



12TH FENIX INFRASTRUCTURE WEBINAR

EBRAINS services deployed on ICEI services: Neuromorphic Computing front-end

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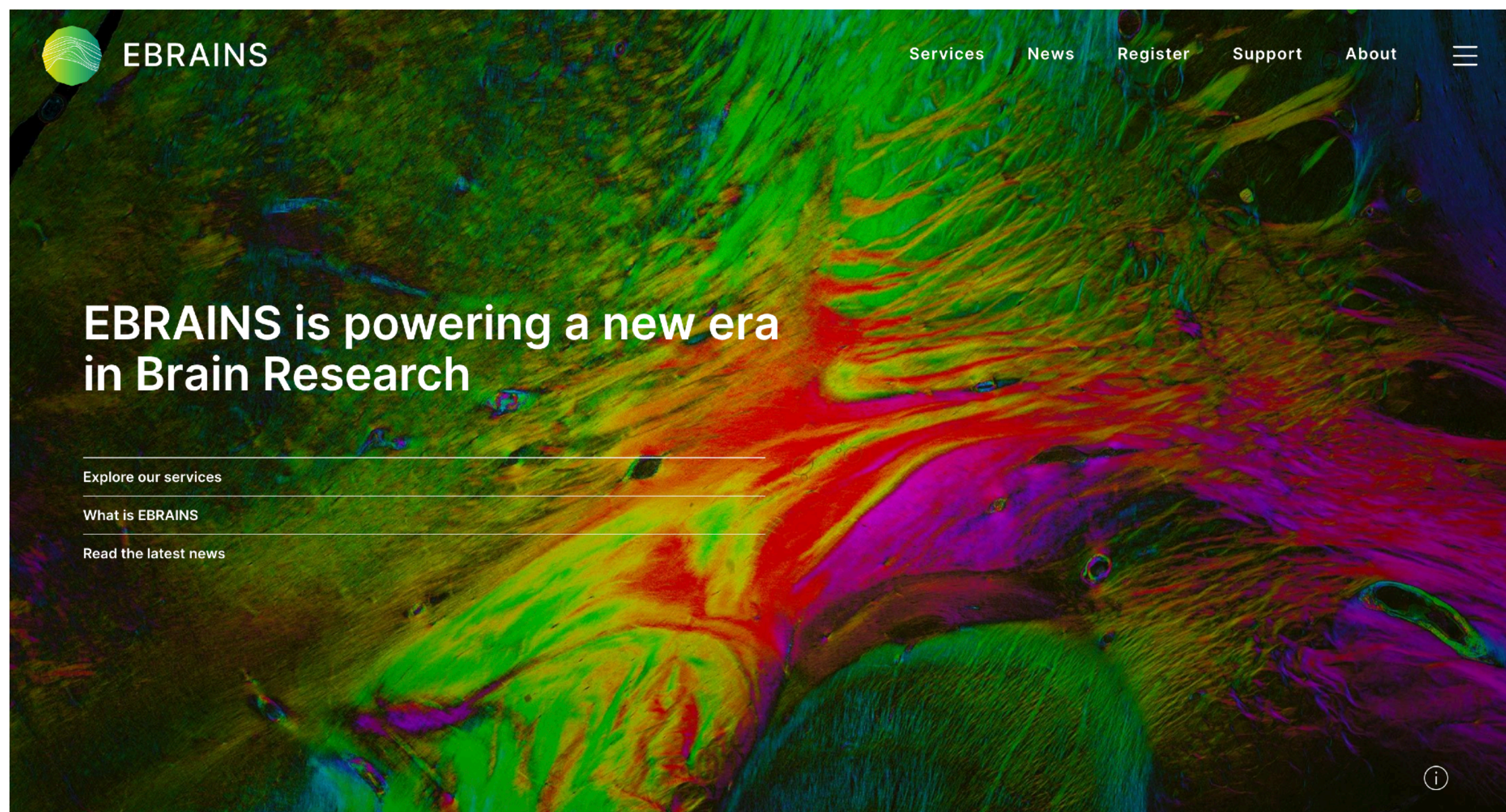
Outline

- Overview of the SpiNNaker and BrainScaleS neuromorphic systems
- How to use these neuromorphic systems
- How we're using the OpenStack infrastructure provided by ICEI to build and deploy our services
- Future plans
- Q&A

Target audience

- Neuroscientists
- HPC infrastructure users
- EBRAINS service developers
- Other platform service developers

Neuromorphic Computing in EBRAINS



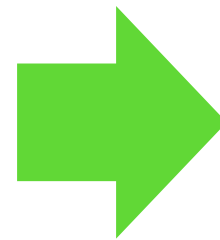
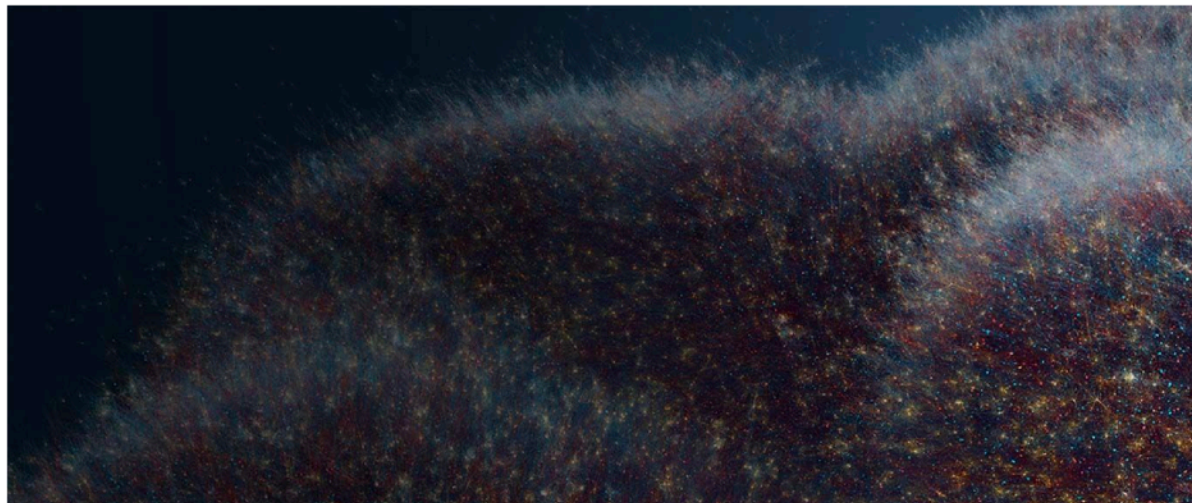
<https://ebrains.eu>

