"Summer" course 2024



Monday 19 Aug.	2		*	١	22°/15°
Tuesday 20 Aug.	2	*	*	2	24°/15°
Wednesday 21 Aug.	*		*	2	21°/15°
Thursday 22 Aug.	2				22°/15°
Friday 23 Aug.		*	4		24°/19°

Summer course in analysis of high throughput sequencing data for population genetics

Organizers/Lecturers



Ida Moltke BIO, UCPH



Anders Albrechtsen BIO, UCPH



Fernando Racimo GLOBE, UCPH

Martin Sikora, GLOBE, UCPH

Garrett Hellenthal, UCL

Cindy Santander BIO, UCPH

Shyam Gopalakrishnan GLOBE, UCPH

Co-organizers and TAs



Sabhrina Aninta



Shixu He



Renzo Balboa Xiaodong Liu



Zilong Li



Carlos Sarabia

Goal of this course

- Comprehensive introduction to a number of topics and common research tools used in analyses of high throughput genetic data for population genetics
- Go beyond the textbooks
 - Use of cutting edge technique and approaches for NGS data analysis
 - Give hands-on experience with complex analysis and NGS data

Networking

Intended learning outcomes

After the course you should be able to

- use population genetic theory to infer basic population general characteristics
- use NGS data incl. low depth for population genetic inference
 - population structure
 - admixture
 - selection
 - demography
- Interpret and discuss the results of own analyses and results the scientific literature

Each day

- 2 teaching sessions
 - Morning (9:00 12:00) & afternoon (13:00 16:00)
 - Mixture of lectures and exercises
- Lunch (on your own)
 - -12:00-01:00
- Research talk
 - 16:15-17:00 (Monday 15:30)
- Social session
 - 17:00-18:00 (Monday 16:00)

Monday

Introduction to population genetics and NGS

Morning session:

Introduction to basic population genetic terms and concepts (Fernando)



Introduction to NGS data (Anders)

Research lecture:

Patrícia Pečnerová, BIO, UCPH Using genomics to study the fine line between extinction and survival in mammoths and elephants

Social:

Reception/social mixer







Tuesday

Analysis of NGS data and population structure

Morning session:

Estimation of allele frequencies, SNP calling and genotype calling, from NGS data (Anders)

Afternoon session:

Inference of population structure I (Ida)

Research lecture:

Rasmus Heller, BIO, UCPH Genomes from the "Asian unicorn" reveal a long-standing decline and two highly differentiated populations of saola

Social:

Boat trip (sightseeing :-))



Wednesday

Population structure and gene flow

- Morning session:
 PCA in population genetics (Anders)
- Afternoon session:
 D/f/F statistics (Martin)

Research lecture:

Martin Sikora, GLOBE, UCPH
Using population-scale ancient genomics to study ancestry,
social organisation and disease in Late Neolithic Scandinavia



Social:

Relaxed gathering

Thursday

Admixture graphs and selection

- Morning session:
 Finestructure (Garrett)
- Afternoon session:
 Detecting genomic regions under positive selection (Cindy)

- Research lecture:
 Kristine Bohmann,GLOBE, UCPH
 Vacuuming animal DNA from thin air
- Social: Relaxed gathering



Friday

Selection and demography

- Morning session:
 Dating selection and admixture events (Garrett)
- Afternoon session:
 Demographic inference (Shyam)
- Research lecture:
 Garrett Hellenthal, UCL
 Inferring overlapping admixture events among human populations
- Social: Relaxed gathering and farewell



