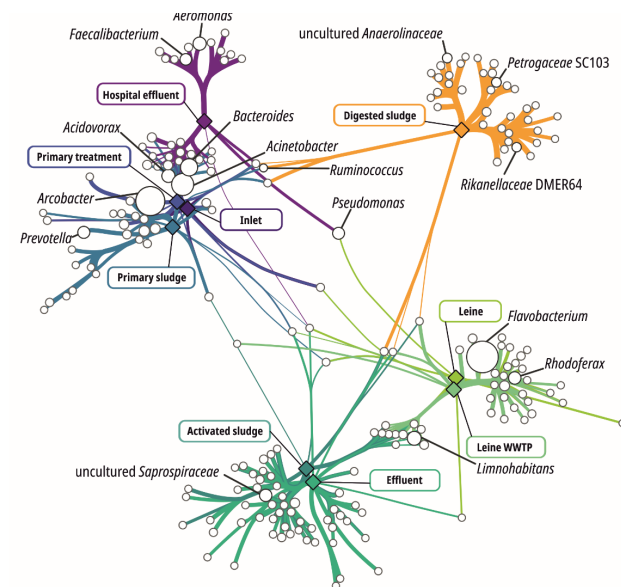
M.Bio.105

# Applied bioinformatics in molecular biosciences



Rolf Daniel, Heiko Liesegang, Department of Genomic and Applied Microbiology  
[rdaniel@gwdg.de](mailto:rdaniel@gwdg.de) or [hlieseg@gwdg.de](mailto:hlieseg@gwdg.de)

- M.Bio.105 addresses available software and databases of – omics driven biological data analysis
  - Application of bioinformatic methods to investigate:
    - **Molecular phylogeny and sequence-based evolution**
    - **Genome dynamics and Meta(omics)**
    - **RNA and protein analysis**
    - **Identification and modelling of protein classes like enzymes and toxins**
    - **Construction of metabolic networks**
- 



# Module M.Bio.105

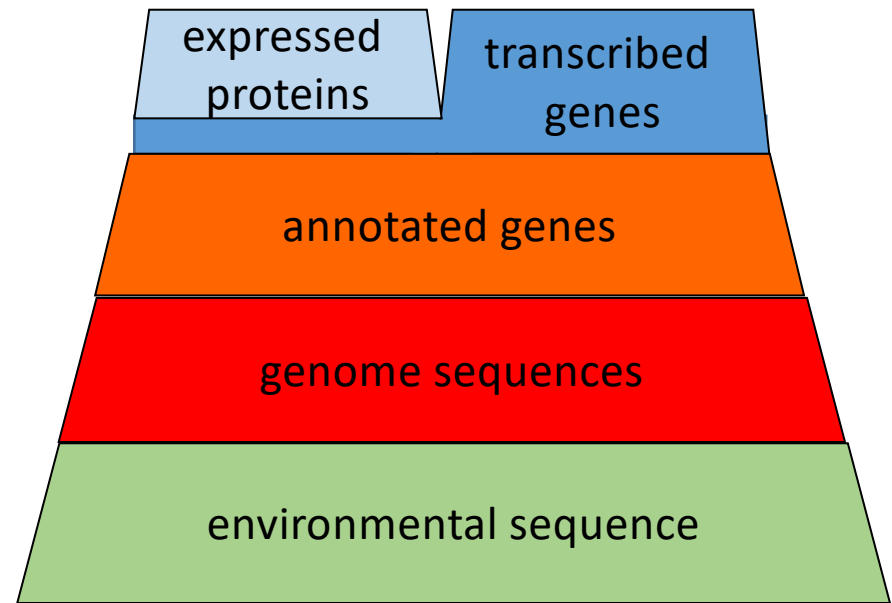
Core of the module is:

analysis, visualization and the integration of large data sets from – omics technologies.