Neuromorphic Computing in EBRAINS



Register for your EBRAINS account today

<https://ebrains.eu/>



BRAIN-INSPIRED TECHNOLOGIES

Neuromorphic Computing

Simulate or emulate spiking neural networks with neuromorphic compute systems

<https://ebrains.eu/service/neuromorphic-computing>

Access BrainScaleS

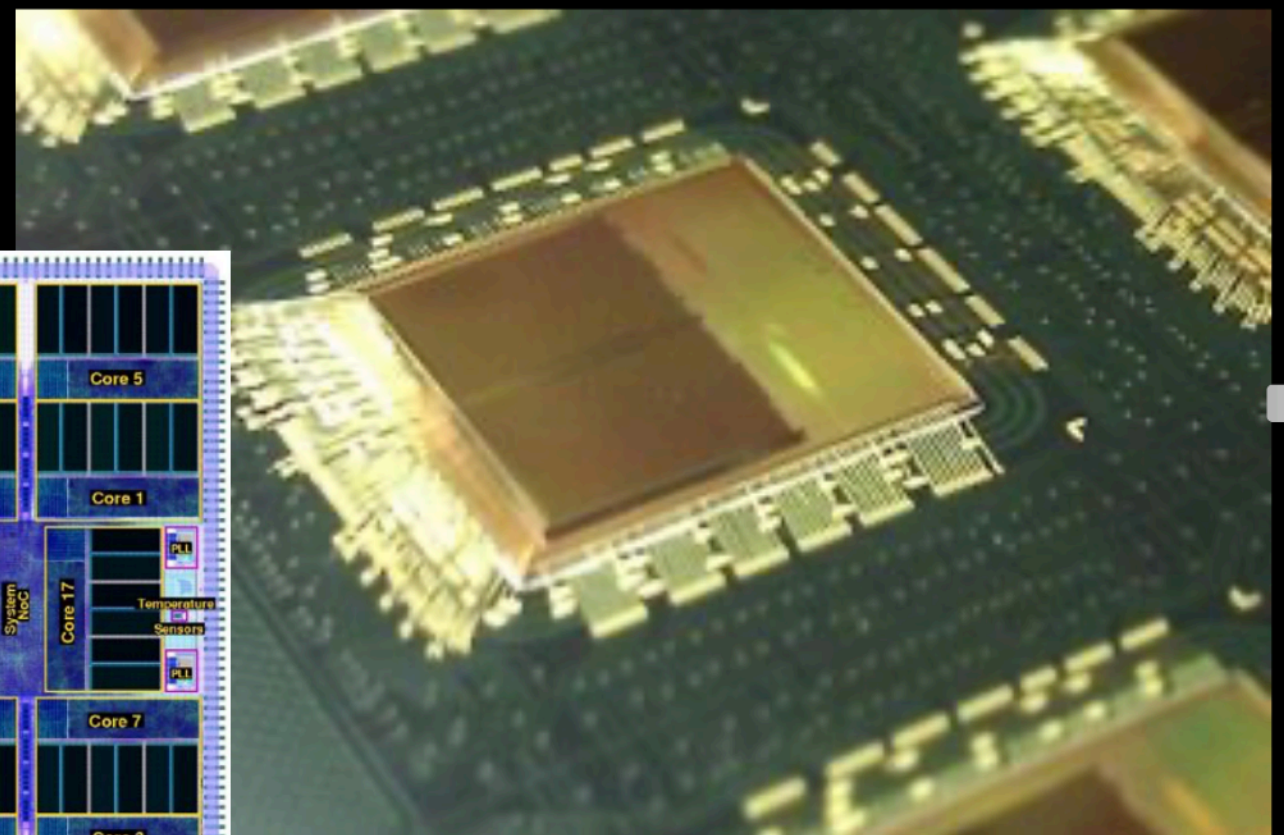
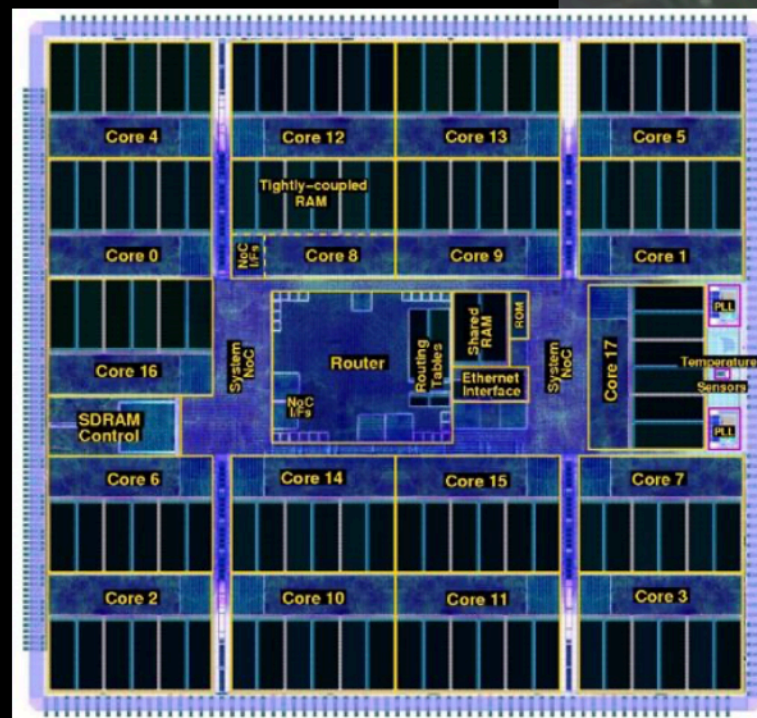
Access SpiNNaker



SpiNNaker

- custom-designed digital chips optimised for spiking neural network models

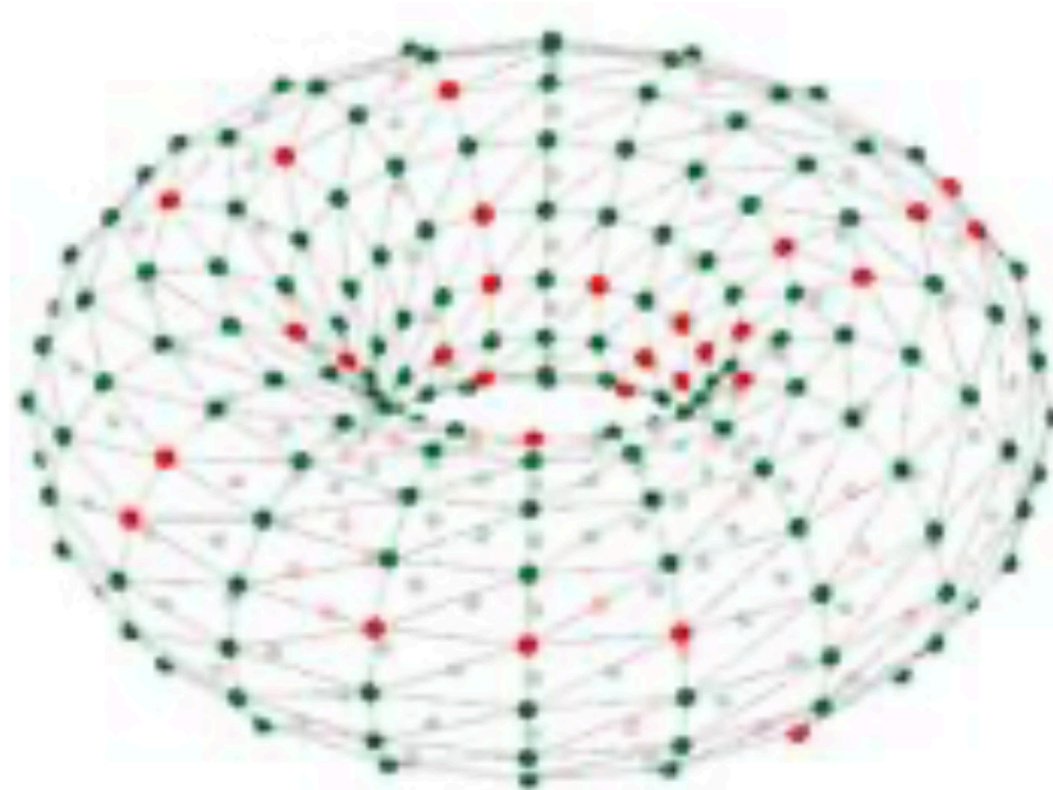
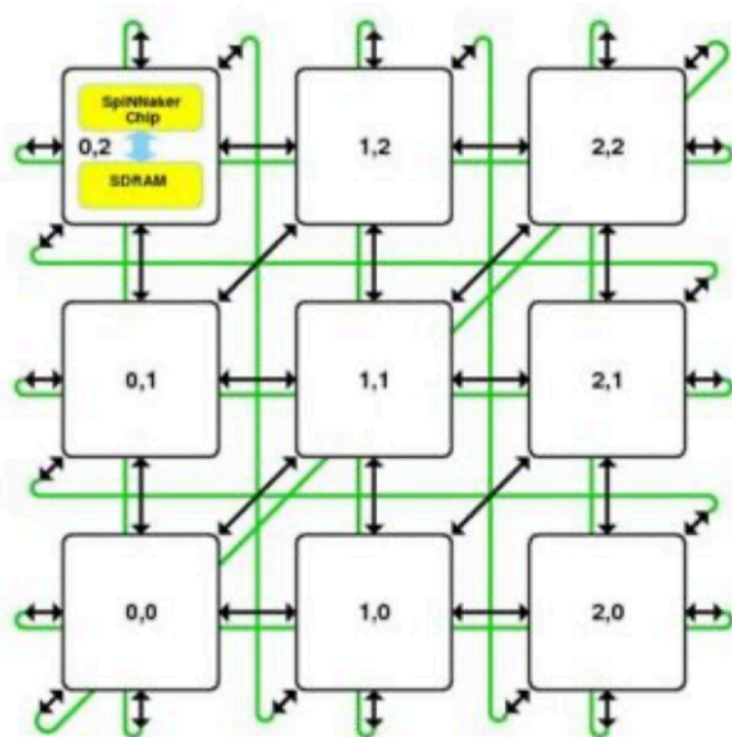
SpiNNaker Group (+ HBP)



