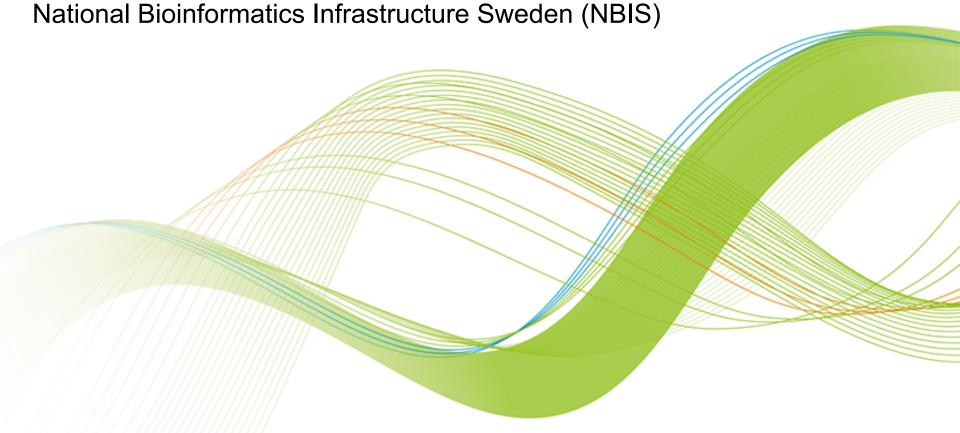


Malin Larsson

SciLifeLab long term bioinformatics support





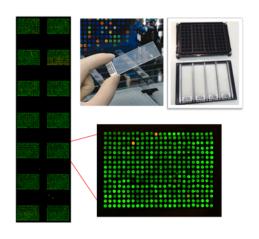
Develops, offers and applies advanced technologies for molecular biosciences with a focus on health and environment



- Joint Uppsala Stockholm center with two nodes
- Hosted by four universities
- Official start, 1 July 2013
- Approximately 1500 researchers by mid 2014
- Infrastructure for molecular bioscience

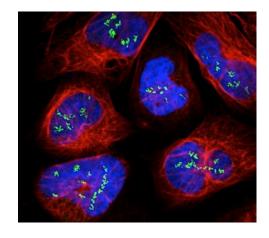
SciLifeLab mission





Technology platforms for national use

- Service
- Technology Development
- Analytical support
- Courses and workshops



Generate a strong research environment

- Affiliated faculty
- SciLifeLab Fellows Program
- National projects
- International collaborations

SciLifeLab platforms

SciLifeLab

VR

National
Genomics
Infrastructure

Joakim Lundeberg Ann-Christine Syvänen Ulf Gyllensten National
Bioinformatics
Infrastructure
Sweden

Bengt Persson

Clinical Diagnostics

Lars Engstrand

! SNIC



Computer resources free for Swedish researchers









Ongoing merge of BILS, WABI and more; complete 2016.
National, distributed













Upcoming courses



Date	Course
April 12-14	RNA-seq
April 26-27	Introduction to Genome Annotation
May 23-27 NOTE! Application deadline April 18	Perl Programming with Application to Bioinformatics
October 10-14	Perl Programming with Application to Bioinformatics
October 25-27	RNA-seq
November 15-16	De novo Genome Assembly
November 22-23	Metagenomics
Application Nov/Dec	The Swedish Bioinformatics Advisory Program











www.scilifelab.se



Course schedule

Monday

Room: Campus Valla, House B, Room Odén

09.00-09.15 Registration and Welcome, presentation of SciLifeLab

09.15-10.00 Lecture: Introduction to Linux (Martin Dahlö)

10.00-12.00 Exercise: Introduction to Linux (Martin Dahlö, Malin Larsson)

12.00-13.00 Lunch

13.00-14.00 Lecture: Introduction to UPPMAX/UPPNEX (Martin Dahlö)

14.00-17.00 Exercise: Introduction to UPPMAX/UPPNEX (Martin Dahlö, Malin Larsson)