Getting connected

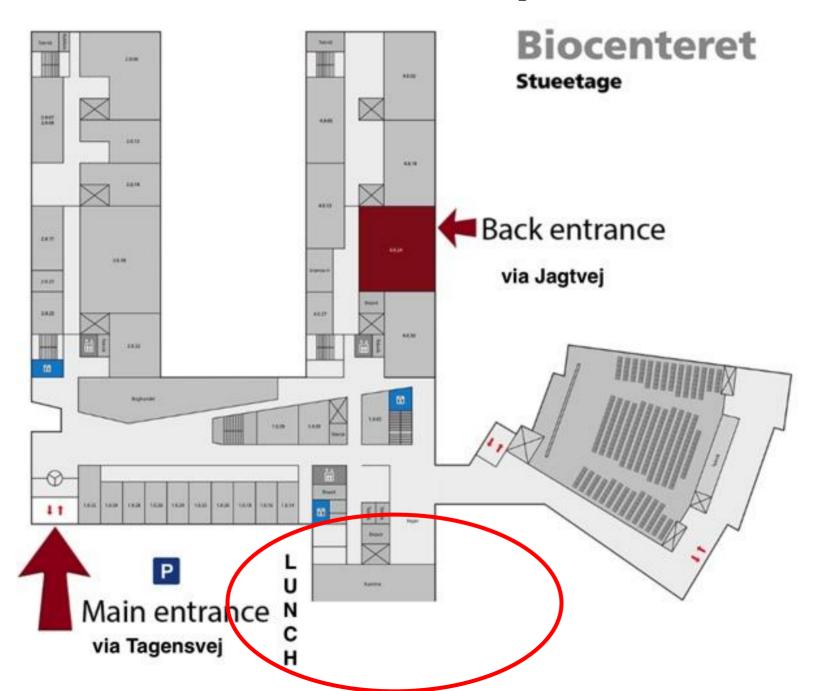
Wifi

- Eduroam
 You can connect via Eduroam (if you have an account)
- KU guest
 Create a guest account with your mobile number

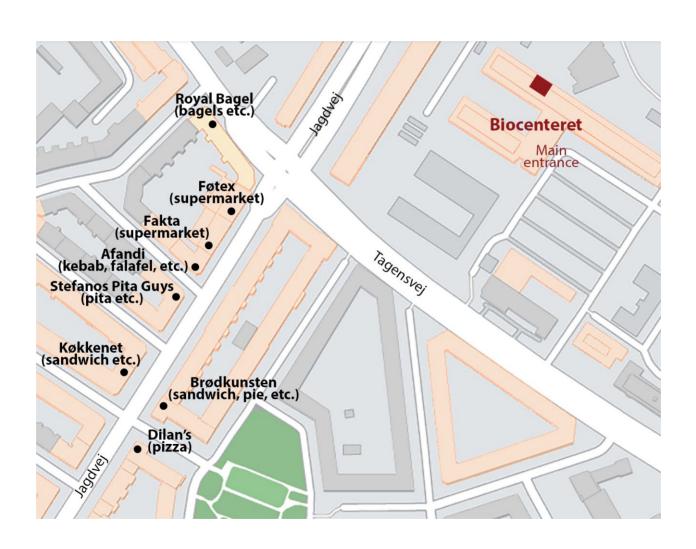
Login to exercises

We will distribute usernames and passwords

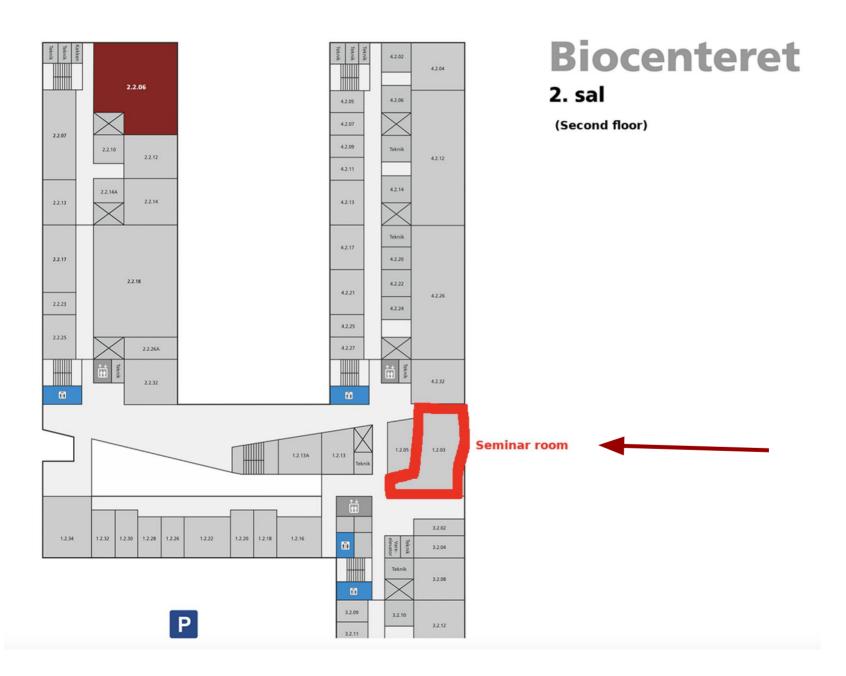
Lunch options I



Lunch options II



Afternoon research lectures



While Fernando gets ready

I'll pass around 2 sheets:

- 1) One where you have to confirm that you are present
- 2) One where you can sign up if you want to join the boat trip

And while those are passed around we'd like to do a really quick round of introduction of you (participants from 15 different countries!):

