



National Bioinformatics Infrastructure Sweden, NBIS

8 September 2021







Analysis of biological data

- Support
- Infrastructure
- Training





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
- Responding swiftly to changes in support needs as new techniques are developed and utilised
- Forming the Swedish ELIXIR node and participating in relevant international



Technical experts are increasingly important in technology- and data-driven life science





170 active support 100 new projects 70 publications 250 drop-in sessions





Overview of NBIS units

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









SciLifeLab

Training events and programs

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

Rapid knowledge-transfer

- Drop-in sessions (Currently online)
- 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

1 Mput gene Sat				0
Species .	Pathene	Testures		
Human Human	web Hit	+ Cancer Genes		
AKT1 ATF1 ASX GMPS MEN1 M SDHD SMARCA	L1 ATM BRD3 BRD4 CR I2 MLH1 MSH2 MSH61 LSMARCR1 TCF3 TPR	EB1 CTNNB1 CBFB CREBBP C MYH9 NRAS NONO NFKB2 N	ARS DOXE DEK DI ID1 NPM1 PTEN P	CERT EWSRT ERCC3 F DRMT ROMTS SFPQ SI









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







Thank you!

Questions? Comments?