

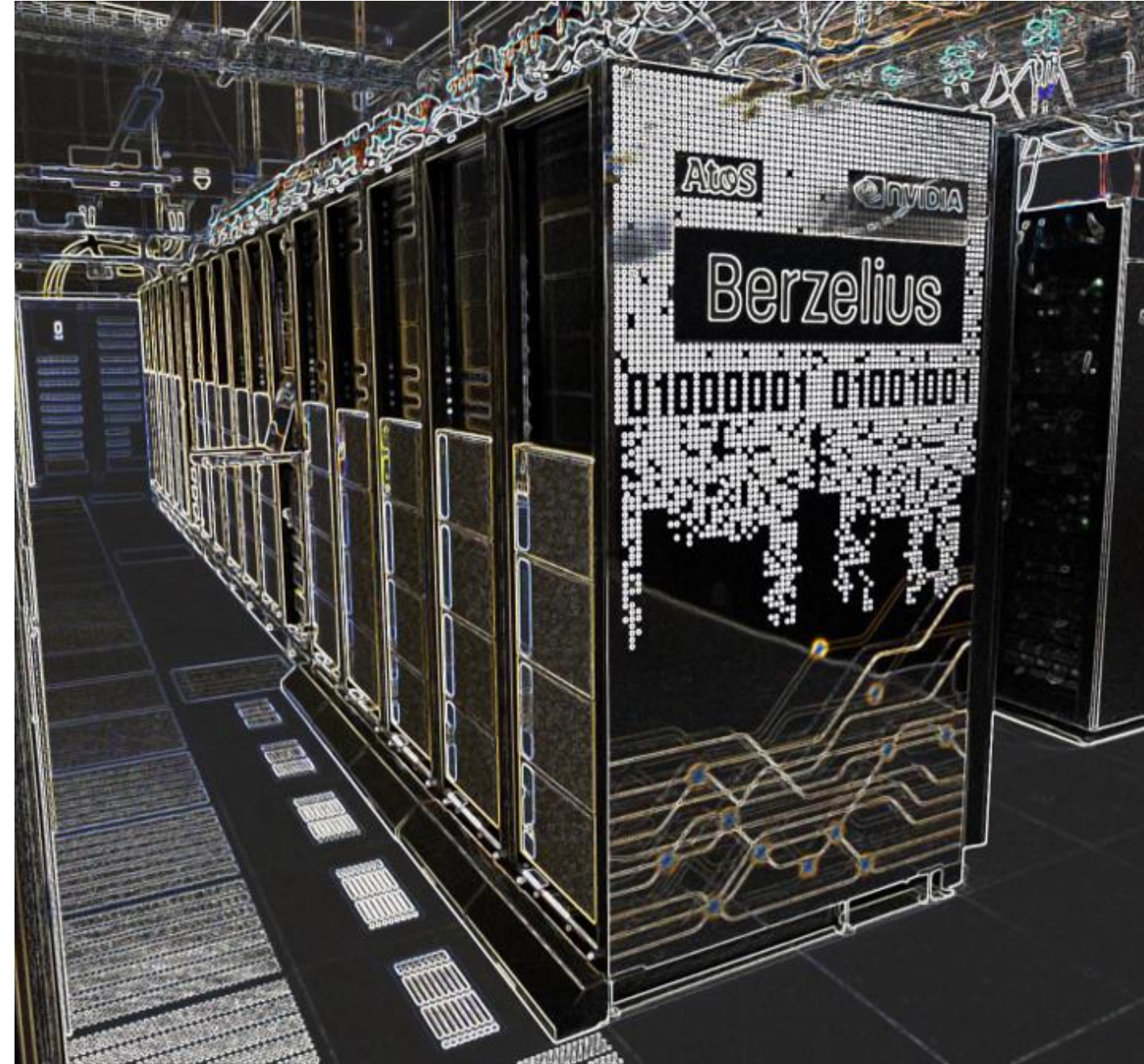
Visual Intelligence



Anders Ynnerman
Norrköping Visualization Center
WASP
Linköping University
Sweden



EXPERIMENT



COMPUTE

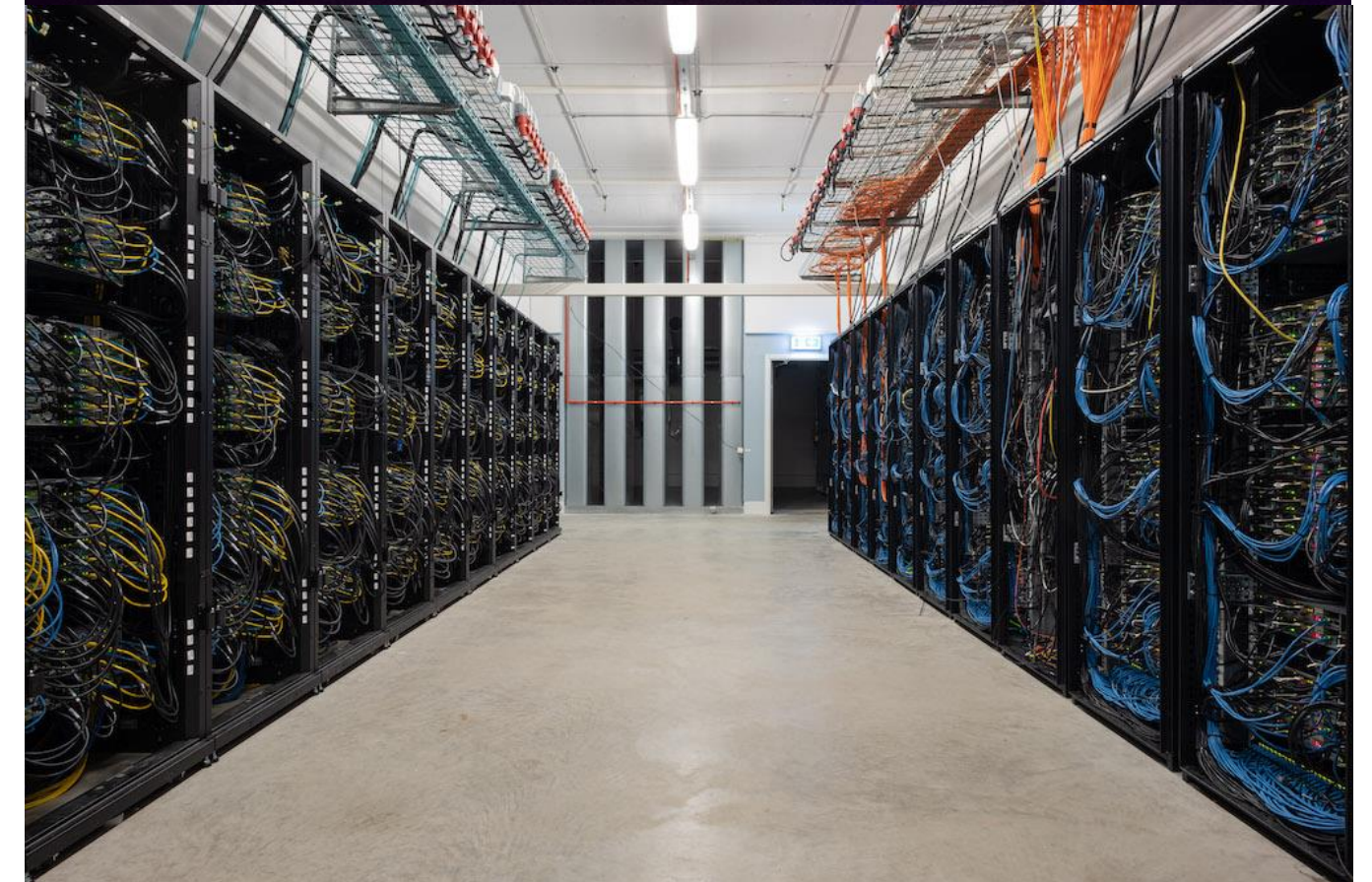
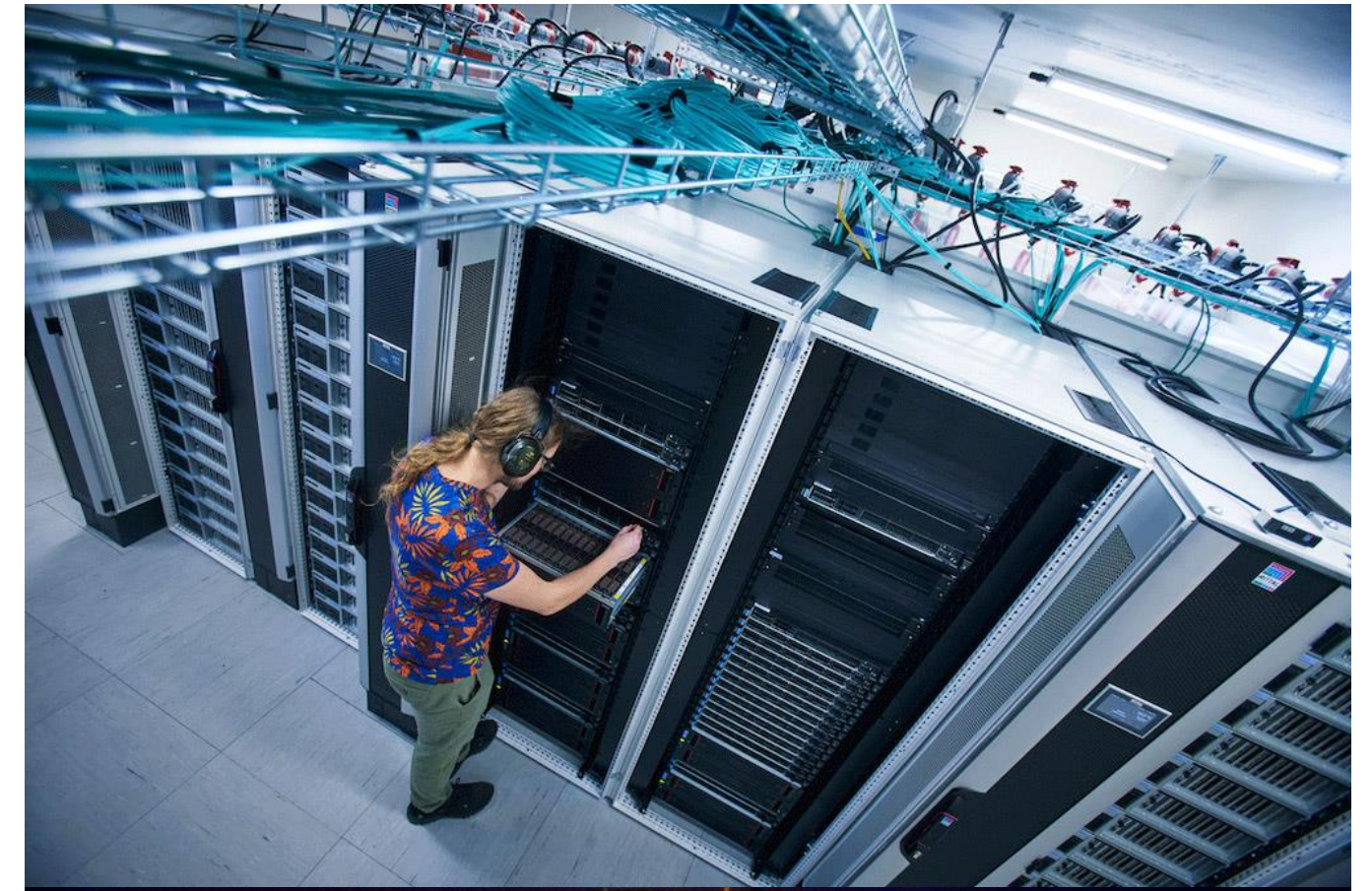