- 18 *ARM 968* Cores per chip
- Integer Arithmetic
- 200 MHz Processor Clock
- Shared system RAM on die

- 128 Mbyte DRAM stacked on die
- Each Chip 6 bi-directional links
- 6 million spikes / s / link
- Real Time Simulator

SpiNNaker



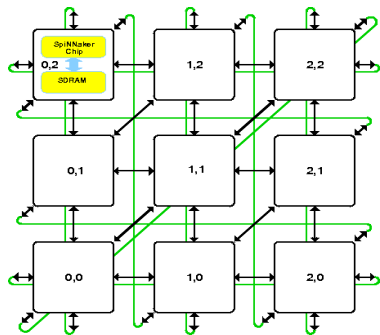
Connectivity

- Small packet based communication network
- Toroidal link geometry
- Maximum of 3 routings between any pair of nodes
- 6 Million spikes per second per bi-directional link

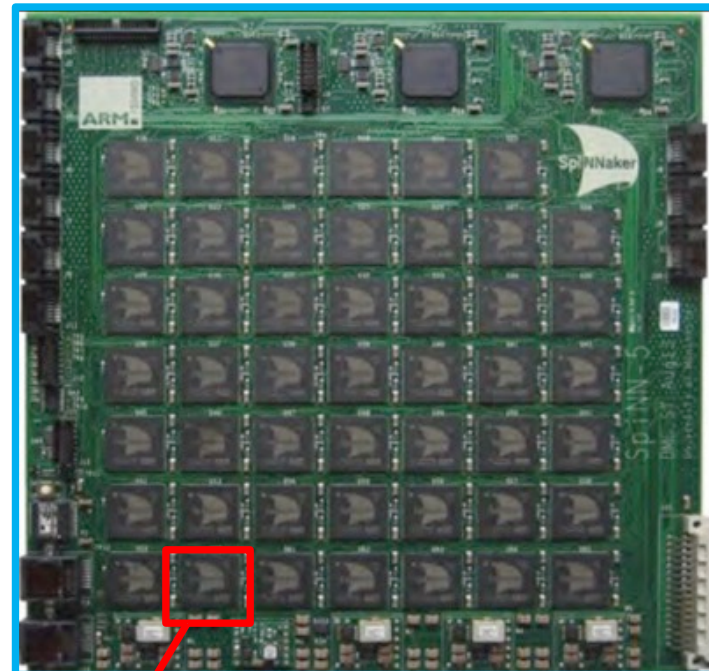
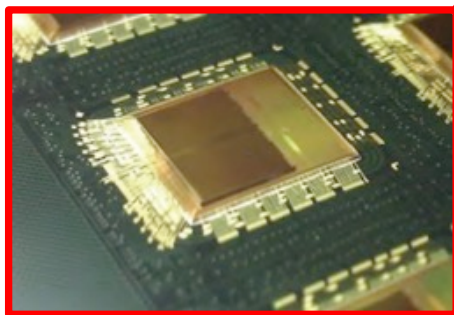


SpiNNaker

SpiNNaker board
(864 ARM cores)



SpiNNaker chip
(18 ARM cores)



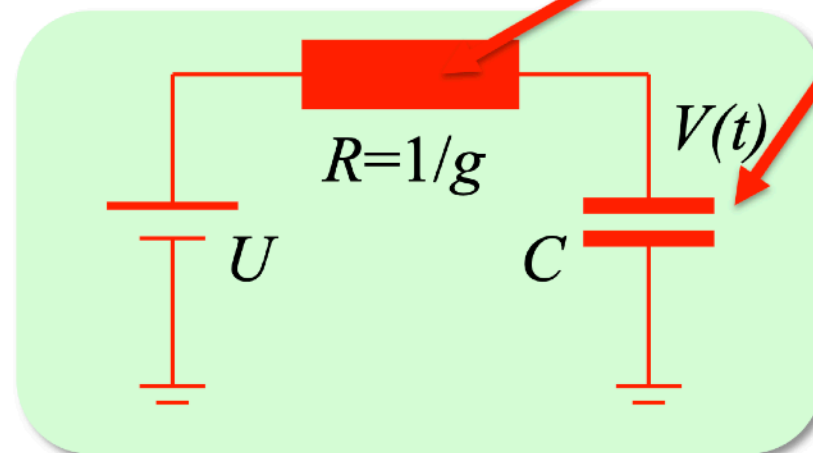
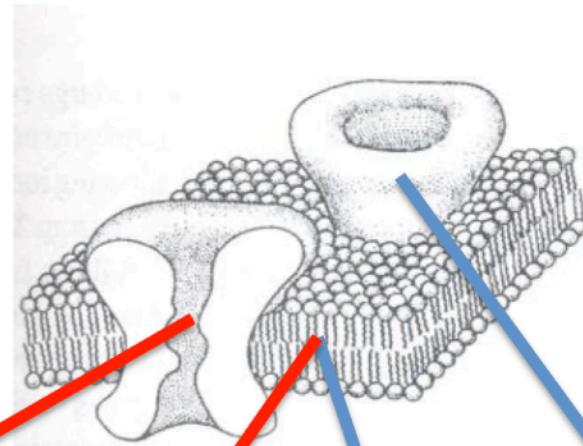
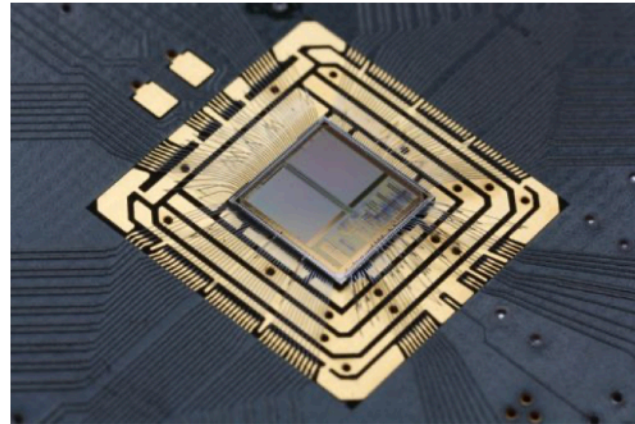
SpiNNaker racks
(1M ARM cores)

