

Computational resources

What is needed? Where do I get it?

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Computational resources

- What do you need
- What options exist
- How do I access SNIC resources
- What next?



Computers as scientific equipment

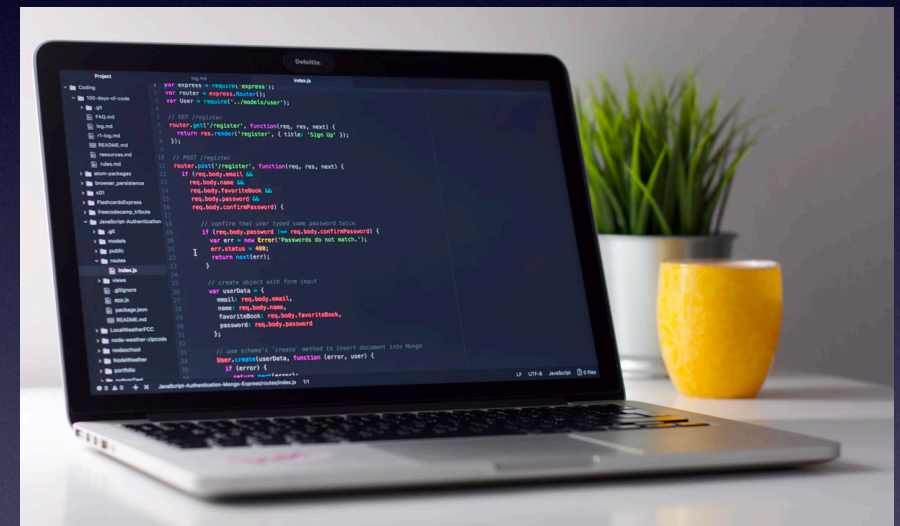
- For a given project, you'll have to evaluate your needs
- Consider what fits your needs:
 - a small self-maintained unit
 - a centrally-funded infrastructure
 - a combination or third option
- As with any equipment, training is beneficial or even necessary

Possible needs

- **Compute** needs: CPU cycles to run programs
- **Storage** needs:
 - Long-term space to store your raw data
 - Shorter-term space for analysis (raw data + results + temporary/intermediate files)
- Archiving (and Publication?)

Your laptop

- Good for:
 - Pre- and post-processing, scripting, etc.
 - A window into more powerful systems
- Limited by:
 - Disk space
 - Don't want to run long-running programs
 - Sensitive personal data not OK
 - Managed by you/local IT



Powerful workstation

- Good for:
 - Running longer programs (can do some genome analysis)
 - Visualisation
- Limited by:
 - Disk space limited to a few TB
 - Managed by yourself/local IT
 - Tricky to prepare for sensitive personal data
 - What happens when system grows old or the main user moves on?



“Shadow IT”

- Buy your own systems for analysis and storage.
- Put disk drives with important data in locked drawer.
- Upside: quick & dirty
- Downside:
 - disorganised
 - unreliable
 - waste of time and resources in the long run

Local IT services

- University and department IT support can often provide various types of storage servers and small clusters
- Better continuity, reliability than workstation

Swedish HPC Resources

- **SNIC** — Swedish National Infrastructure for Computing
 - Free to use
 - Collaboration
- **UPPMAX** — HPC center with large storage system
- **SNIC-SENS** — Secure system for sensitive personal data (WGS of living humans)
- **Compute and Storage** — SciLifeLab facility, part of Bioinformatics Platform, supports Life Science users at SNIC resources



SNIC Centra

- **HPC2N** - Very large memory, GPU's
- **UPPMAX** - Large storage
- **PDC** - Highly parallel processing
- **NSC** - Large general-purpose system
- **C3SE** - Göteborg
- **Lunarc** - Lund

Cloud

- **SNIC Science Cloud** (Free! Open to provisioning Software-as-a-Service and Platform-as-a-Service)
- **SUNet SafeSpring** (not free, ok for sensitive personal data)
- International science clouds
- Commercial providers:
 - Amazon web services (AWS)
 - Microsoft Azure
 - Google Cloud Platform