DATA

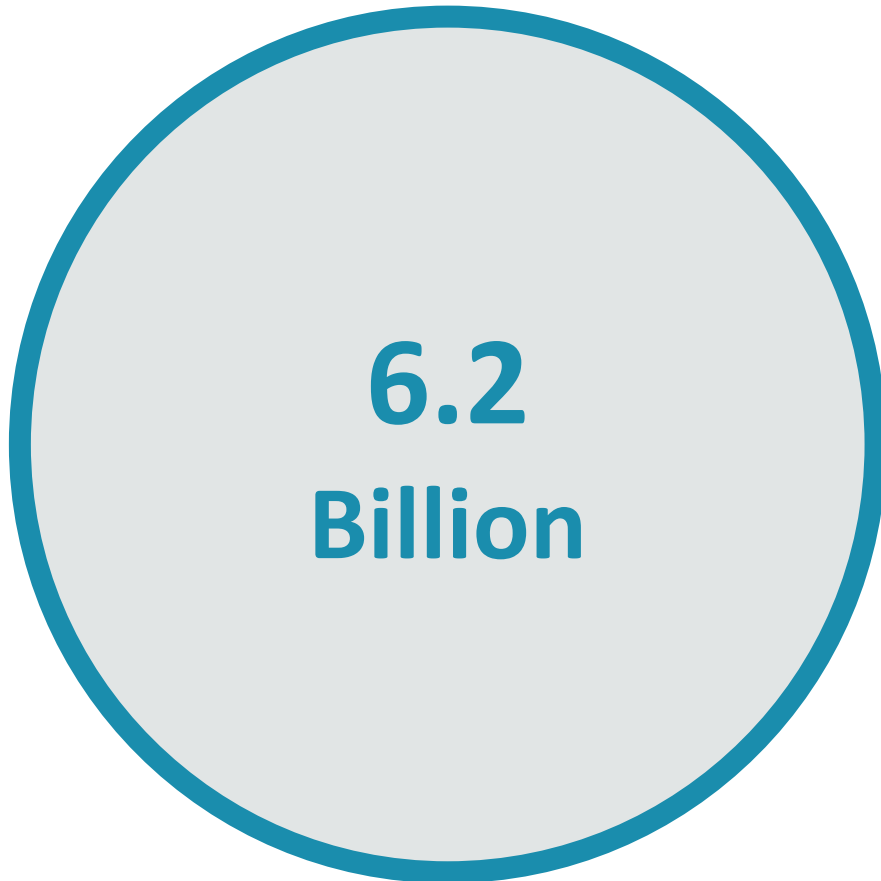
NAISS

National Academic Infrastructure for Supercomputing in Sweden

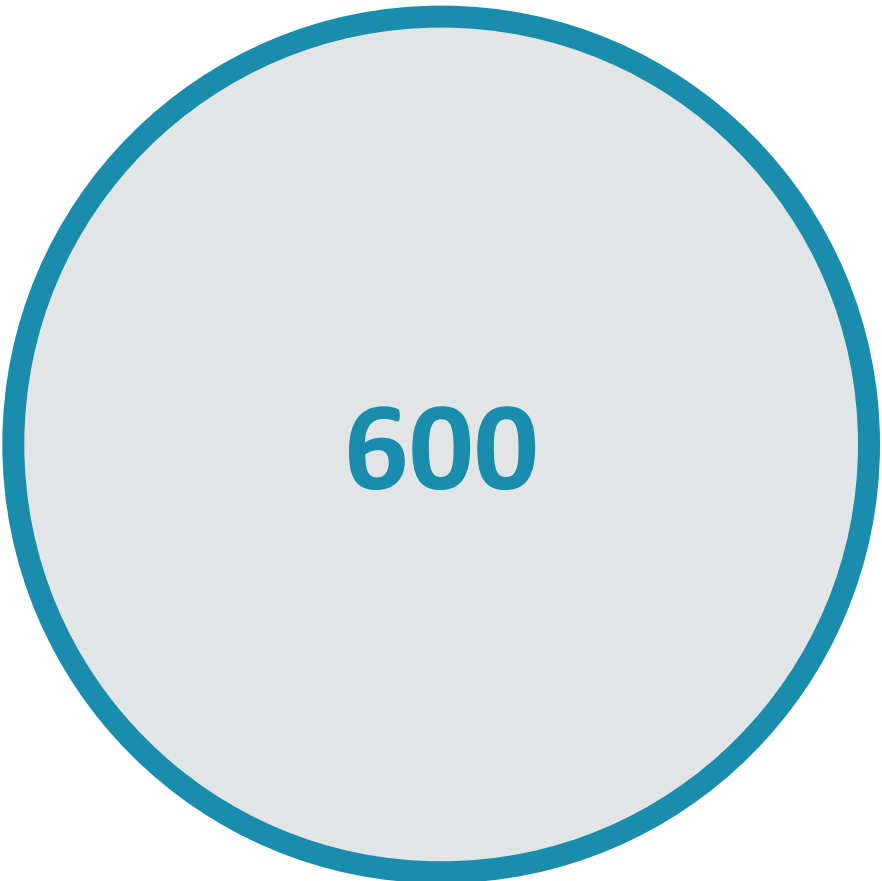
- The new national research infrastructure organisation for HPC in Sweden
- Hybrid model – centralized hardware and distributed support
- NAISS is extending user support and applications expertise across Sweden with branches in partnership with all large universities
- Linköping University is the host, the Swedish research council the main financier



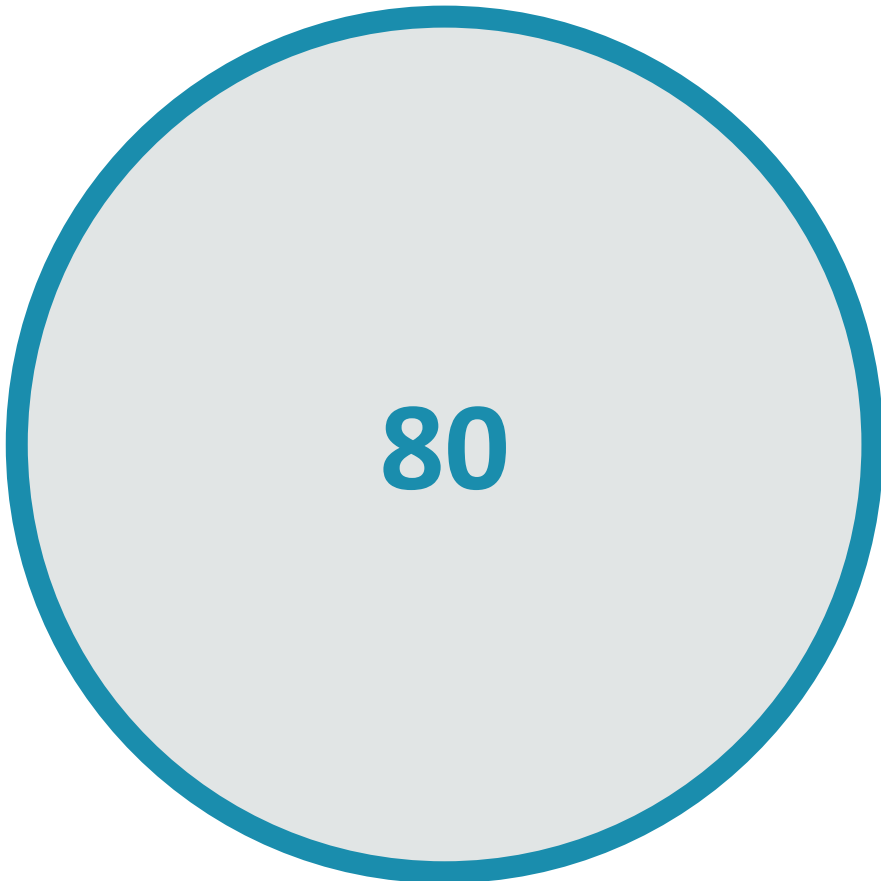
Wallenberg AI, Autonomous Systems and Software Program



6.2 billion SEK for 15 years until 2031



600 Graduated PhDs



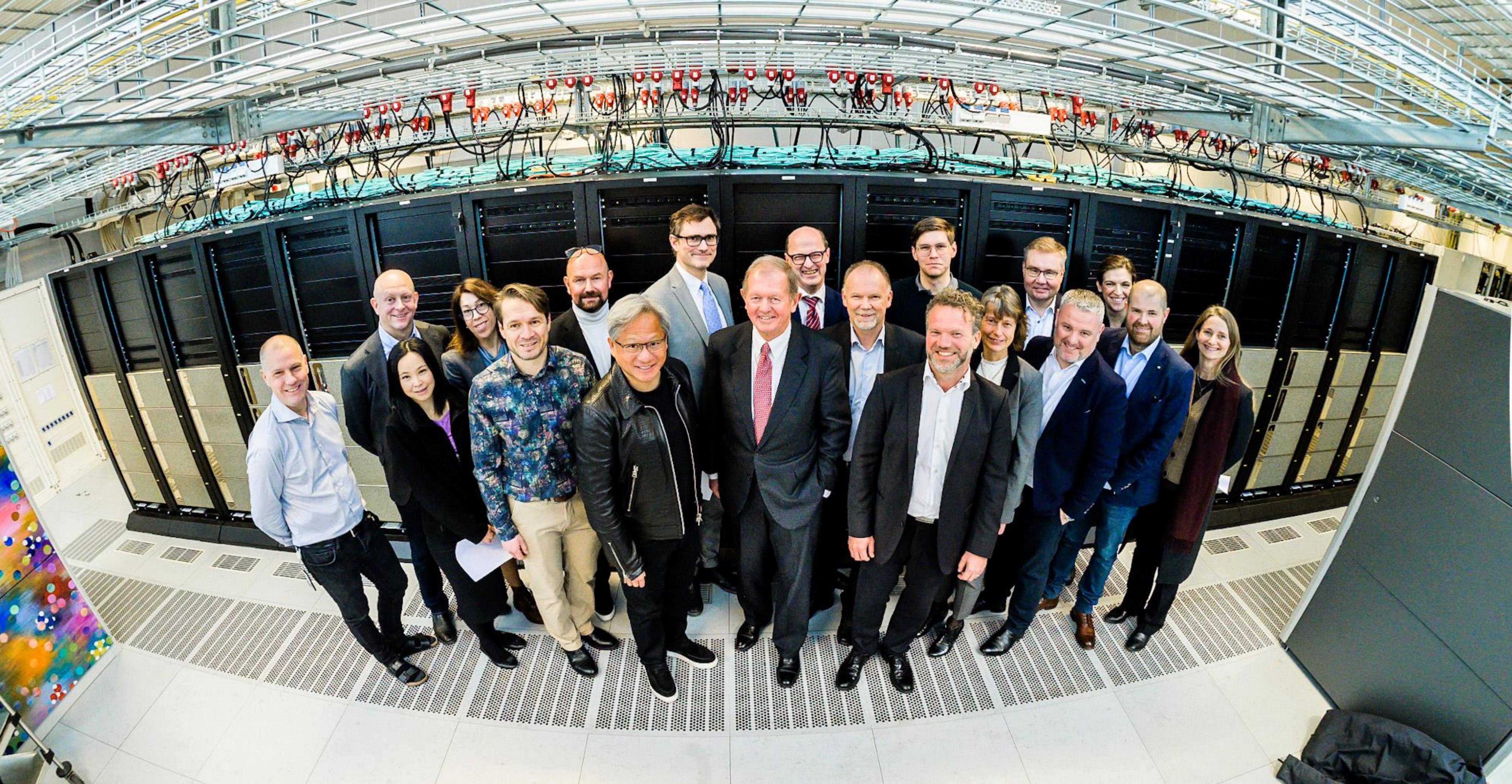
80 Faculty Recruitments

Berzelius - AI Resource

- ATOS - NVIDIA SuperPod
- 94 DGX A100 systems
- 8 A100 GPUs/node
- 5 PetaFLOPS/node
- 1800 Gb/s interconnect
- Hosted by the National Supercomputer Centre (NSC) and integrated in the HPC environment for compute and storage solution

- Fully utilized as of April 2022
 - Natural Language Processing (GPT-SW3)
 - Protein Folding
 - Computer Vision
 - ...