- HBP platform
 - 1M cores
 - 11 cabinets (including server)



BrainScaleS

- custom-designed analog chips for emulating spiking neurons



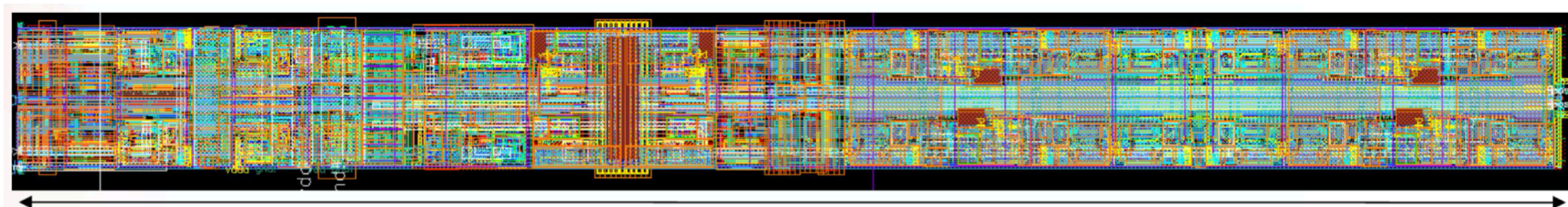
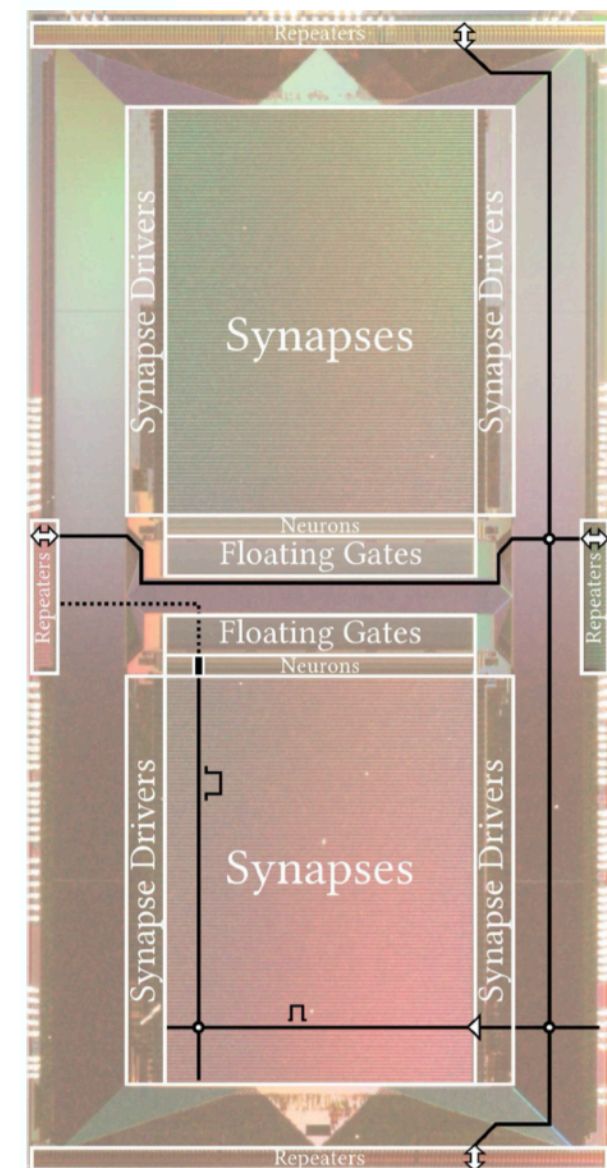
Physical Model

$$C \frac{dV(t)}{dt} = -g (V(t) - U)$$

Mathematical Model

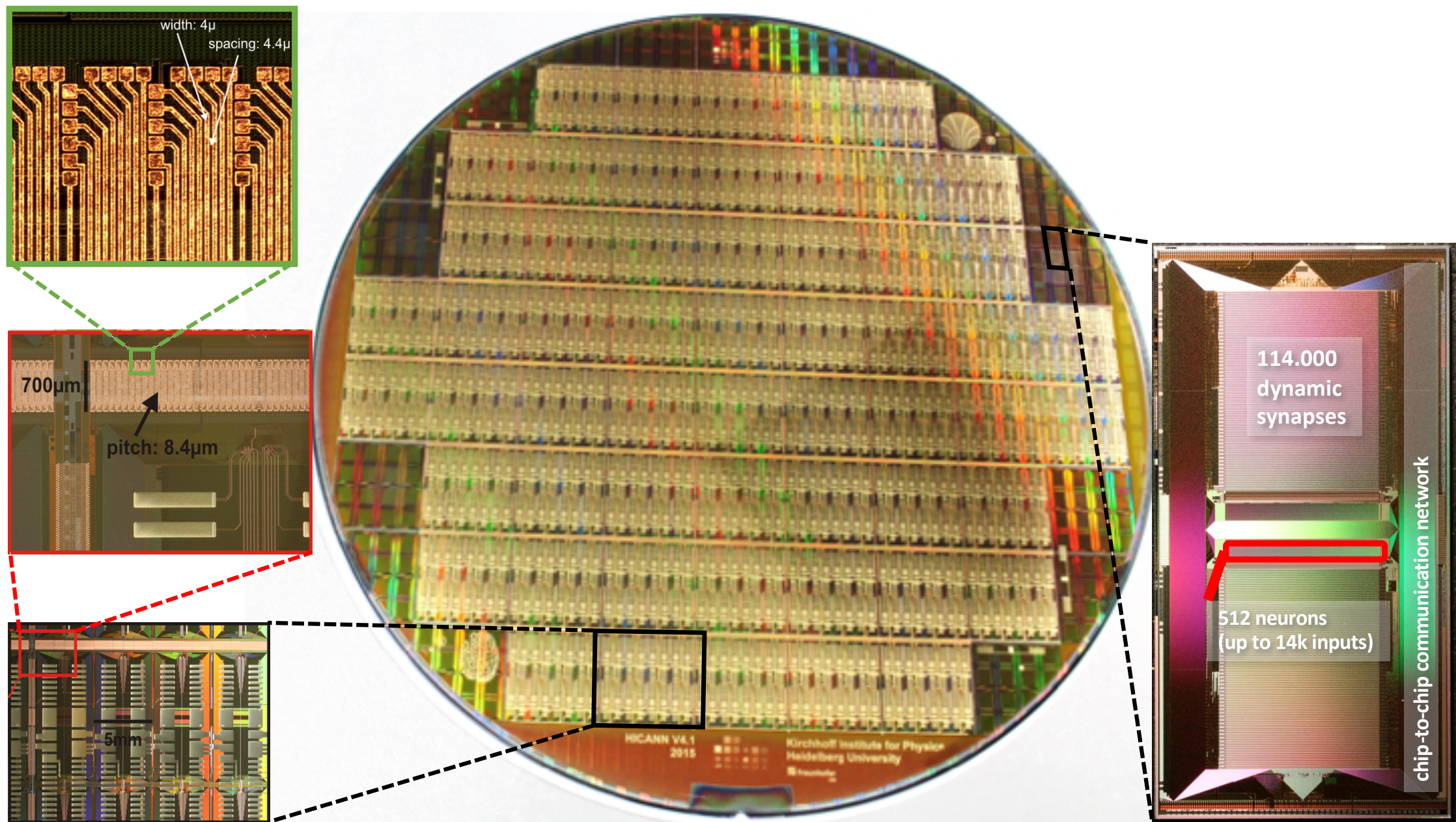
BrainScaleS

- Adaptive Exponential Integrate and Fire (AdEx) Model
- Accelerated dynamics compared to biological real-time (1000-10000 x)
- 512 analog neurons, 110000 plastic synapses
- Digital communication → mixed-signal system
- Sparse crossbar switches connecting buses → programmable network connections
- Analog parameter storage (floating gates)