What SNIC can/can't be used for

- **Do:**
 - Analyse data
 - Store data while you're working with it
- **Do not:**
 - Use to (only) store/backup/move a dataset
 - Do patient diagnosis
 - Run a business



Applying for SNIC project

- At UPPMAX and elsewhere, the procedure is the same
- Projects are managed in SUPR (supr.snic.se)
- Resources are granted to projects
- Project membership determines which files are readable and how much compute power is available
- <https://www.uppmax.uu.se/support/getting-started/applying-for-projects/>

Types of projects



- Compute project
- Storage project
- SNIC SENS
- Cloud

Scale of project

	Compute (core-hours/ month)	Storage (TB)	SENS
Small	< 10,000	< 10	< 10 kch/m < 20 TB
Medium	< 100,000	< 50	< 100 kch/m < 150 TB
Large	> 200,000	> 50	> 100 kch/m > 150 TB

The cost?

- Free to you but...
- Rackham's extension cost (very roughly, only initial investment):
 - 375 kr/TB per year
 - 0.1 kr/core-hour
- A typical PhD-student's project (1 TB, 1000 core-hours/month for four years): 6,300 kr.
- An ongoing potato genomics project (20 TB, 1000 core-hours/month): 8,700 kr each year.
- A large shotgun genomics project (5 TB, 30,000 core-hours/month) for half a year: 19,000 kr.
- The creation of a new reference genome, requiring 100 TB and 50,000 core-hours/month for one year: almost 100,000 kronor.
- Large molecular dynamics simulations, using 200,000 core-hours/month: 240,000 kr/year.

So what do I need?

- If in doubt: apply for a **SNIC Small Compute** (at UPPMAX if you want a large software library)
- Apply for **storage** for:
 - Your sequencing data (NGI can tell you how much)
 - Plus an expansion factor (2-5x)
- Apply for both ideally about 1 month before NGI is ready to deliver data

Using SNIC/UPPPMAX

- Linux command-line
- Graphical remote Linux desktop
- Training available via SNIC, UPPMAX, and NBIS

Getting data to UPPMAX

- From **NGI**: Data is placed on the Grus server and a “delivery project” is created on SUPR. You can then move the data to Rackham or Bianca by clicking a button in SUPR.
- From **elsewhere**: upload/download as normal using scp, rsync, sftp, wget, git, etc

