Epigenomics Data Analysis

Short course

August, 26th - 30th, 2019

Stockholm, SciLifeLab

National course open for PhD students, postdocs, researchers and
other employees within Swedish universities.

https://www.scilifelab.se/events/epigenomics-
data-analysis/

Best practices for data processing and
analysis for ChIP-seq, ATAC-seq,
Illumina EPIC arrays, Bisulfite-seq.
Quality control. Peak calling. Motif
enrichment. Differential binding,
accessibility and methylation. Functional
analyses. RNA-seq integration.
Integrative visualizations.

NBIS training events
Up-coming events

RNA-seq data analysis
starts May, 13th, Uppsala

Intro. to Bioinformatics using NGS data
starts May, 20th, Göteborg

Biostatistics Essentials
starts May, 21st, Uppsala

Data integration
starts September, 9th, Stockholm

Bioinformatics for Principal Investigators
starts September, 17th, Stockholm

Python programming
starts September, 23th, Uppsala

More events & information:
https://nbis.se/training/events.html