— Ours — ASLA — SCM

Berzelius and Computer Vision

- Berzelius is our backbone for large scale experiments on machine learning for computer vision.
- Data-driven models for detection and semantic segmentation in static image, tracking and segmentation of objects in video sequences, and for solving recognition and segmentation problems on 3D point clouds.
- Leads to publications at top tier conferences in engineering and computer science, such as CVPR, NeurIPS, ICCV, ICML, and ECCV.

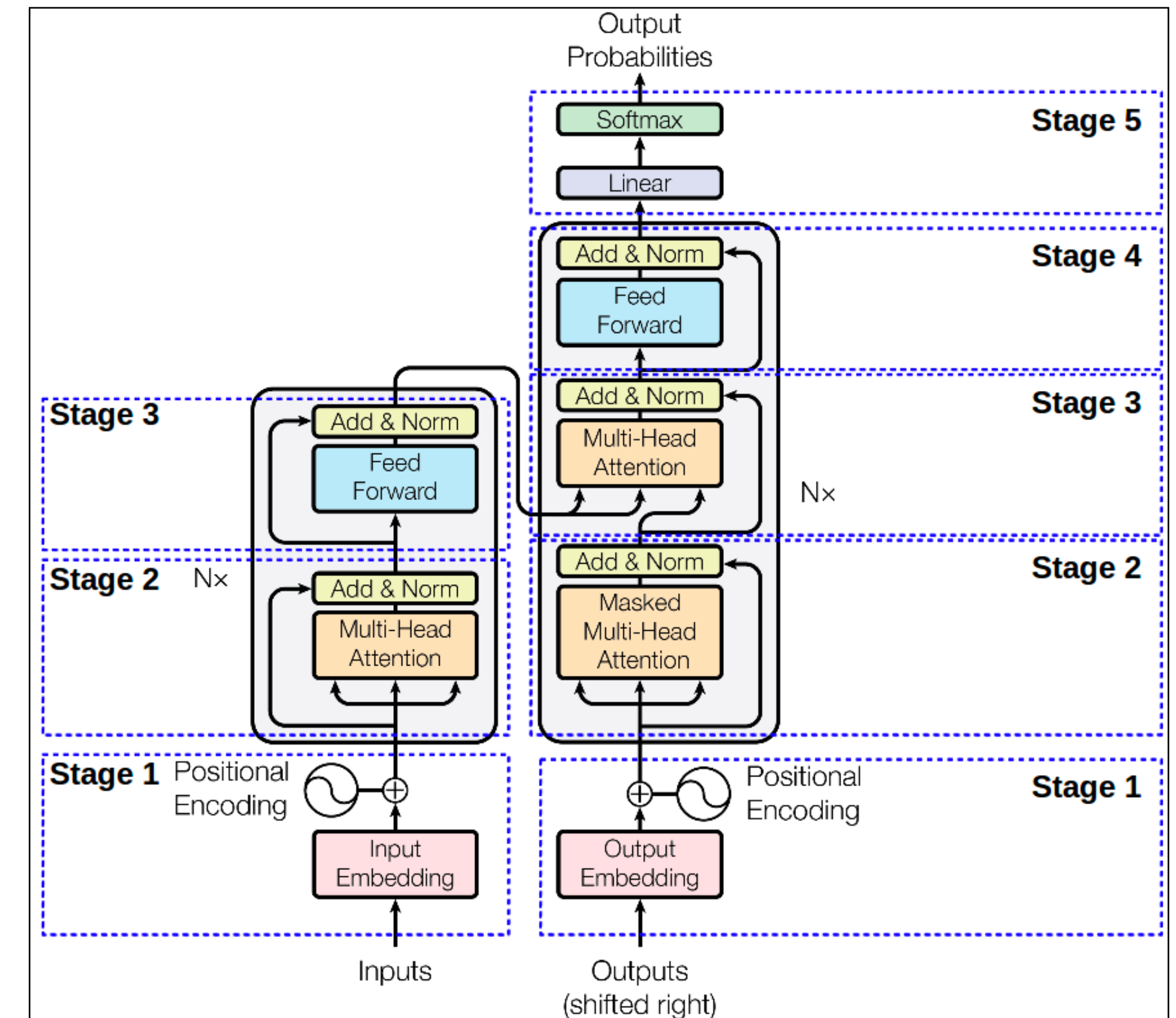


Berzelius and Protein Folding

- AlphaFold, an AI breakthrough from DeepMind, is revolutionizing structural biology by predicting protein structures with remarkable accuracy.
- Innovative sampling strategy involving the activation of dropout layers within AlphaFold's neural network.
- The approach led to increased structural diversity, discovering alternative solutions that not only matched the correct answer but were also consistently ranked high by the self-assessment scoring function.