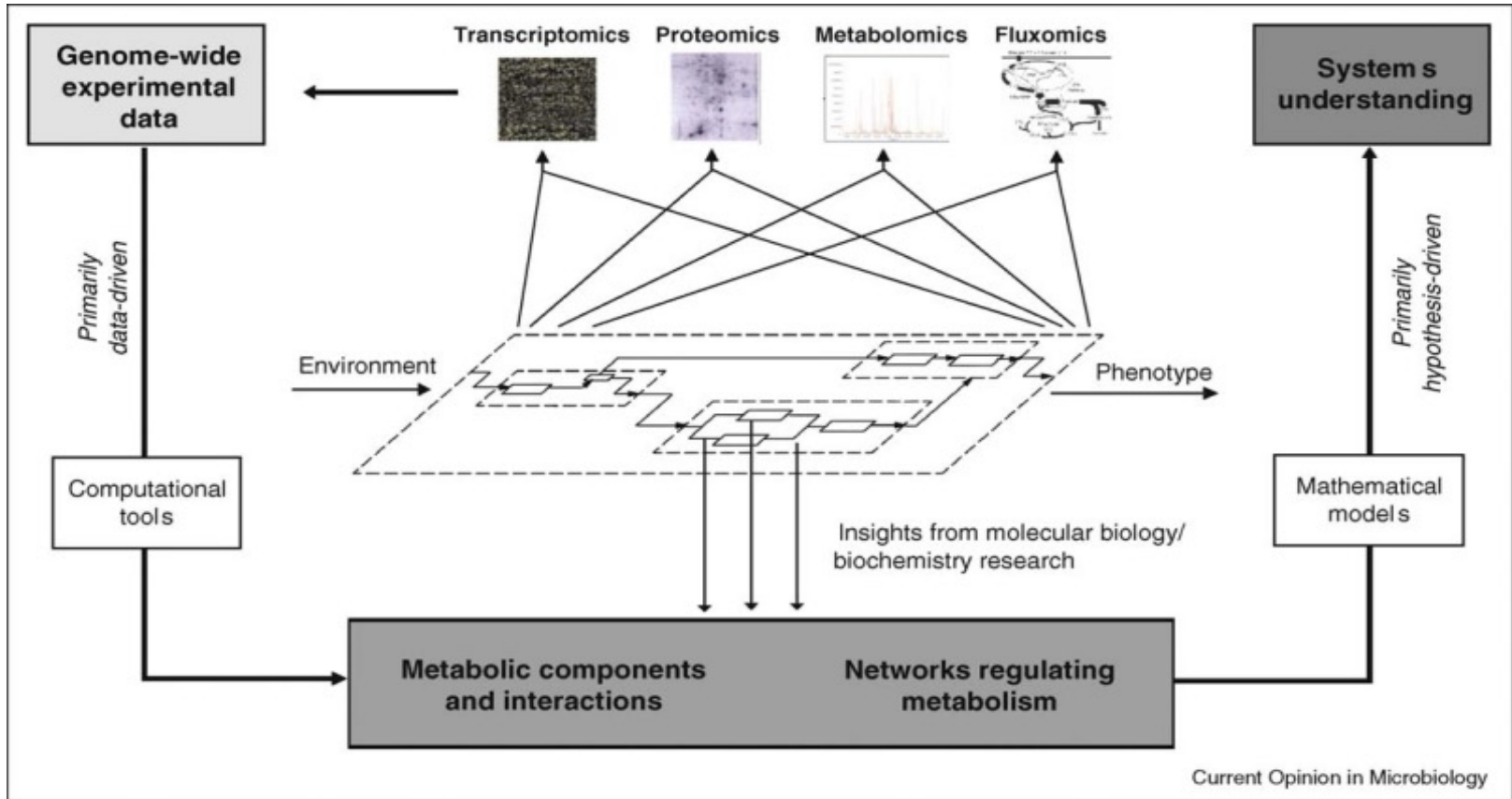
Goal: to acquire the data analysis abilities for the understanding of systems biology from single organisms to entire environmental communities.

# -omics data describes the entire set of a biological entity:

- **Metagenomics:**  
all organisms present in a sample
- **Genomics:**
  - Replicons in a genome
  - genes (protein and sRNA) present in organisms
- **Transcriptomics:**
  - All genes active at a given condition
- **Proteomics:**
  - All proteins expressed at a given condition



# -omics data and systems biology



# M.Bio.105: summary

- Organized in:
  - Lectures (3 SWS)
  - Literature seminars (1 SWS)
  - Practical exercises (10 SWS)
  - Limited to 10 students
- Uses real data from current projects
- Bioinformatic background
  - Linux skills
  - B.Bio.117 genome analysis or equivalent
  - Recommended: Python- and R- scripting
  - Fits to Computational Biology (M.CoBi.501)
- Scheduled each winter term
- Language: English
- Exam requirements: knowledge on how to apply bioinformatic methods focused on (Meta)omics data. Ability to present and understand scientific literature
- Exam: oral presentation on methods and results applied on –omics data (80%) plus seminar talk (20%)
- Workload: 5 weeks block course, start 19.1.2026; 360h (196 presence/164 self-study)
- Supervisors: Prof. Jan de Vries, Dr. Oliver Valerius, PD Dr. Heiko Liesegang, Dr. Anja Poehlein, Prof. Rolf Daniel, Dr. Ines Friedrich

This is a module designed also  
as preparation for data-driven lab  
rotations or master thesis projects

In case of questions:

**rdaniel@gwdg.de**

**hlieseg@gwdg.de**