

Course KB8019 Comparative Genomics, 7.5 hp

Preliminary schedule for 2023, version 3/23/23. *Do not make a copy of this file* as it may change.

Hosted by Stockholm University, DBB.

Course responsible: Prof. Erik Sonnhammer, erik.sonnhammer@scilifelab.se

Course goals: to learn current techniques for analysing genomes and how comparative genomics can be used to understand the organisation, evolution, and function of genomic sequences.

Course literature:

- Web resources.
- Zvelebil and Baum, [Understanding bioinformatics](#). Not as up to date as the web resources but has more in-depth explanations of many concepts and algorithms.

Course begin/end: May 2 – June 2 2022

- Classes at Science for Life Lab, in Air&Fire on Gamma 2.
- The listed literature should be read before each class. Time is reserved for this in the morning of the class day.
- Practicals are done via remote login on a server or locally on your own computer (not recommended). Accounts on server will be provided at the course start. Teacher assistants will be available in the Gamma 7 lunch room on non-class days 13-16.
- Reports for practicals should be submitted during the week they are listed, but at the latest the Monday after. Practicals are graded passed / not passed.
- Slides will be provided before the class.
- Teacher assistants: Davide Buzzao and Emma Persson
- Course information on [Google drive](#)
- Course information and reporting via Athena; log in with your SU account on <https://athena.itslearning.com>

Week 1. The structure of prokaryotic and eukaryotic genomes; Gene prediction

May 2, 10.00-12.00: Roll call, Introduction to course, and start of practicals

Practical 1: Basic genome analysis. Briefing May 2, ~11.00

Practical 2: Gene prediction. Briefing May 2, ~11.00

May 3, 14-16:

Class 1. Genome organisation

Intended Learning Objectives:

- Introduction to genomes
- Genome properties in different species
- Gene content of genomes
- Regulatory sequences
- Non-coding sequences
- Metagenomics

Class 2. Gene prediction

Intended Learning Objectives:

- Gene structure in different species
- Gene prediction in different species
- Promoter prediction in different species

Literature:

https://en.wikipedia.org/wiki/List_of_biological_databases

[https://en.wikipedia.org/wiki/Domain_\(biology\)](https://en.wikipedia.org/wiki/Domain_(biology))

<https://en.wikipedia.org/wiki/Genome>

<https://en.wikipedia.org/wiki/C-value>

[https://en.wikipedia.org/wiki/Translation_\(biology\)](https://en.wikipedia.org/wiki/Translation_(biology))

https://en.wikipedia.org/wiki/Genome_evolution

https://en.wikipedia.org/wiki/Gene_prediction

<https://www.ncbi.nlm.nih.gov/blast/tutorial/>

Optional: https://en.wikipedia.org/wiki/Introduction_to_genetics

Optional: https://en.wikipedia.org/wiki/Human_genome

Optional: <https://en.wikipedia.org/wiki/Bioinformatics>

Optional: https://en.wikipedia.org/wiki/Genetic_code

Optional: https://en.wikipedia.org/wiki/Non-coding_DNA

Zvelebil:

Chapter 3 Dealing with Databases

Chapter 9 Revealing Genome Features

Chapter 10 Gene Detection and Genome Annotation

Week 2. Evolution of genes and genomes

May 8, 14-16:

Class 3. Phylogenetics

Intended Learning Objectives:

- Understand principles of phylogenetic trees
- Understand methods of phylogenetic tree building
- Understand methods to assess the quality of phylogenetic trees

Class 4. Phylogenomics

Intended Learning Objectives:

- Understand principles of phylogenomics
- Understand applications of phylogenomics
- Understand phylogenetic profiles, metaphylogeny, and tree of life
- Understand reasons for phylogenetic disagreement

Practical 3: Phylogenetic reconstruction. Briefing May 8, 16.00

Practical 4: Phylogenomics. Briefing May 8, 16.00

Literature:

https://en.wikipedia.org/wiki/Phylogenetic_tree

https://en.wikipedia.org/wiki/Substitution_model

<https://en.wikipedia.org/wiki/UPGMA>

https://en.wikipedia.org/wiki/Neighbor_joining

<https://www.nature.com/articles/nrg1603.pdf>, boxes 1 and 2 (Delsuc et al., 2005)