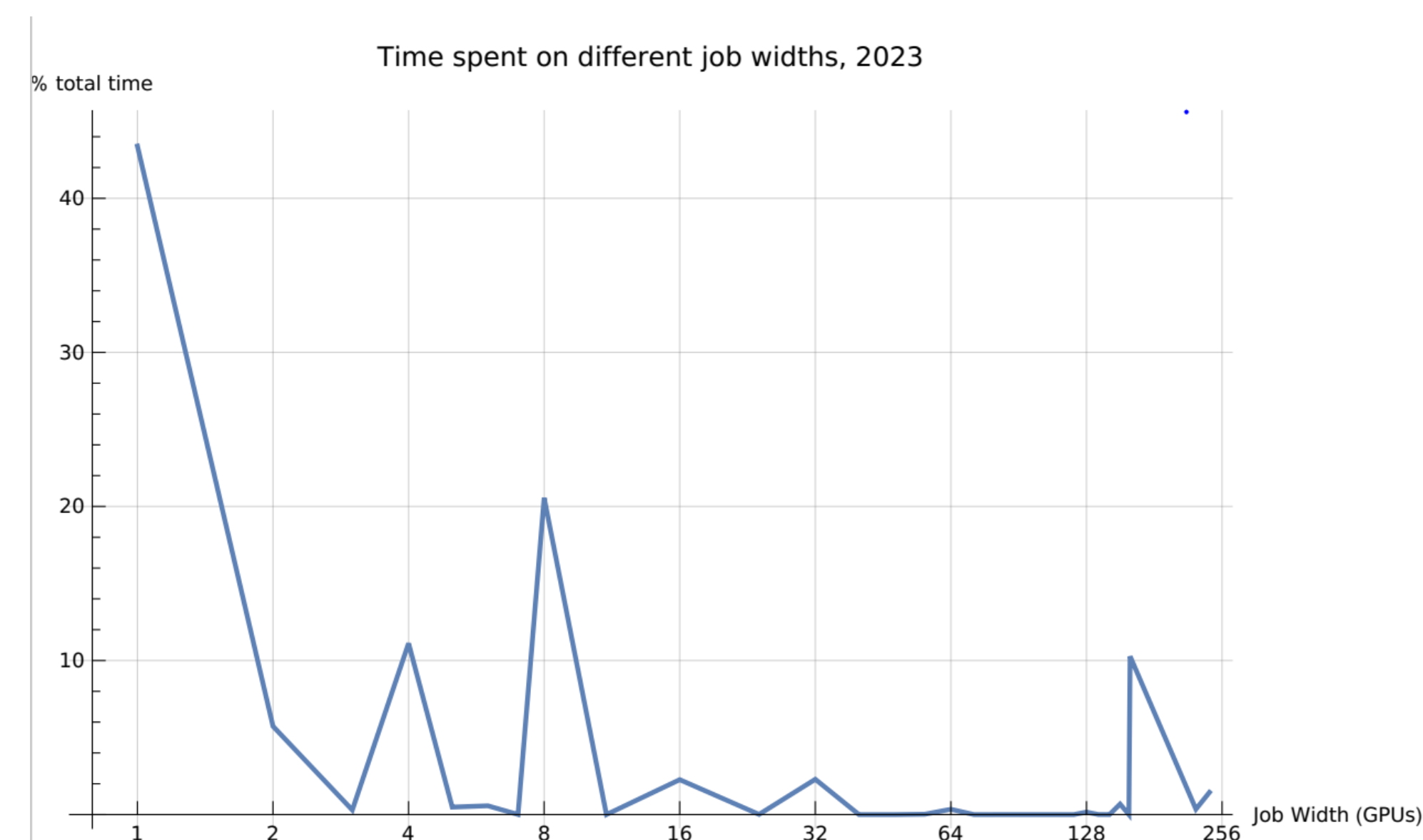
Berzelius and NLP

- Nordic language models of sizes 126m 356m 1.3B, 7B, 20B and 40B on Berzelius.
- “Closed open access”, i.e., on request.
- Trained on The Nordic Pile, a dataset of 1.2TB freely available data in the Nordic languages, collected from e.g. ArXiv, PubMed, Reddit, Flashback, etc.
- The largest model, i.e., of 40B required 9.11×10^{22} FLOPs for training on Berzelius.
- The value of models are found in the openness and transparency, as well as the competence build-up associated with the work on the models.



