

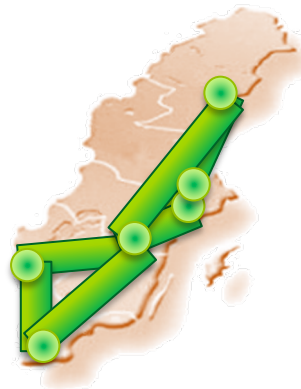
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# National Bioinformatics Infrastructure Sweden NBIS

**Dag Ahrén**  
**26 November 2021**

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



Distributed infrastructure with nodes at each of the 6 large university towns and totally ~100 staff



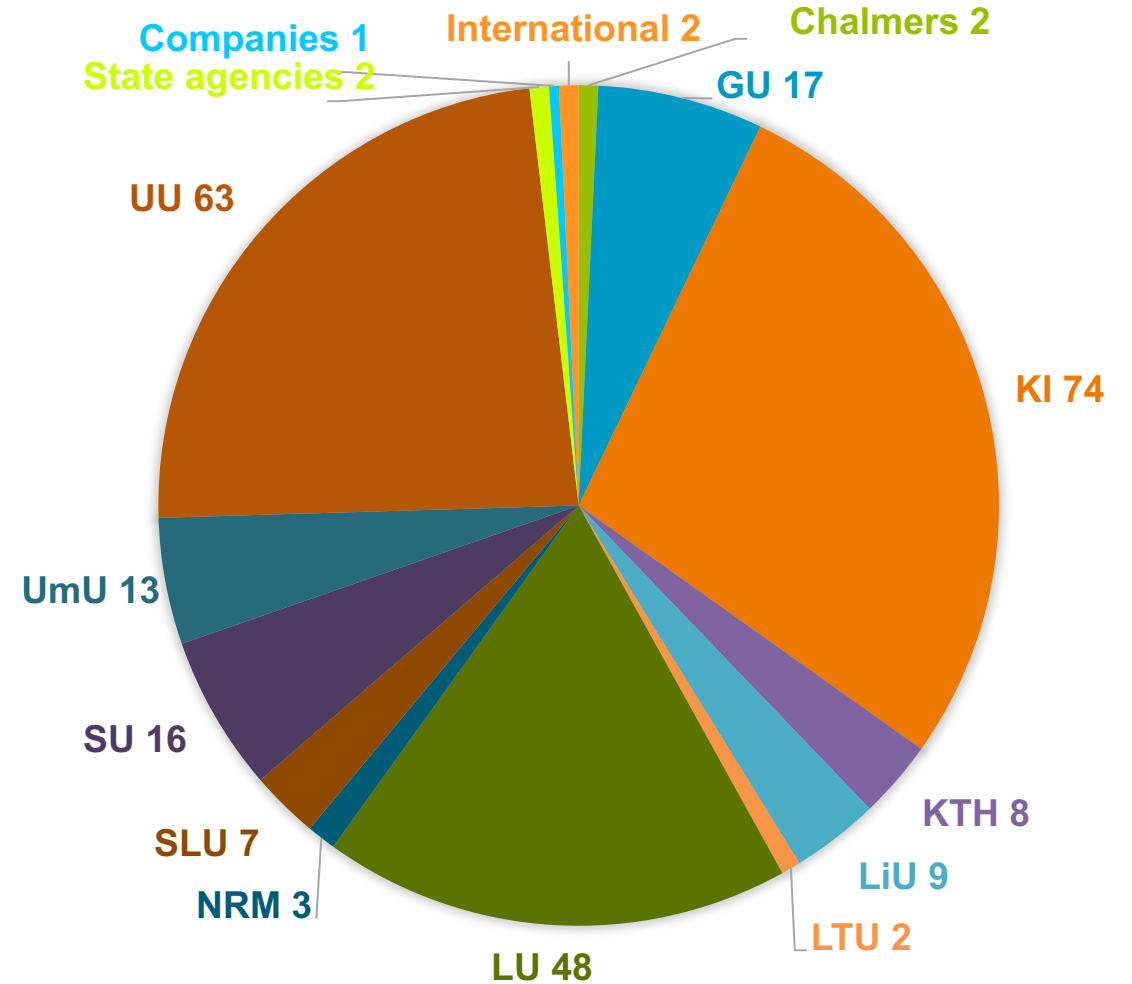
## Analysis of biological data

- **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

Funding from:



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



# Overview of NBIS units

<https://nbis.se/>



The screenshot shows the NBIS website homepage. At the top left is the NBIS logo. The top right navigation menu includes: Support, Infrastructure, Training, News, Events, About, and the SciLifeLab logo. The main banner features the large NBIS logo with the text 'NATIONAL BIOINFORMATICS INFRASTRUCTURE SWEDEN' below it. A dark blue banner below the logo contains the text: 'NBIS is a distributed national bioinformatics infrastructure, supporting life sciences in Sweden'. Below this is a red banner with the text: 'See [here](#) for information on how our services are affected by the [covid19](#) outbreak'. At the bottom, there are three green boxes representing the main service areas: Support, Infrastructure, and Training.