<https://genome.cshlp.org/content/8/3/163.long> (Eisen, 1998)

https://en.wikipedia.org/wiki/Phylogenetic_profiling

https://en.wikipedia.org/wiki/Phylogenetic_network

Optional: https://en.wikipedia.org/wiki/Models_of_DNA_evolution

Optional: https://en.wikipedia.org/wiki/List_of_phylogenetics_software

Optional: https://en.wikipedia.org/wiki/Phylogenetic_tree_viewers

Optional: [https://en.wikipedia.org/wiki/Bootstrapping_\(statistics\)](https://en.wikipedia.org/wiki/Bootstrapping_(statistics))

Optional: <https://en.wikipedia.org/wiki/Phylogenomics>

Optional: <https://en.wikipedia.org/wiki/Phylogenetics>

Zvelebil:

Chapter 7: Recovering Evolutionary History

Chapter 8: Building Phylogenetic Trees

Week 3. Synteny and orthology analysis

May 15, 14-16:

Class 5. Gene order

Intended Learning Objectives:

- Introduction to gene order conservation (synteny)
- Analysing synteny with dotplots
- Using synteny for inferring phylogeny

Class 6. Orthology

Intended Learning Objectives:

- Definitions & assumptions
- Applications
- How to identify orthologs
- Tree reconciliation
- Graph-based methods; Tree-based methods; Other methods
- Comparison of ortholog databases

Practical 5: Gene order analysis. Briefing May 15, 16.00

Practical 6: Orthology. Briefing May 15, 16.00

Final project assignment: Briefing May 15, 16.00

Literature:

<https://www.ncbi.nlm.nih.gov/pubmed/31278664> (Altenhoff et al., 2019)

<https://inparanoid.sbc.su.se/cgi-bin/faq.cgi>

<https://en.wikipedia.org/wiki/Synteny>

[https://genomevolution.org/wiki/index.php/Synteny: Getting the Big Picture](https://genomevolution.org/wiki/index.php/Synteny:_Getting_the_Big_Picture)

https://en.wikipedia.org/wiki/Sequence_homology

[https://en.wikipedia.org/wiki/Dot_plot_\(bioinformatics\)](https://en.wikipedia.org/wiki/Dot_plot_(bioinformatics))

<https://www.ncbi.nlm.nih.gov/pubmed/11934753> (Tesler, 2002)

Optional: <https://questfororthologs.org/>

Optional: <https://orthology.benchmarkservice.org/>

Zvelebil:

Chapter 7.2 Molecular Evolution and its Consequences

Week 4. Interaction networks

May 22, 14.00-16.00:

Class 7. Interaction networks

Intended Learning Objectives:

- Introduction to network biology
- Discovering protein interactions
- Network properties
- Network databases (STRING, FunCoup)

Class 8. Applications of network analysis

Intended Learning Objectives:

- Network analysis for gene identification

- Pathway analysis
- Gene regulatory networks

Practical 7: Interaction networks. Briefing May 22, 16.00

Literature:

https://en.wikipedia.org/wiki/Biological_network

<https://en.wikipedia.org/wiki/Interactome>

https://en.wikipedia.org/wiki/Network_science

https://en.wikipedia.org/wiki/Scale-free_network

https://en.wikipedia.org/wiki/Clustering_coefficient

https://en.wikipedia.org/wiki/Systems_biology

[The Architecture of Biological Networks \(Wuchty et al., 2006\)](#)

[FunCoup 5: Functional Association Networks... \(Persson et al., 2021\)](#)

https://en.wikipedia.org/wiki/Pathway_analysis

Optional: <https://www.ncbi.nlm.nih.gov/pubmed/27151197> (Ogris et al., 2016)

Zvelebil:

Chapter 17: Systems Biology

Week 5. Final project assignment

(Briefing in week 3)

June 2, 10.00-13.00, Air&Fire Gamma 2: exam