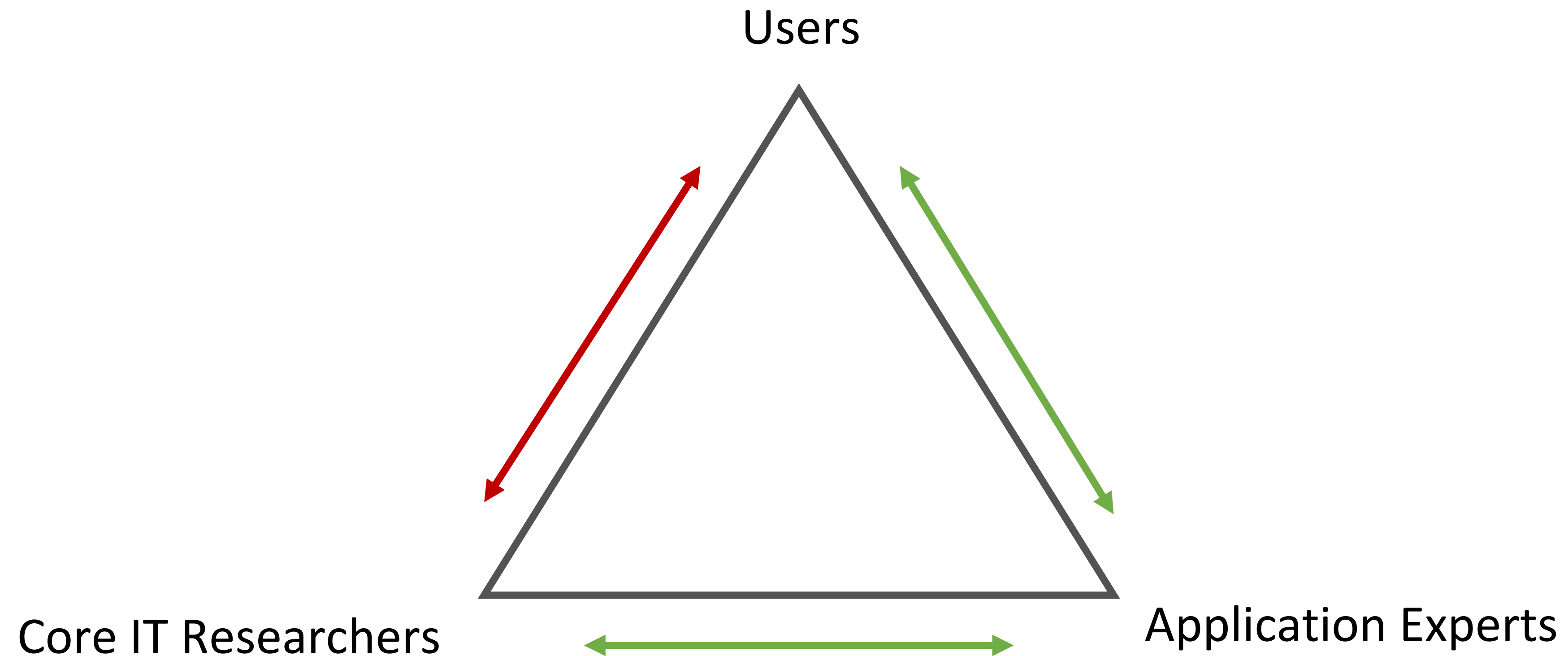
Current situation

- Load varies and peaks during daytime with shorter jobs.
- Shorter and small jobs do not utilize the capacity of the system in terms of scaling and addressing large problems.
- Scaling of jobs to make full use of Berzelius requires user training.
- A new generation of application support staff is needed to address large scale AI problems.

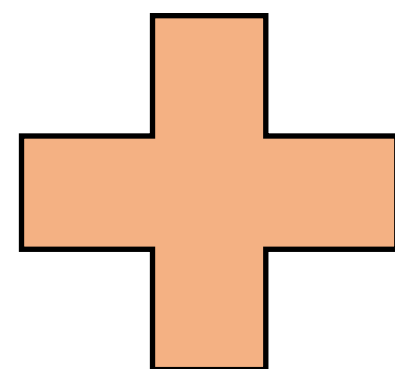
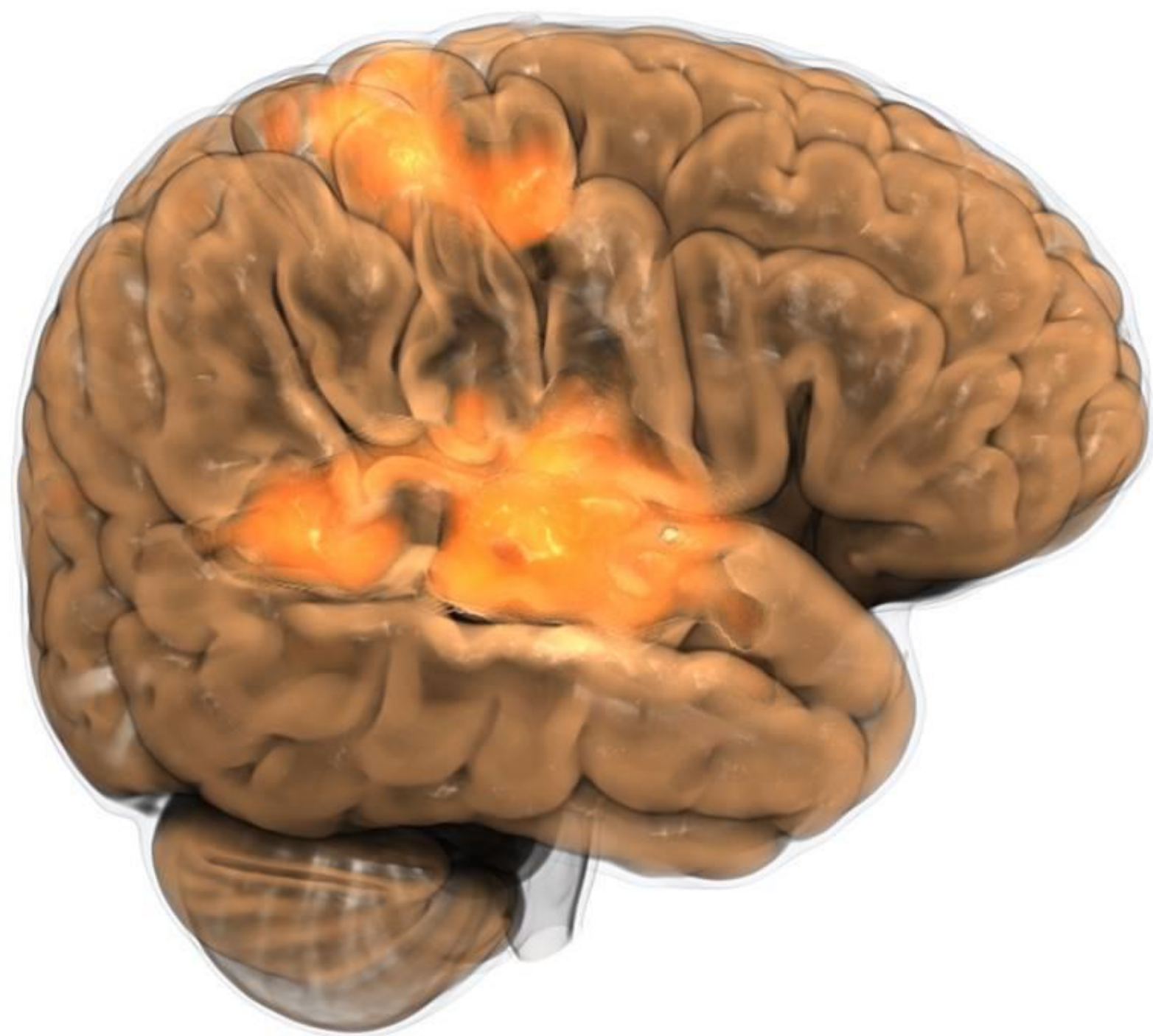