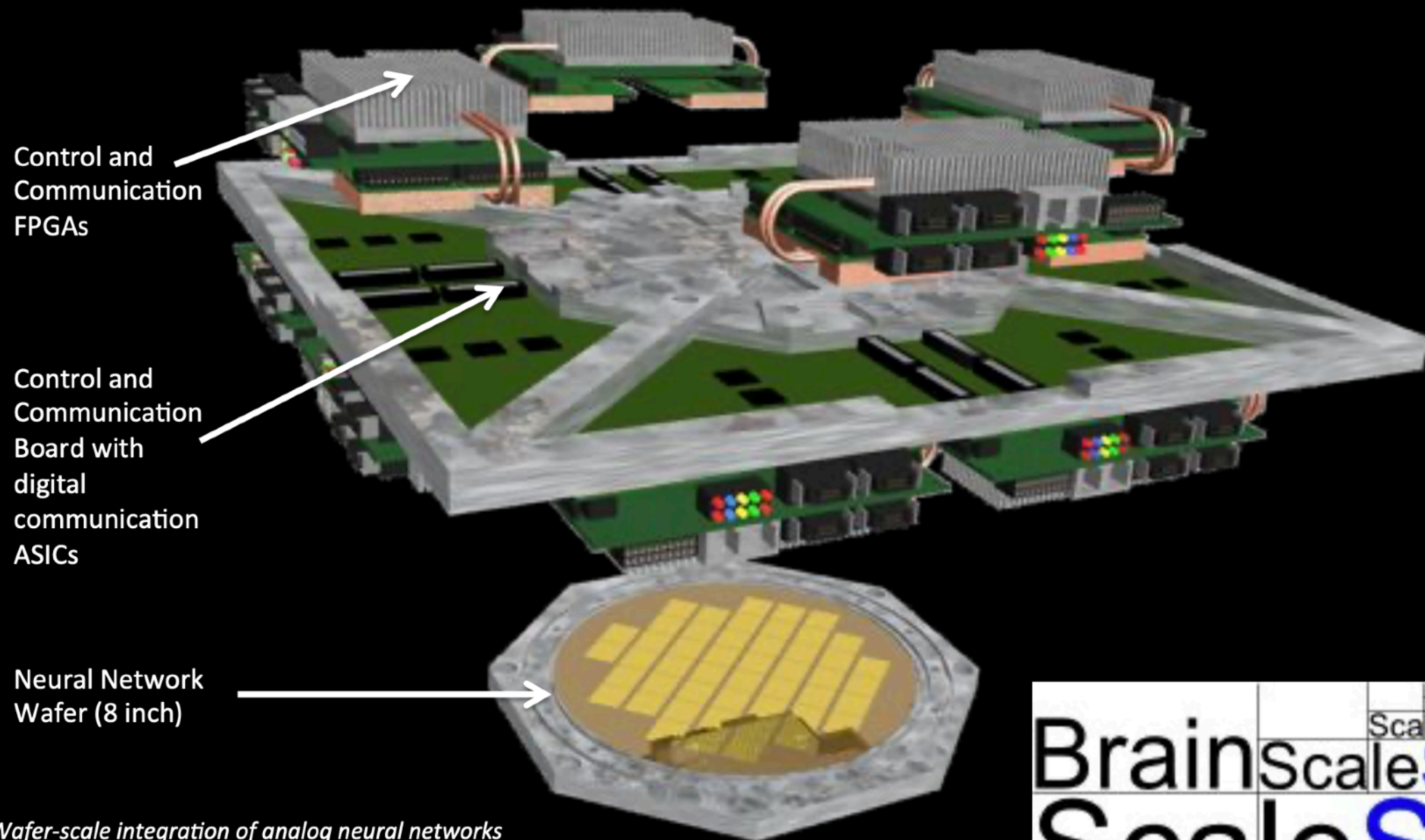


layout drawing of two neurons: 150x20 μm^2

BrainScaleS



Neural Processing Unit,
up to 200.000 Neurons, 50.000.000 plastic Synapses
Separation of Neural Circuits and Monitoring/Readout/Control



Control and
Communication
FPGAs

Control and
Communication
Board with
digital
communication
ASICs

Neural Network
Wafer (8 inch)

Wafer-scale integration of analog neural networks
J. Schemmel, J. Fieres and K. Meier
In : Proceedings of IJCNN (2008), IEEE Press, 431