**NBIS**  
NATIONAL BIOINFORMATICS  
INFRASTRUCTURE SWEDEN

NBIS is a distributed national bioinformatics infrastructure, supporting life sciences in Sweden

See [here](#) for information on how our services are affected by the [covid19](#) outbreak

**Support**  
Support services ranging from short consultation, consultancy to long-term embedded bioinformaticians.

**Infrastructure**  
Providing infrastructure in the form of services, computational resources, tools and guidelines to the life science community.

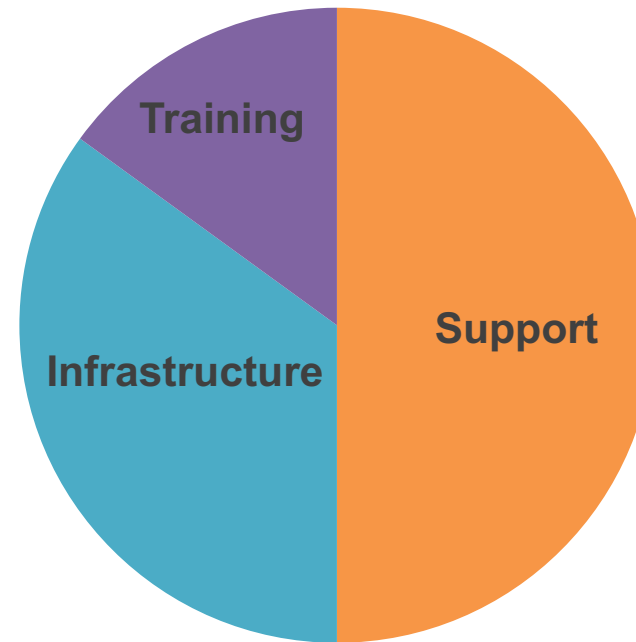
**Training**  
Training events in advanced and applied bioinformatics.



20 national courses  
5 international courses  
625 students  
30 PhD:s in mentor program



Development for sensitive data  
Coordination of e.g. 1M Genomes Initiative  
Guidance to FAIR data  
Maintaining selected Swedish tools



*Resources 2019*



245 unique PI:s  
170 active support projects  
100 new projects  
70 publications  
120 booked consultations  
250 drop-in sessions

## Training events and programs

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

[Advisory Programme open now! Deadline 22<sup>nd</sup> of December](#)

## Rapid knowledge-transfer

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



- **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  $\geq 12$  person months over the project life-time, running 2–5 years calendar time
  - user fee to cover staff costs



