



National Bioinformatics Infrastructure Sweden NBIS

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NBIS - The SciLifeLab Bioinformatics Platform

nbis.se

many different omics-related areas













Overview of NBIS units

https://nbis.se/













Drop-in sessions

Ask your question! Bioinformatics, sequencing, data management, and more Online - Tuesdays at 14.00







Study Design & project consultation	3 h per project No hands-on work	
Short or medium term support (Fee-for-service)	User fee 800 SEK/h * All projects welcome, queuing system	
Long-term support (KAW funded)	500 h support over ~ 1-2 years, for free Scientific selection by committee, 3 application rounds/year Involvement from research group mandatory	a second
Partner Projects	For projects with large bioinformatic component "Hire" an NBIS bioinformatician Full-cost model: utilize and pay for e.g. 50% of one NBIS FTE	

*The BioImage Informatics Facility has a slightly different financing policy















Infrastructure, Data management, Tools 🌖 SciLifeLab

Infrastructure development

- International repositories (e.g. Federated EGA)
- Hosting/interoperability of national resources of major interest (e.g. Human Protein Atlas, SweFreq, Metabolic Atlas)

Data management (close collaboration with Data Centre)

- Representing Sweden in international data initiatives
- Data Stewards to guide users in data management and FAIR data publishing

Tools

- Maintaining a few selected major Swedish software
- Continuous development of software, workflows and services













- Data stewards support researchers
- Collaboration with SciLifeLab Data Centre (supports platforms)







About 6 staff members

A bridge between NBIS and SNIC-funded UPPMAX

- SNIC free to use national infrastructure for computing, funded by Swedish Research Council
- UPPMAX Uppsala high-performance computing centre
- Provides a digital research environment serving 700+ PIs with over 1000 projects

Focus on HPC systems provided by UPPMAX



NBIS coordinates the Swedish node in ELIXIR – SciLifeLab Stream Eleveretics the European infrastructure for biological information

22 members + 1 observer

Five technical platforms for

Compute Data Tools Interoperability and Training

Complemented by **Communities** in

Marine metagenomics

Rare diseases

Human data

Plants sciences

Proteomics

Metabolomics

Galaxy and more to come

Swedish contributions: -- Human Protein Atlas

- -- Human Data Community
- -- Federated EGA and Beacons
- -- Data Management
- -- Biodiversity
- -- Training (Jessica Lindvall Ex-Co)
- Bengt Persson, Head of Node
- Jessica Lindvall, Deputy Head of Node and Training Coordinator
- Johan Viklund, Technical Coordinator
- Jessica Lindberg, Node Coordinator

At the global level interactions with: RDA, NIH BD2K, GA4GH













Training events and programs

- Courses, teacher-dense for good networking
- Mentor program, boosting PhD skills

Rapid knowledge-transfer

- Drop-in sessions (Currently online, Tuesdays at 14:00)
- Catalyse rapid knowledge-transfer across projects/research areas
- Portable tools and workflows to empower users







Training opportunities:

https://uppsala.instructure.com/courses/48087/pages/nbis-training-catalogue

Example workshop topics:

- Python
- Data visualisation in R
- Biostatistics
- Workflows (snakemake) Bring your own code
- Reproducible research
- RNASeq & scRNASeq
- Epigenomics





• Last minute registration for the 3 day RNASeq course in Uppsala

Application deadline: February 12, 2023

https://nbis.se/training/events.html https://www.scilifelab.se/event/rna-seq-data-analysis/

- RNAseq quality control analysis
- Mapping of short reads
- Quantification of gene expression
- Exploratory data analysis
- Differential gene expression
- Gene set enrichment analysis







Any questions?