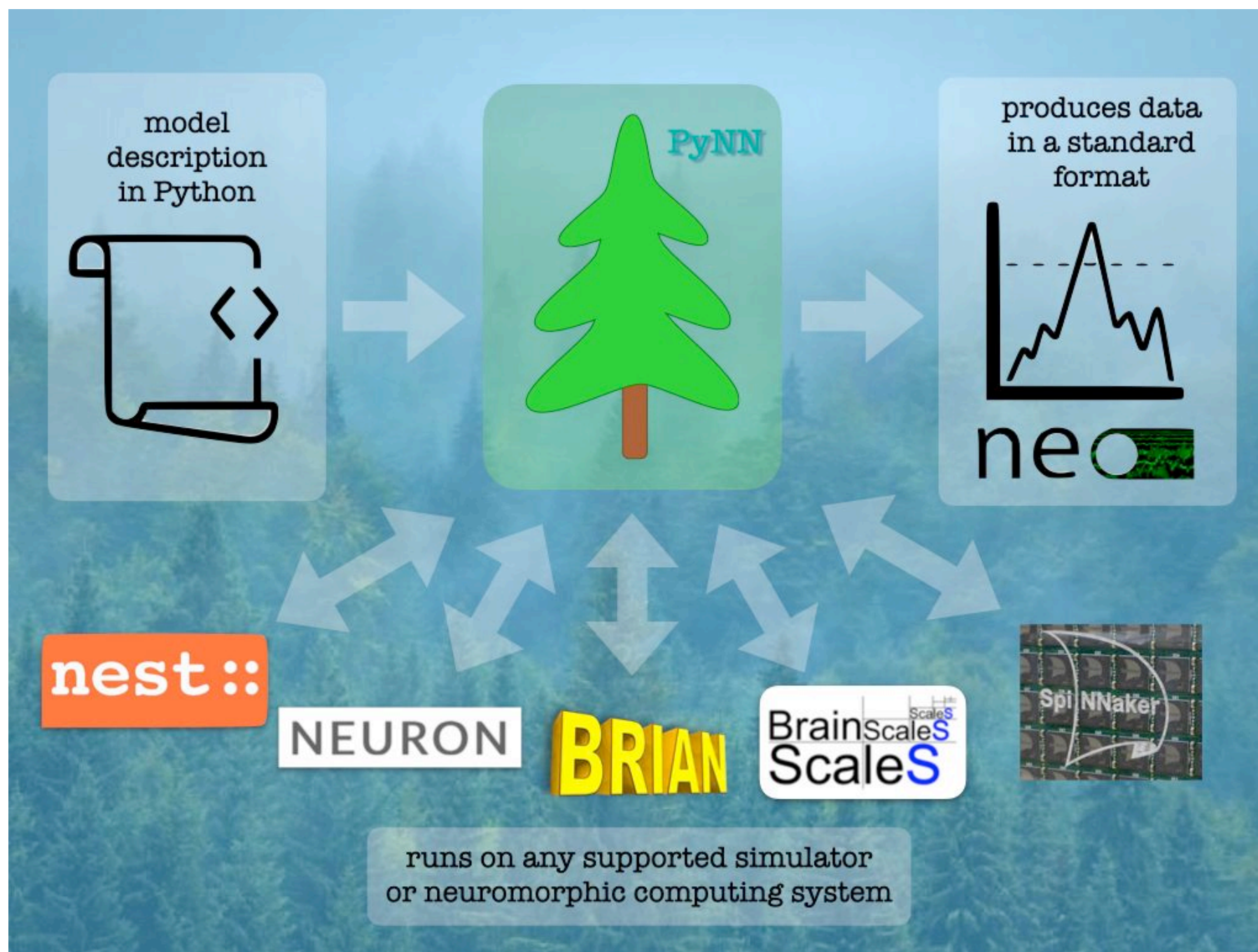
BrainScaleS
ScaleS



BrainScaleS
ScaleS

<https://ebrains.eu/service/neuromorphic-computing#BrainScaleS>

Programming neuromorphic computers



Programming neuromorphic computers

```
import pyNN.nest as simulator
import pyNN.neuron as simulator
import pyNN.brainscales as simulator
import pyNN.spiNNaker as simulator
```

```
cell_type = ...
```

```
p1 = simulator.Population(size1, cell_type, structure)
p2 = simulator.Population(size2, another_cell_type,
                           structure)
```

```
all = p1 + p2
all.record(['spikes', 'v'])
```

```
connections = simulator.Projection(
    p1, p2,
    connection_rule,
    synapse_type)
```

```
simulator.run(1000.0)
```



How to access SpiNNaker and BrainScaleS

- **Batch mode**
 - Web app
 - Python client (Jupyter notebooks)
 - Cmdline client
 - UNICORE
- **Interactive mode**
 - Jupyter notebooks

Job Manager app

EBRAINS Neuromorphic Computing Service: Job Manager

Jobs Quotas +

New job

Hardware Platform

Hardware
SpiNNaker

Please choose a simulation platform

Code

EDITOR FROM GIT REPOSITORY OR ZIP ARCHIVE FROM THE DRIVE GRAPHICAL MODEL BUILDER

```
1 import pyNN.spiNNaker as p
2 from pyNN.random import RandomDistribution
3 import matplotlib.pyplot as pylab
4 import numpy
5 from pyNN.utility.plotting import Figure, Panel
6 p.setup(timestep=0.1)
7 n_neurons = 100
8 n_exc = int(round(n_neurons * 0.8))
9 n_inh = int(round(n_neurons * 0.2))
10 weight_exc = 0.1
11 weight_inh = -5.0 * weight_exc
12 weight_input = 0.001
13 pop_input = p.Population(100, p.SpikeSourcePoisson(rate=0.0), label="Input")
14 pop_exc = p.Population(n_exc, p.IF_curr_exp, label="Excitatory",
15                       additional_parameters={"spikes_per_second": 100})
16 pop_inh = p.Population(n_inh, p.IF_curr_exp, label="Inhibitory",
17                       additional_parameters={"spikes_per_second": 100})
18 stim_exc = p.Population(
```

Command

Command:

run.py {system} --option1=42

Optional: specify the path to the main Python script, with any command-line arguments.

- Write Python code directly in the app, or load from a Git repository or the EBRAINS Drive

Hardware Configuration

Hardware config

```
{
  "spynnaker_version": "master",
  "spinnaker_tools_version": "3.1.0",
  "extra_pip_installs": ["elephant"],
  "extra_git_repositories": ["https://github.com/SpiNNakerManchester/SpiNNakerGraphFrontEnd"],
  "extra_makes": ["SpiNNakerGraphFrontEnd/spinnaker_graph_front_end/examples"],
  "extra_python_setups": ["SpiNNakerGraphFrontEnd"]
}
```

Please type a JSON-formatted object. See the Guidebook for more details

Tags

tags

Tag1,Tag2;This is Tag3



Please type job tags, separated by a comma, or semicolon. Tags can have spaces.

CANCEL SUBMIT

Job Manager app

EBRAINS Neuromorphic Computing Service: Job Manager

Jobs Quotas +

 	ID	 Status	 System	 Code	 Submitted on	 Submitted by
	153802	 submitted	SpiNNaker	https://github.com/msenoville/SpNN-test.git...	2021/04/09 10:24:30	msenoville
	153801	 queued	SpiNNaker	https://github.com/msenoville/SpNN-test.git...	2021/04/09 10:24:08	msenoville
	153799	 error	SpiNNaker	<code>import pyNN.spiNNaker as pynn import numpy as np ...</code>	2021/04/09 10:15:42	msenoville
	153798	 finished	SpiNNaker	<code>import pyNN.spiNNaker as pynn import numpy as np ...</code>	2021/04/09 10:15:17	msenoville
	153797	 finished	BrainScaleS	<code>import pyNN.spiNNaker as pynn import numpy as np ...</code>	2021/04/09 10:14:50	msenoville
	153796	 finished	SpiNNaker	<code>import pyNN.spiNNaker as pynn ...</code>	2021/04/09 10:10:33	msenoville
	153795	 finished	SpiNNaker	<code>import pyNN.spiNNaker as pynn import numpy as np ...</code>	2021/04/09 10:09:53	msenoville
	153794	 finished	BrainScaleS	<code>from neo.io import NeoMatlabIO import pynn_brainsc...</code>	2021/04/09 10:00:32	msenoville
	153793	 finished	BrainScaleS	<code>from neo.io import NeoMatlabIO import pynn_brainsc...</code>	2021/04/09 10:00:09	msenoville

Job Manager app

EBRAINS Neuromorphic Computing Service: Job Manager

Resource requests for 'neuromorphic-testing-private'

Title
testing Collaborato
testing Collaborato
Still testing Collabo
Still testing Collabo
Still testing Collabo
Testing Collaborato
yet another test of

+ NEW REQUEST

Editing

Title

Summary

Full description of project

Full description is not required for test quotas

CANCEL SAVE CHANGES SUBMIT REQUEST

Job Manager app

EBRAINS Neuromorphic Computing Service: Job Manager

JobsQuotas+

Job 153802

finished

Submitted on 2021/04/09 10:24:30 by msenoville to SpiNNaker
Completed on 2021/04/09 10:26:55

Output files

<http://spinnaker.cs.man.ac.uk/services/rest/output/incf-nmpi-2021/153802/reports.zip>