Tuesday

Room: Campus Valla, House B, Room Odén

09.00-10.00 Lecture: Alignment with BWA; Data Processing with Picard; Variant Calling with GATK; SAM/BAM and VCF Formats

10.00-12.00

Exercise: Alignment with BWA; Data Processing with Picard; Variant Calling with GATK; SAM/BAM and VCF Formats part I (Malin Larsson, Mihaela

Martis)

12.00-13.00 Lunch

13:00-17:00 Exercise: Alignment with BWA; Data Processing with Picard; Variant Calling with GATK; SAM/BAM and VCF Formats part II

Wednesday

Room: Campus Valla, House B, Room Odén

09.00-11.30 Lecture: Next Generation Sequencing Overview

11:30-12.00 Lecture: The National Bioinformatics Infrastructure Sweden

12.00-13.00 Lunch

13.00-14.00 Lecture: NGS and bioinformatics analysis pipelines

14.00-17.00 Project discussion

18:00- Course dinner at De Klomp

<u>Gr</u>

Thursday

Room: Campus Valla, House B, Room Odén

09:00-10.00 Lecture: ChIP sequencing and data analysis

10.00-12:00 Exercise: ChIP sequencing and data analysis part I

12.00-13.00 Lunch

13.00-17:00 Exercise: ChIP sequencing and data analysis part II

Friday

Room: Campus Valla, House B, Room Odén

09:00-10:00 Lecture: RNA sequencing, transcriptome and expression quantification

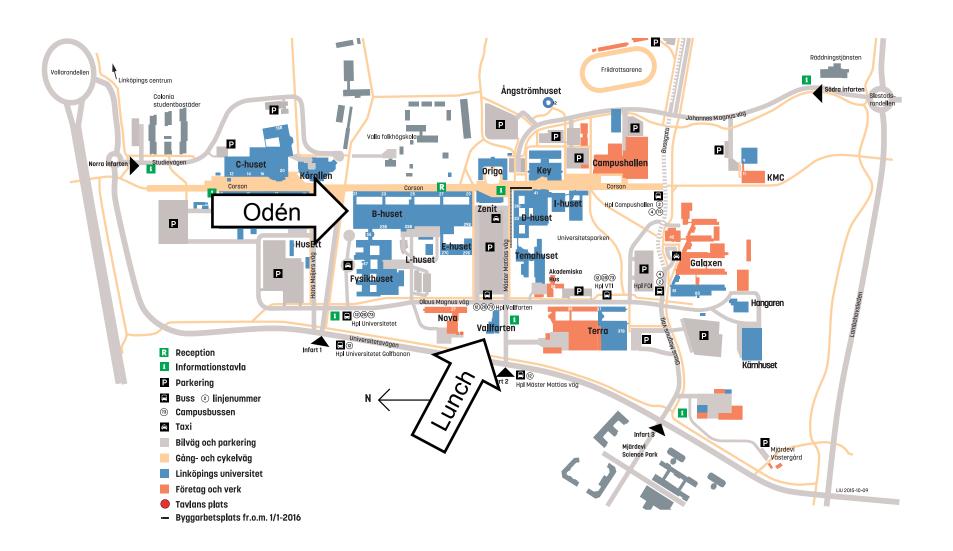
10:00-12:00 Exercise: RNA sequencing, transcriptome and expression quantification part I

12.00-13.00 Lunch

13:00-17.00 Exercise: RNA sequencing, transcriptome and expression quantification part II

Project discussions in smaller groups

Lunch in Universitetsklubben



Project discussions on Wednesday

Groups based on your interest. Proposed groups:

- RNA-seq and epigenetics (Agata and Mihaela)
- Variant calling including somatic variants (Malin and Adam)
- Experimental setup (Olga)

Please plan questions for the discussion:

- technology-related
- analysis-related
- if there is something your would like to discuss 1:1 with a teacher only
- ?

Time: 14:00 -17:00 on Wednesday

Place: Odén, Fick and Dalton

Groups will be decided during Monday/Tuesday (let me know which group you want to be in)