## 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 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



**Mr Bayes**  
Bayesian Inference of Phylogeny



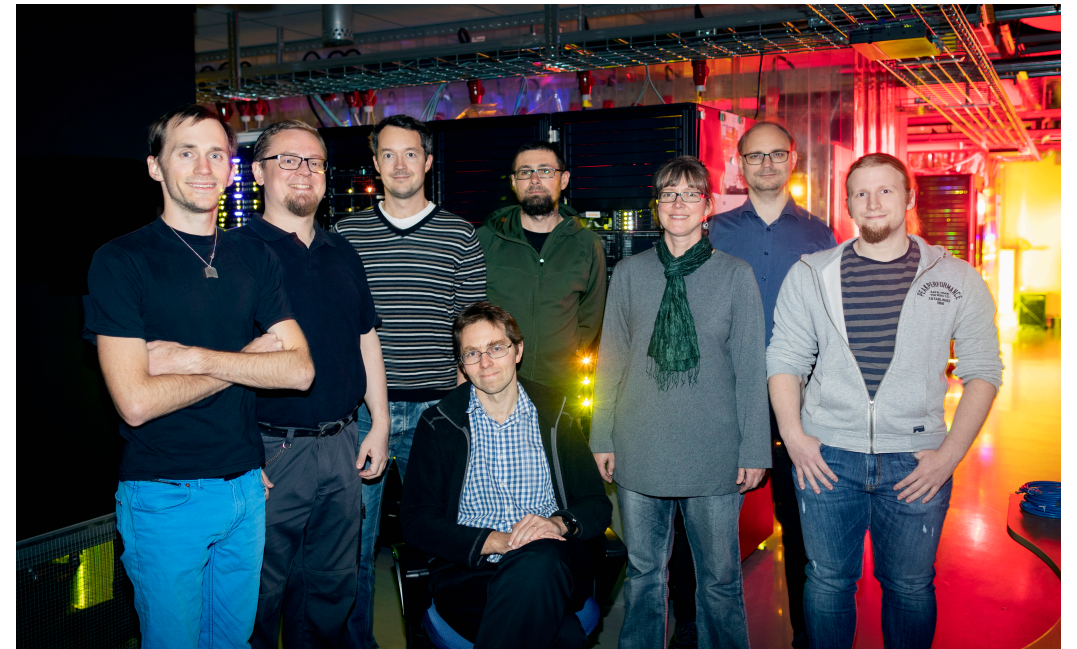
About 6 staff members

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

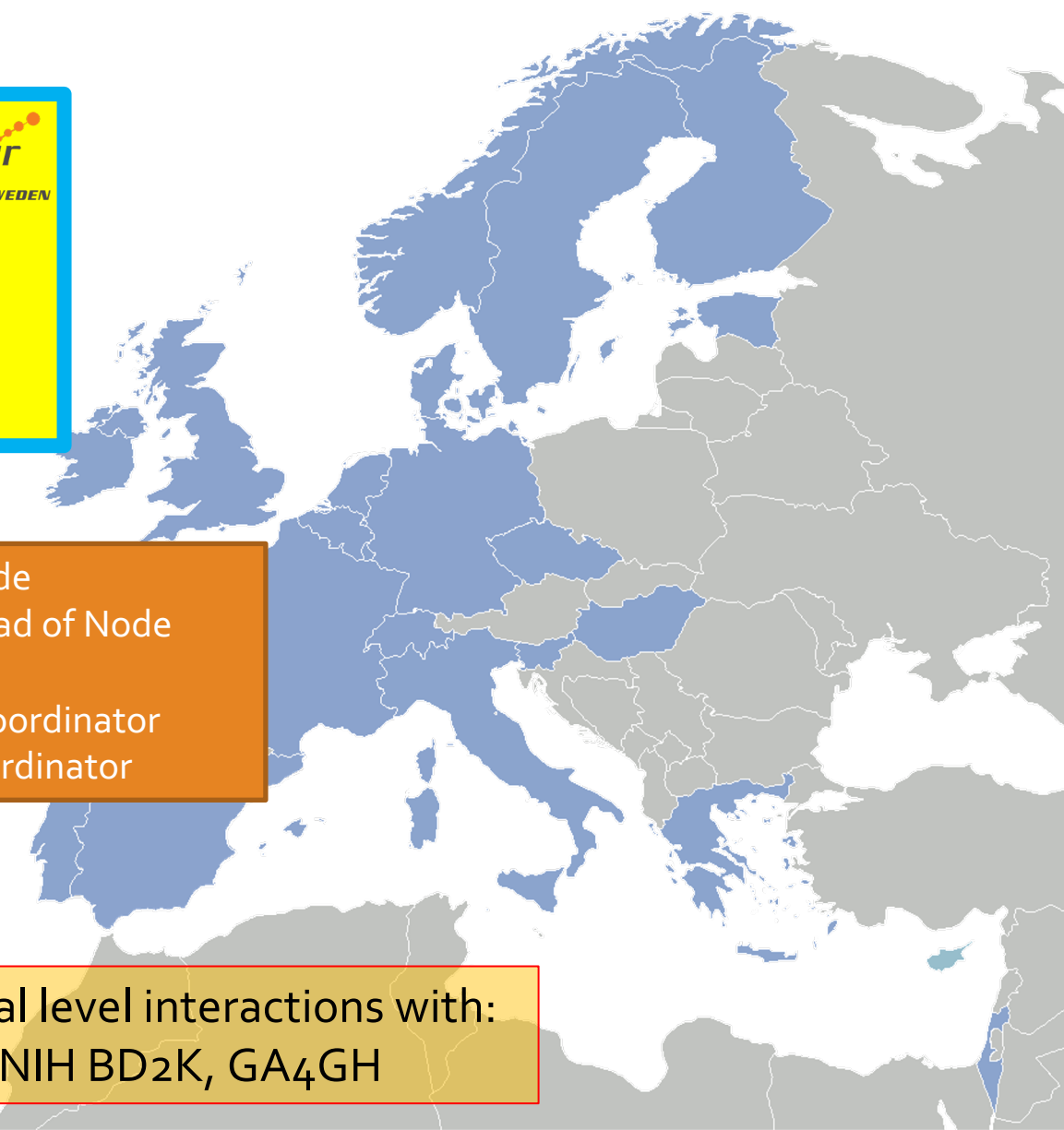
## Swedish contributions:

- 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



Any questions?



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