Distribution of job sizes on Berzelius during 2023. Approximately 42% of the used compute time is single GPU jobs

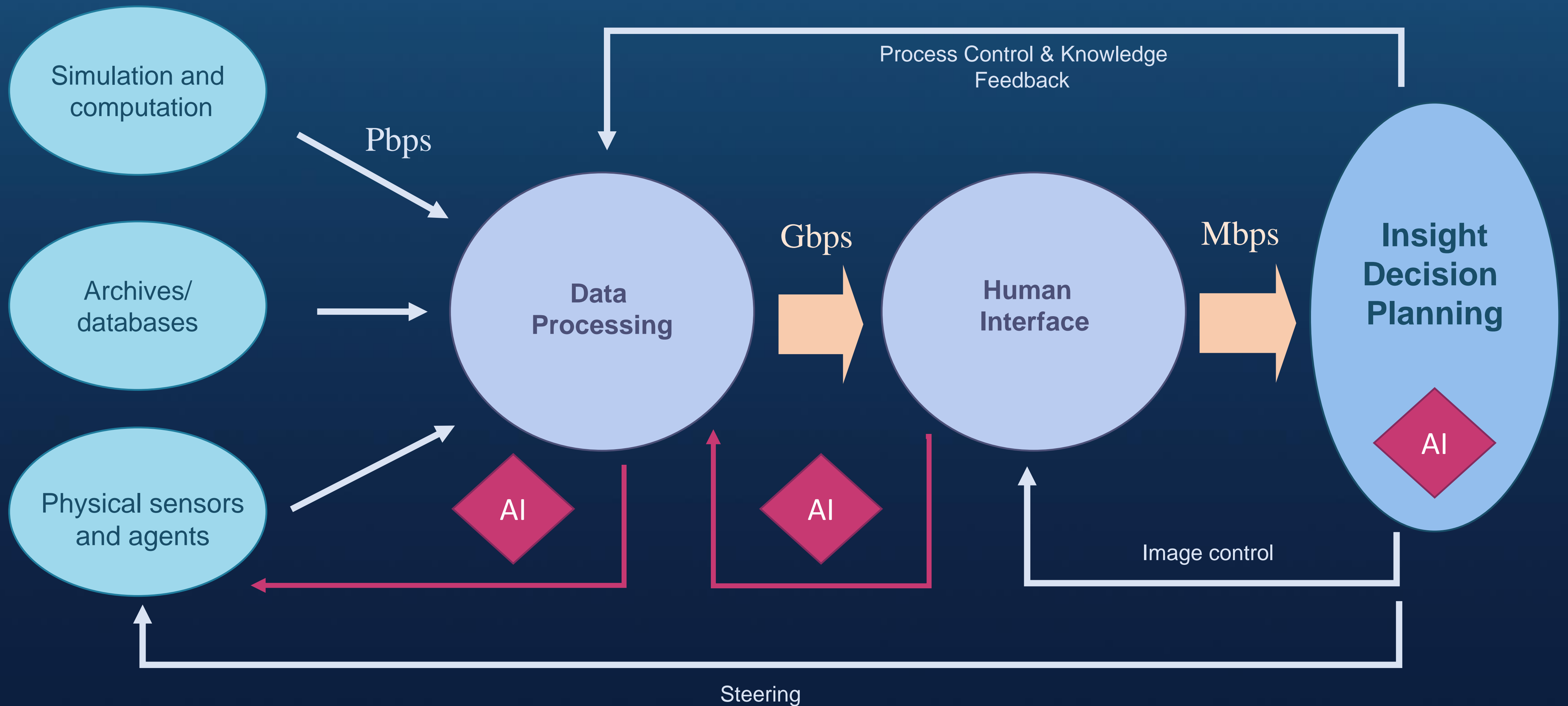
The SeRC Skill Triangle



Berzelius - AI Resource



Visual Data Analysis

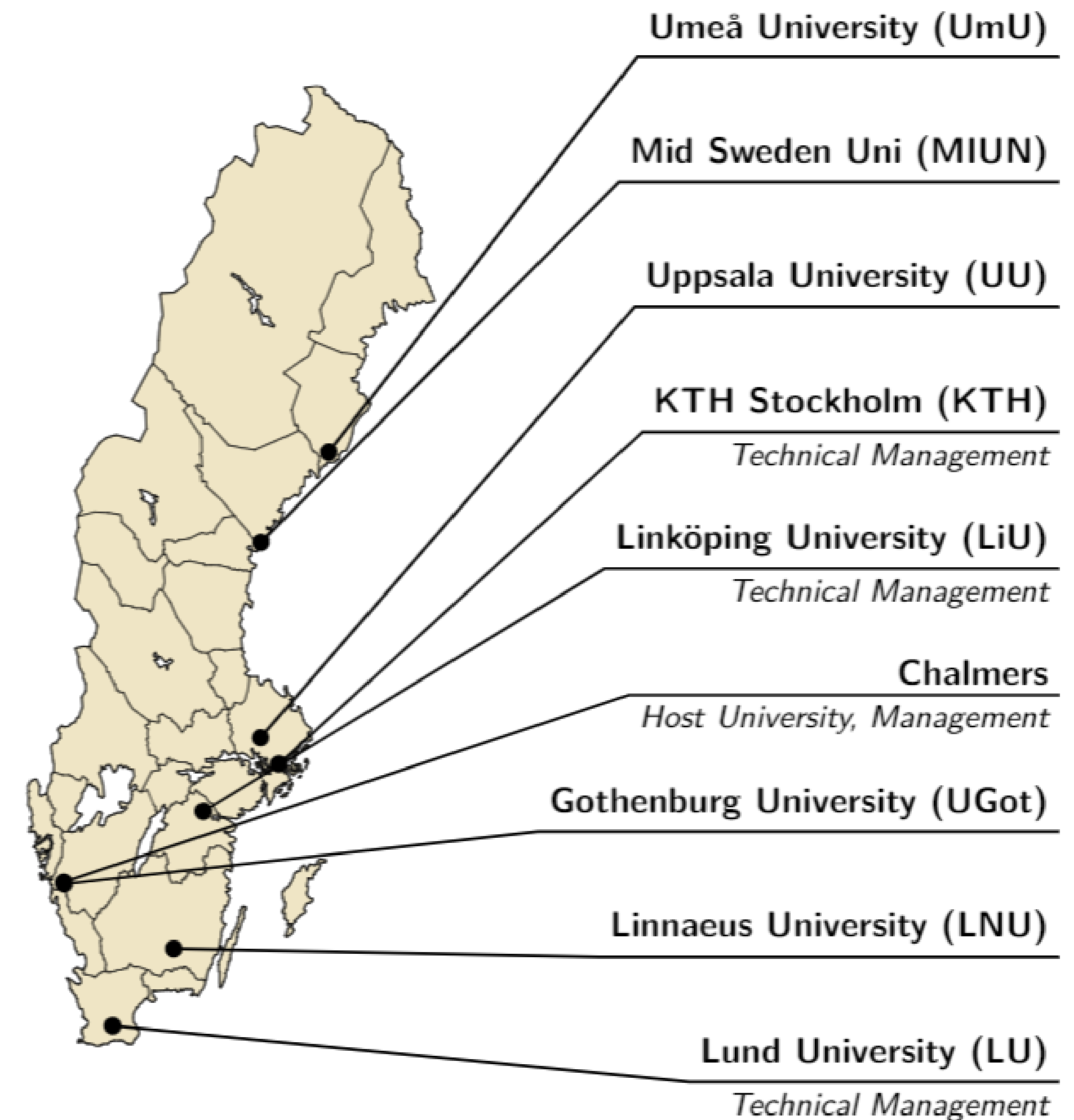




InfraVis

Supporting Scientific Discovery

- Visualization Infrastructure
 - Humans
 - Software
 - Services
 - Hardware



Swedish
Research
Council



CHALMERS
UNIVERSITY OF TECHNOLOGY



UNIVERSITY OF
GOTHENBURG



KTH
VETENSKAP
OCH KONST



li.u LINKÖPING
UNIVERSITY



LUND
UNIVERSITY



Linnaeus University



Mittuniversitetet
MID SWEDEN UNIVERSITY



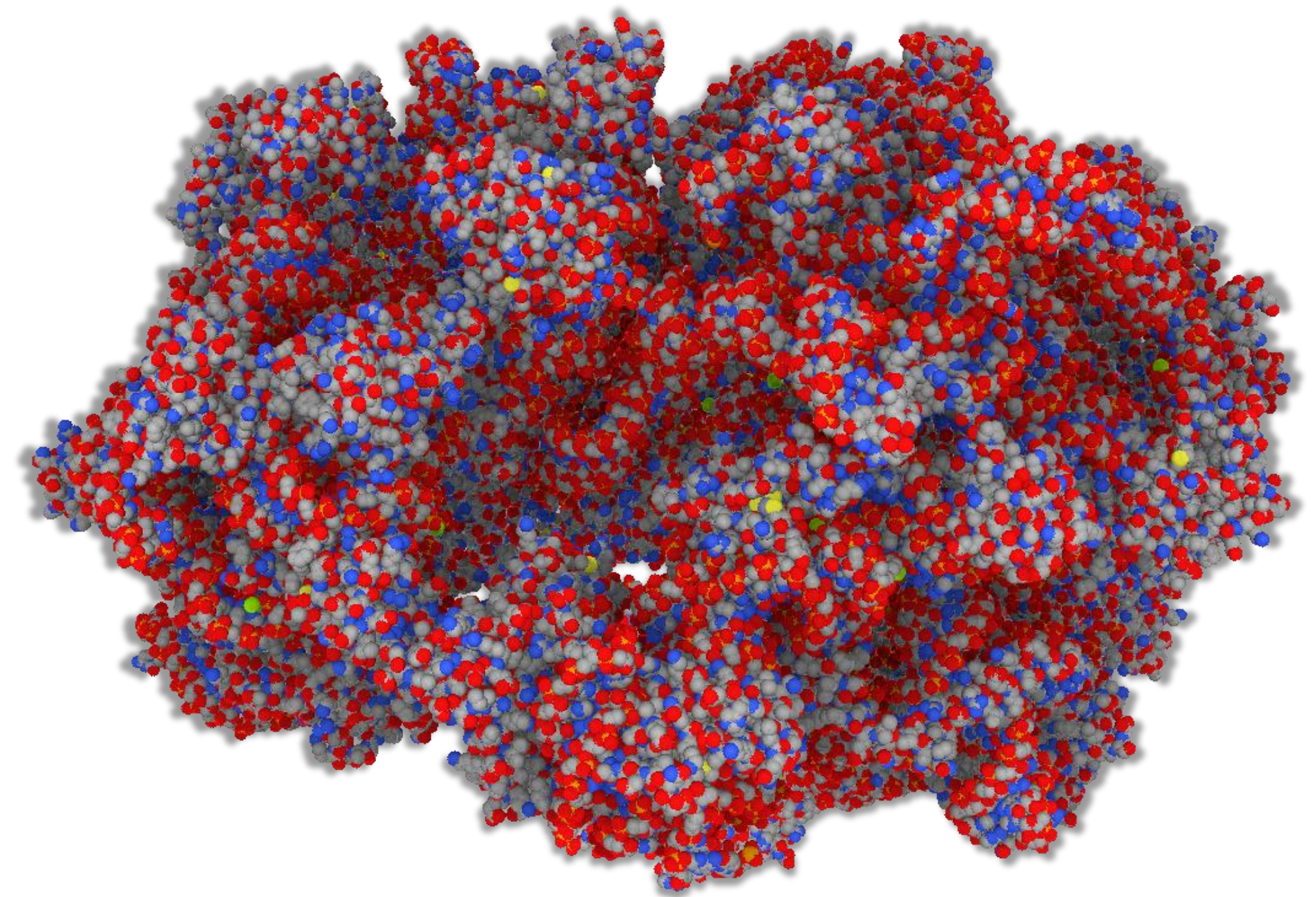
UMEÅ
UNIVERSITY



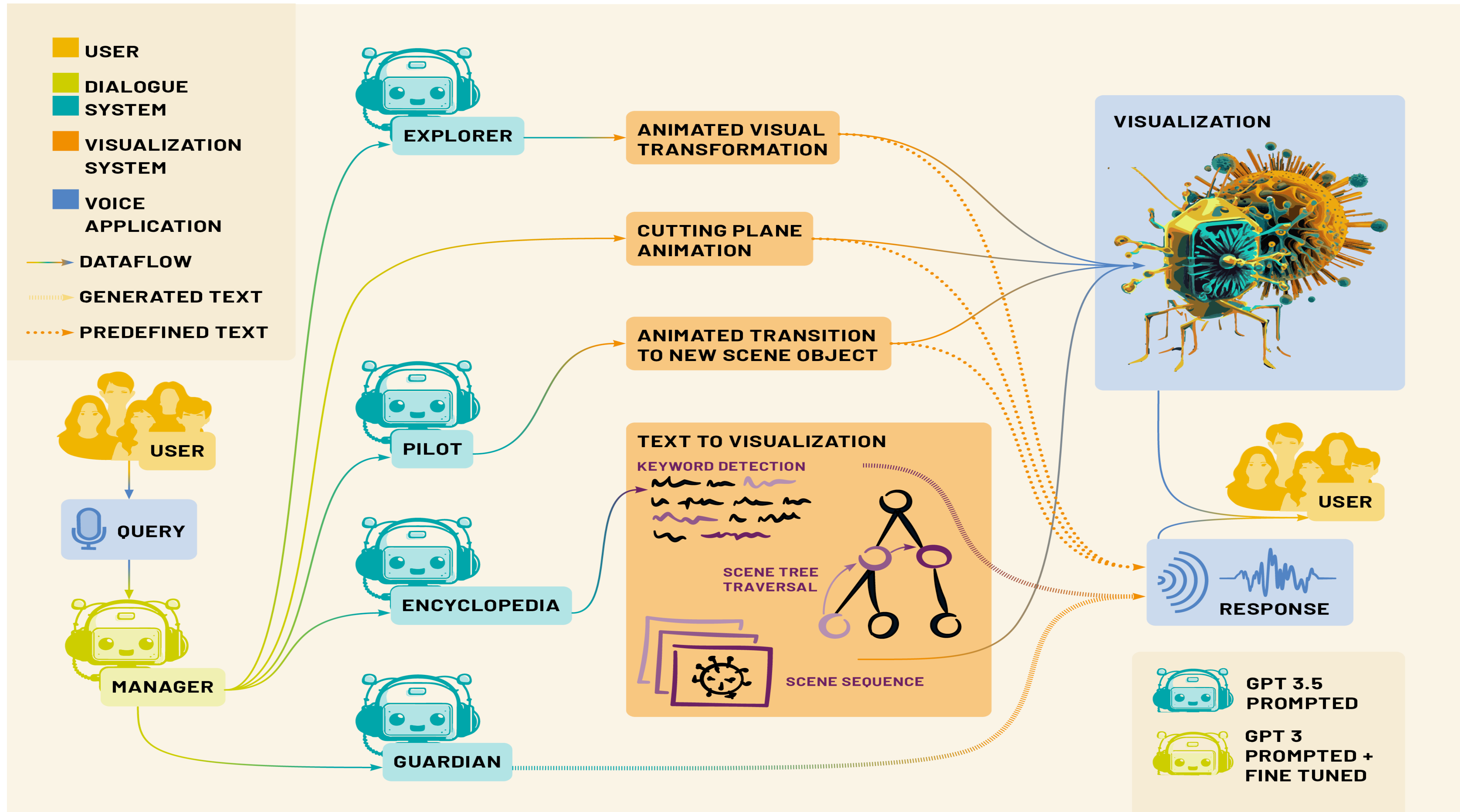
UPPSALA
UNIVERSITET

InfraX4Science

- X tools for data driven scientific discovery
- An infrastructure of application experts is needed with a (new) wider set of skills:
 - large scale ML and AI deployment
 - interactive human in the loop data analysis
