



National Bioinformatics Infrastructure Sweden NBIS

Dag Ahrén 21 May 2021





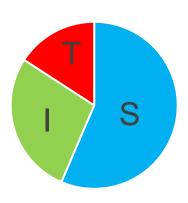
NBIS – SciLifeLab Bioinformatics Platform



Enable world-class life science research and maximise scientific and societal impact of collected data by:

- Providing expert knowledge, innovative data integration, advanced training, efficient data publication for open science, and access to high-performance data analysis methods
- Coordinating bioinformatics support within Sweden and making bioinformatics easily accessible for life science researchers
- Swiftly responding to changes in support needs as new techniques are developed and utilised
- Forming the Swedish ELIXIR node and participating in relevant international projects





Analysis of biological data

Support

- Genomics/NGS/Metagenomics
 - Genome annotation and assembly
- MS-Proteomics and Protein bioinformatics
- Systems biology and Metabolomics
- Integrative bioinformatics
- Biostatistics
- Data publication and FAIRification of data
- Data management and Data stewardship (collaboration with SciLifeLab Data Centre) https://nbis.se/support

Infrastructure

- services, computational resources, data management, tools and guidelines
- https://nbis.se/infrastructure

Training

- https://nbis.se/training
- The Swedish node in ELIXIR -- the European infrastructure for biological information



Utiftelse

Funding from:























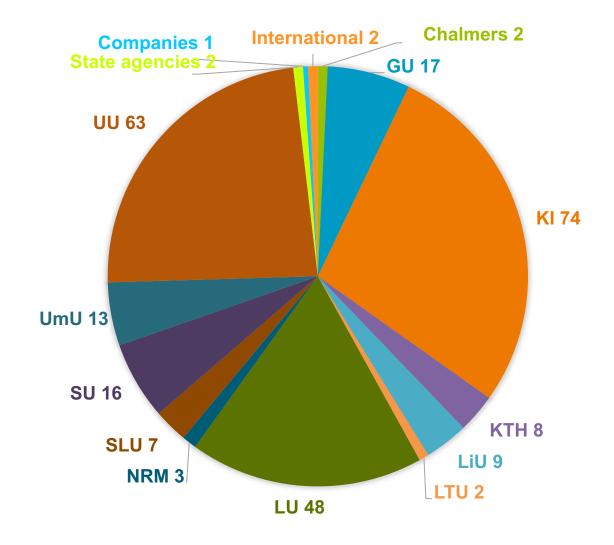




Distribution of support during 2020



- **267 Pls** (104 female, 163 male)
- 212 active support projects
- 104 study design consultations
- Several advanced training courses







Overview of NBIS units

https://nbis.se/





Four support tracks



Study design consultation

- provided for free (up to 3 hours)
- help our users to order the right type of data for their analysis
- consider bioinformatics and data management issues at an early stage
- extra drop-in sessions at VR grant application periods

Short- & Medium-term support

- academic user fee 800 SEK (~80 EUR) per hour
- short waiting times
- 50% support between universities, most suitable expert

Long-term support

- KAW funding, up to 500 hours for free
- scientific evaluation; application rounds 3 times per year

Partner Projects (PP)

- intended for projects with a large bioinformatics component
- NBIS support ≥12 person months over the project life-time, running 2–5 years calendar time
- user fee to cover staff costs





Training: A key strategic area



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





Infrastructure, Data management, Tools



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

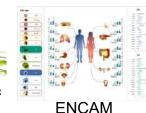
- Maintaining a few selected major Swedish software
- Continuous development of software, workflows and services













Compute & Storage



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 – 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



Thank you!!



Any questions?