Code

<https://github.com/msenoville/SpNN-test.git>

Command

Hardware Config

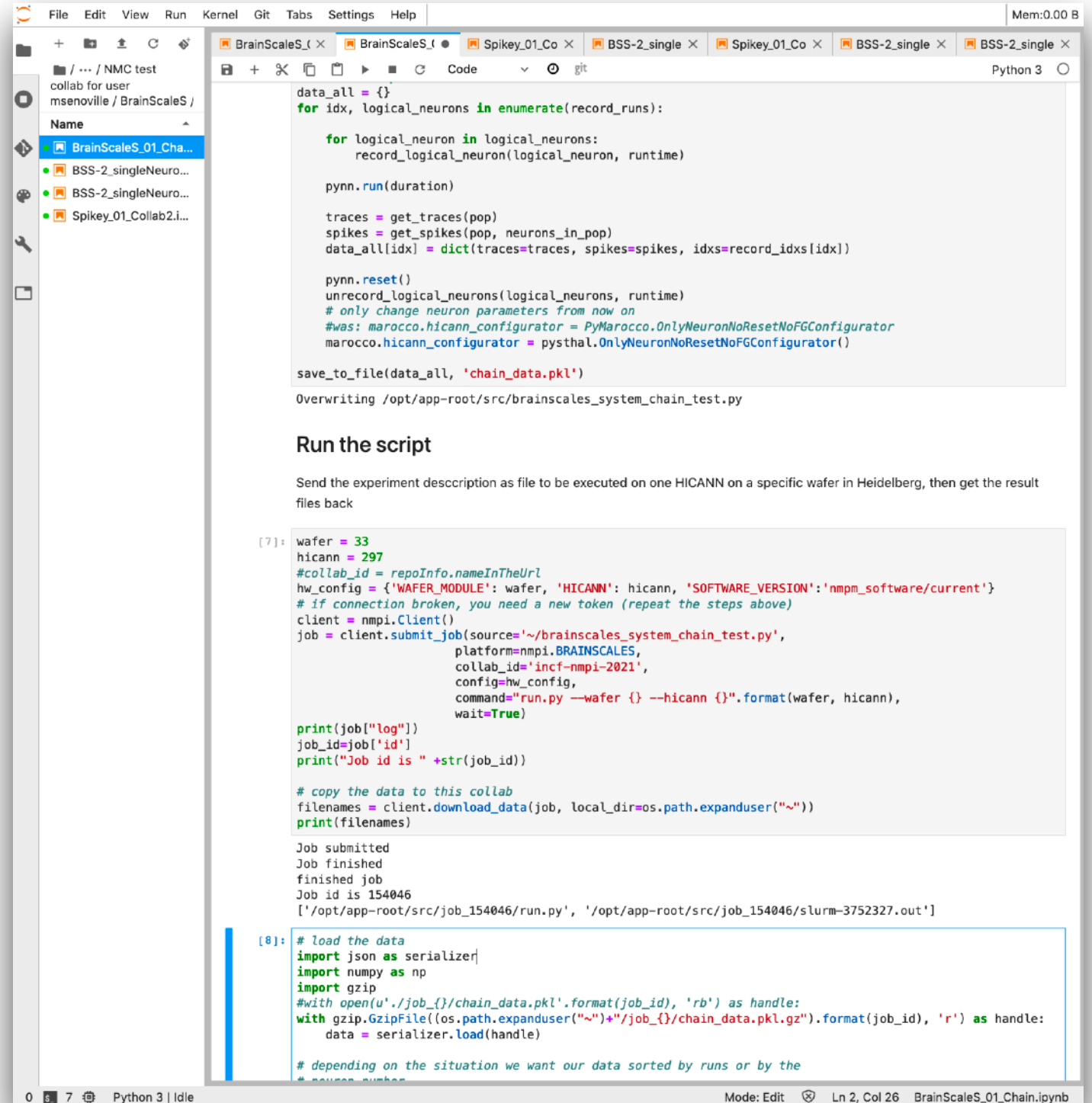
Provenance

Log

```
# [Machine]
# #spalloc_server=192.168.240.254
# spalloc_server=10.11.192.11
# spalloc_user=JupyterUser

[Reports]
reports_enabled = True
write_energy_report = TrueRunning from /tmp/job7208250740777422271/tmp/SpNN-test/chan...
```

Python client



```
data_all = {}
for idx, logical_neurons in enumerate(record_runs):
    for logical_neuron in logical_neurons:
        record_logical_neuron(logical_neuron, runtime)

pynn.run(duration)

traces = get_traces(pop)
spikes = get_spikes(pop, neurons_in_pop)
data_all[idx] = dict(traces=traces, spikes=spikes, idxs=record_idx[idx])

pynn.reset()
unrecord_logical_neurons(logical_neurons, runtime)
# only change neuron parameters from now on
# was: marocco.hicann_configurator = PyMarocco.OnlyNeuronNoResetNoFGConfigurator
marocco.hicann_configurator = pynthal.OnlyNeuronNoResetNoFGConfigurator()

save_to_file(data_all, 'chain_data.pkl')

Overwriting /opt/app-root/src/brainscales_system_chain_test.py

Run the script

Send the experiment description as file to be executed on one HICANN on a specific wafer in Heidelberg, then get the result files back

[7]: wafer = 33
      hicann = 297
      #collab_id = repoInfo.nameInTheUrl
      hw_config = {'WAFER_MODULE': wafer, 'HICANN': hicann, 'SOFTWARE_VERSION': 'nmpm_software/current'}
      # if connection broken, you need a new token (repeat the steps above)
      client = nmpi.Client()
      job = client.submit_job(source='~/brainscales_system_chain_test.py',
                             platform=nmpi.BRAINSCALES,
                             collab_id='incf-nmpi-2021',
                             config=hw_config,
                             command="run.py --wafer {} --hicann {}".format(wafer, hicann),
                             wait=True)

      print(job["log"])
      job_id=job['id']
      print("Job id is " +str(job_id))

      # copy the data to this collab
      filenames = client.download_data(job, local_dir=os.path.expanduser("~"))
      print(filenames)

      Job submitted
      Job finished
      finished job
      Job id is 154046
      ['/opt/app-root/src/job_154046/run.py', '/opt/app-root/src/job_154046/slurm-3752327.out']

[8]: # load the data
      import json as serializer
      import numpy as np
      import gzip
      #with open(u'./job_{}/chain_data.pkl'.format(job_id), 'rb') as handle:
      with gzip.GzipFile((os.path.expanduser("~")+"/"+job_{}/chain_data.pkl.gz").format(job_id), 'r') as handle:
          data = serializer.load(handle)

      # depending on the situation we want our data sorted by runs or by the
      # neuron number
```

`pip install hbp-neuromorphic-platform`

Command-line client

- provided with the Python client
- develop code locally, submit to remote NMC system, results downloaded to local directory

```
$ nmpi run --help
Usage: nmpi run [OPTIONS] SCRIPT
```

Run a simulation/emulation

Options:

-p, --platform TEXT	Spinnaker, BrainScaleS, BrainScaleS-2, or Spikey
-t, --tag TEXT	Add a tag to the job
-b, --batch	Submit job then return immediately
-o, --output-dir TEXT	Output directory
-e, --server-env TEXT	
--help	Show this message and exit.