Software at UPPMAX

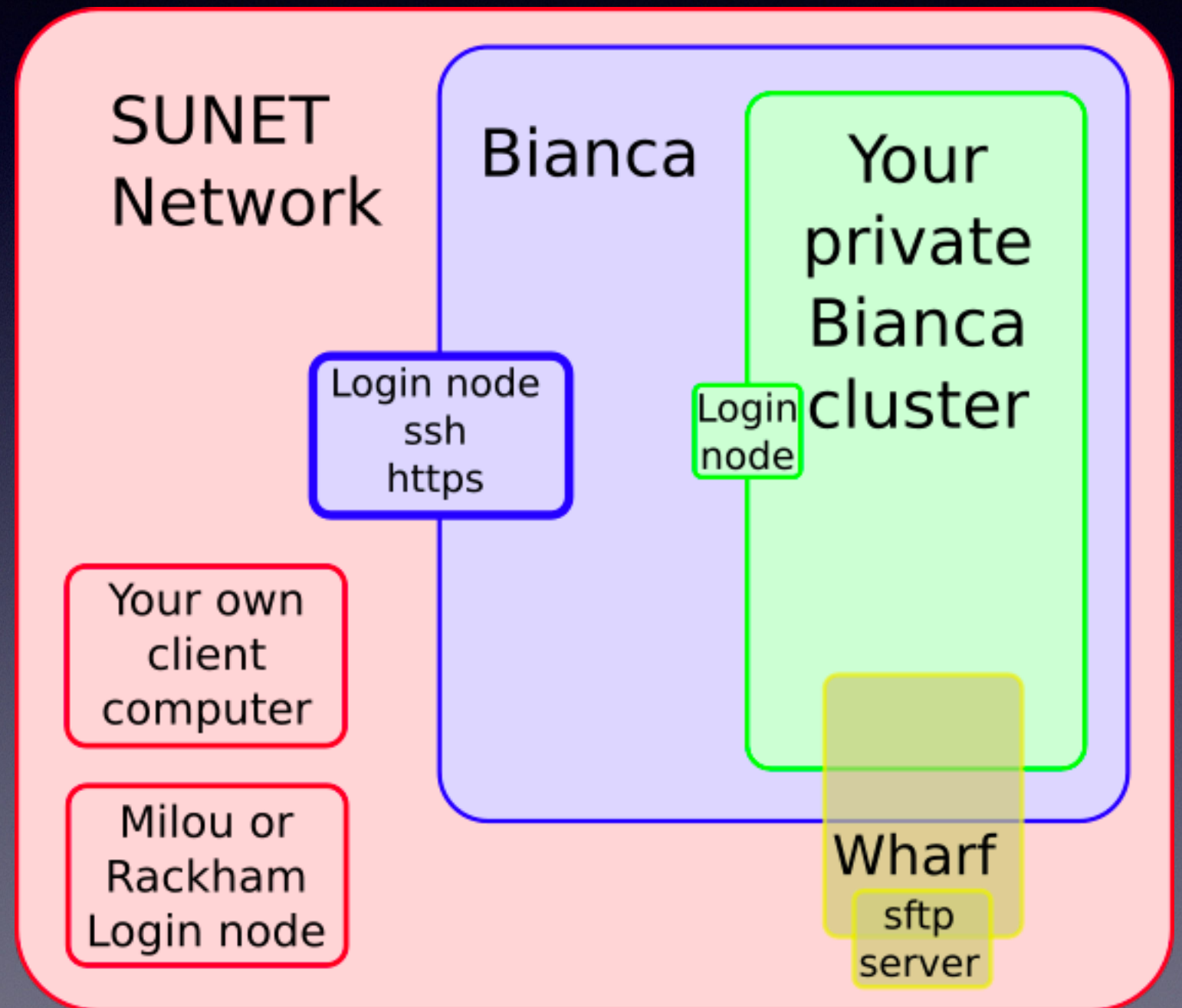
- A huge software library is installed and maintained
- Additional software can be requested, but be aware this can take a while
- Most software can be installed by yourself

Sensitive personal data

- What is (or might be) sensitive personal data?
- Why do you need additional technical measures?

Bianca (SNIC SENS)

- “Lätt att göra rätt”
- A secure HPC environment
- Collaboration-friendly



transit.upppmax.uu.se

- A server is set up that allows easy “mounting” of Bianca wharfs.
- Useful for:
 - Copying data between SENS projects
 - Downloading data from internet to a SENS project

When does a project end?

- SNIC project lasts 12 months (typically).
- Must be extended or replaced, or access will be revoked.
- SNIC Common User Agreement states that data may be deleted after 6 months.

What next?

- After analysis, your data continues to live. Think about:
 - Publishing
 - Depositing into a repository
 - Hand it over to a university archive
 - Storing it yourself
 - Cloud (e.g. SUNet SafeSpring)
 - UPPMAX Offload Storage (Lutra)
 - SNIC Swestore
 - USB drive or HDD on a shelf in your office

Lutra (UPPMAX Offload Storage)

- For larger projects or groups
- 50 TB increments, 4 years
- Annual cost is 25,000 SEK/increment

Compute and Storage Resources

- The landscape is always changing
- Expertise is available
- Just ask if you have any questions:
 - support@uppmax.uu.se
 - marcus.lundberg@nbis.se
 - Anyone else you've heard from today

SNIC User Forum @ KI

- For current and future users and owners of SNIC projects
- September 27th. Theme is Bioinformatics.
- Program at: <https://staff.ki.se/snuc-user-forum-at-karolinska-institutet-2019>
 - Posters from related infrastructures
 - Information from SNIC
 - Presentations from users

Thank you for listening