 - data collection and curation
 - statistical data analysis
 - interactive and collaborative visual analysis
 - regulatory aspects
 - ...



AI-Enabled Conversations



AI-Enabled Visual Conversations

The image shows a 3D molecular model of a SARS-CoV-2 virus particle. The model is spherical and composed of several components: a grey outer shell representing the envelope, a blue inner shell representing the membrane, and numerous orange and red spike proteins protruding from the surface. Labels identify the following parts: Envelope protein, Membrane, Spike protein (closed), Spike protein (open), and Membrane protein. A yellow label points to a specific spike protein structure. A scale bar in the bottom right corner indicates 10 nm. The interface includes a top menu with 'Program', 'Widgets', 'Themes', and 'Views', and a 'HOME >' button. A chatbox on the right side contains the following text:

20:14

Welcome!

I am your personal tour guide through SARS-CoV-2 model.

You can ask me questions about it and I will do my best to answer them.

Let's start exploring together!

Model: SARS-COV-2 | SpeechToText | Whisper | ManagerBot | ChatGPT

✓ GPT-4 | ✓ Parallel | ✓ Interactive Content | ✓ Overview Orbiting | Subtitles | SEND

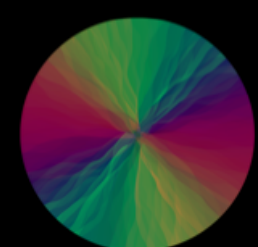
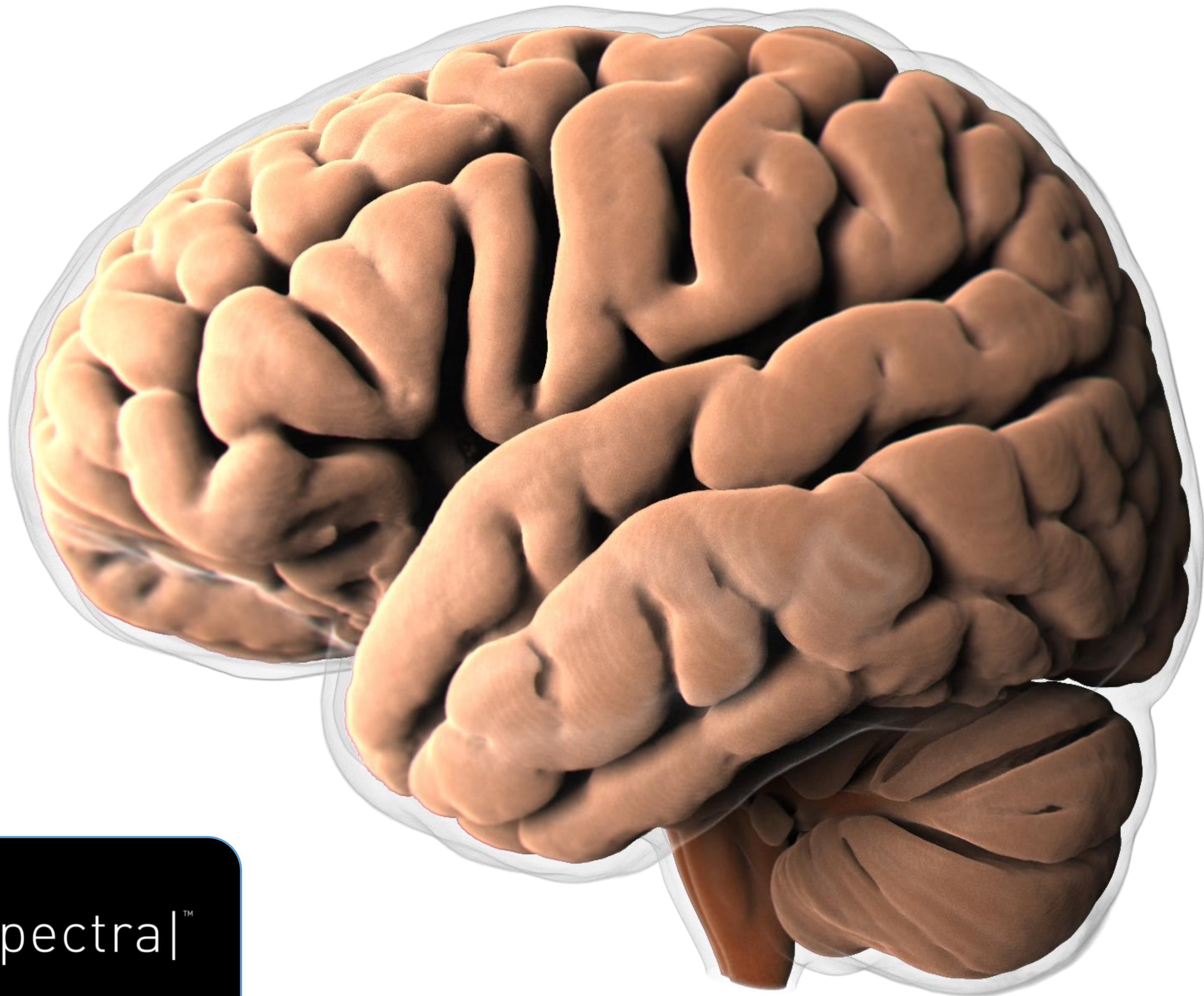
Chatbox | Selections

Pixel: (0.000, 0.000, 0.000, 0.000) | (0.00, -0.00, 2500.00) | 40.6 FPS

Future Scientific Discovery

AI will fundamentally change scientific discovery workflows and communication of results

- Reduce data flows**
- Lower cognitive load**
- Raise level of abstraction**
- Enable Advanced interfaces**



Interspectra™

Large Scale Human Data

- **European Synchrotron in Grenoble**
- **Human Organ Atlas**
- **HiP-CT Hierarchical Phase-Contrast Tomography**
 - 25 micron resolution (100 times higher resolution than medical CT)
 - Organ overview and cell detail
 - Volumetric rendering by spin-off company Interspectral from Visualization Center C