UNICORE

- in development for BrainScaleS
- submit a neuromorphic computing job in the same way as any other HPC job

<https://sourceforge.net/p/unicore/>

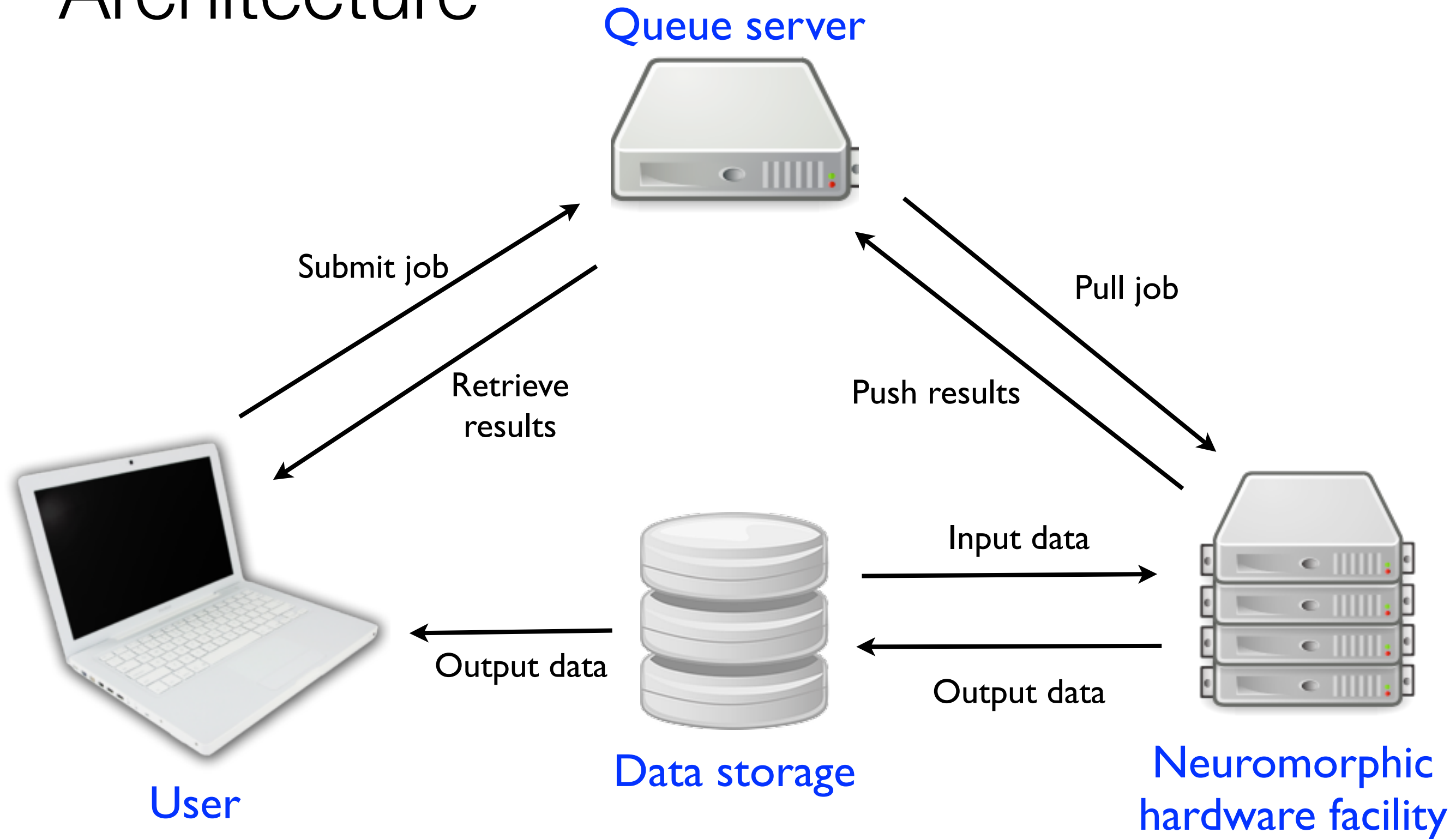
Interactive access



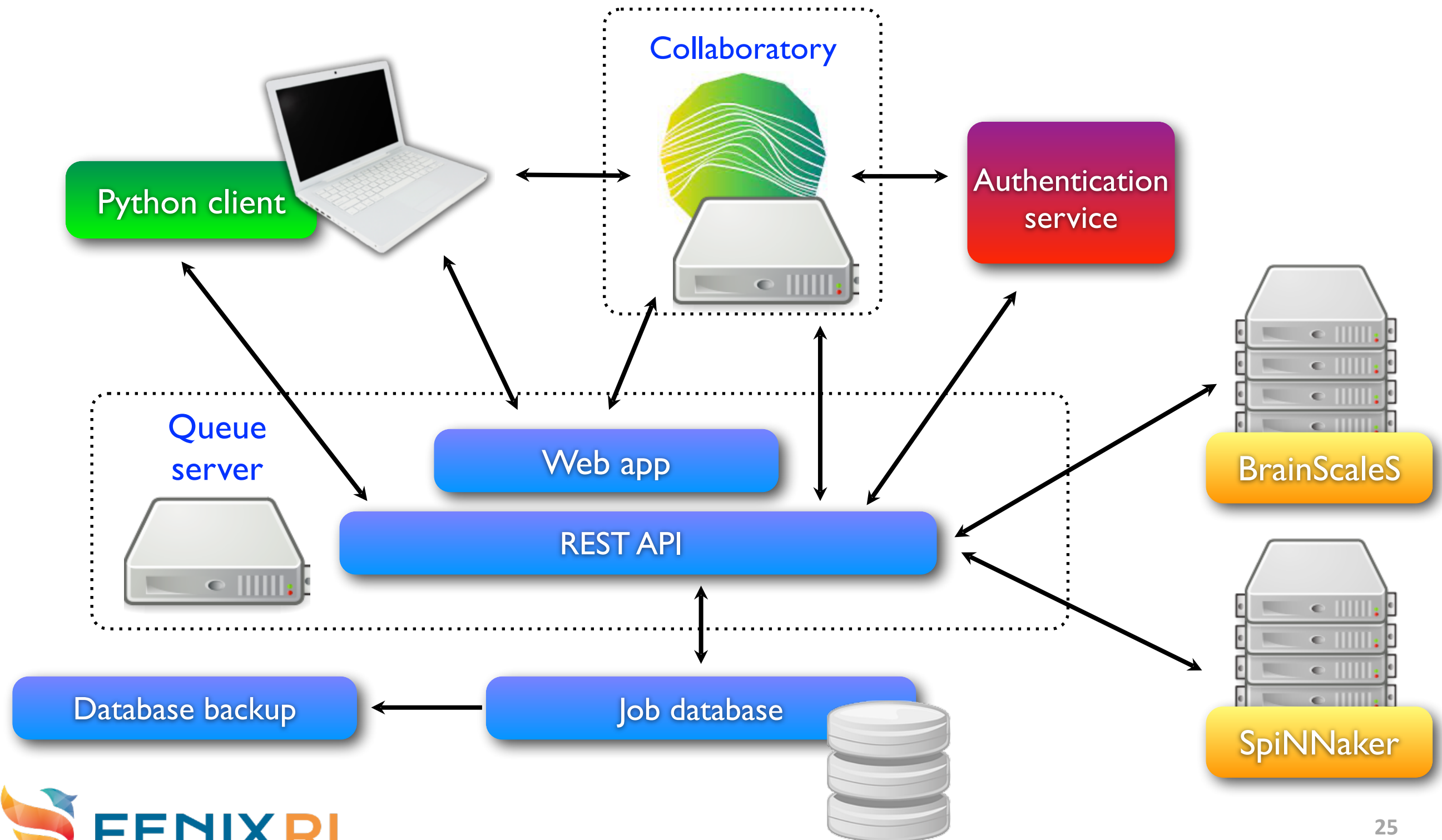
- Jupyter notebooks running live in Manchester, giving interactive access to SpiNNaker
- PyNN commands are executed directly, allowing exploration and iterative development (*cf* submitting an entire PyNN script as a batch job)

<https://spinn-20.cs.man.ac.uk/>

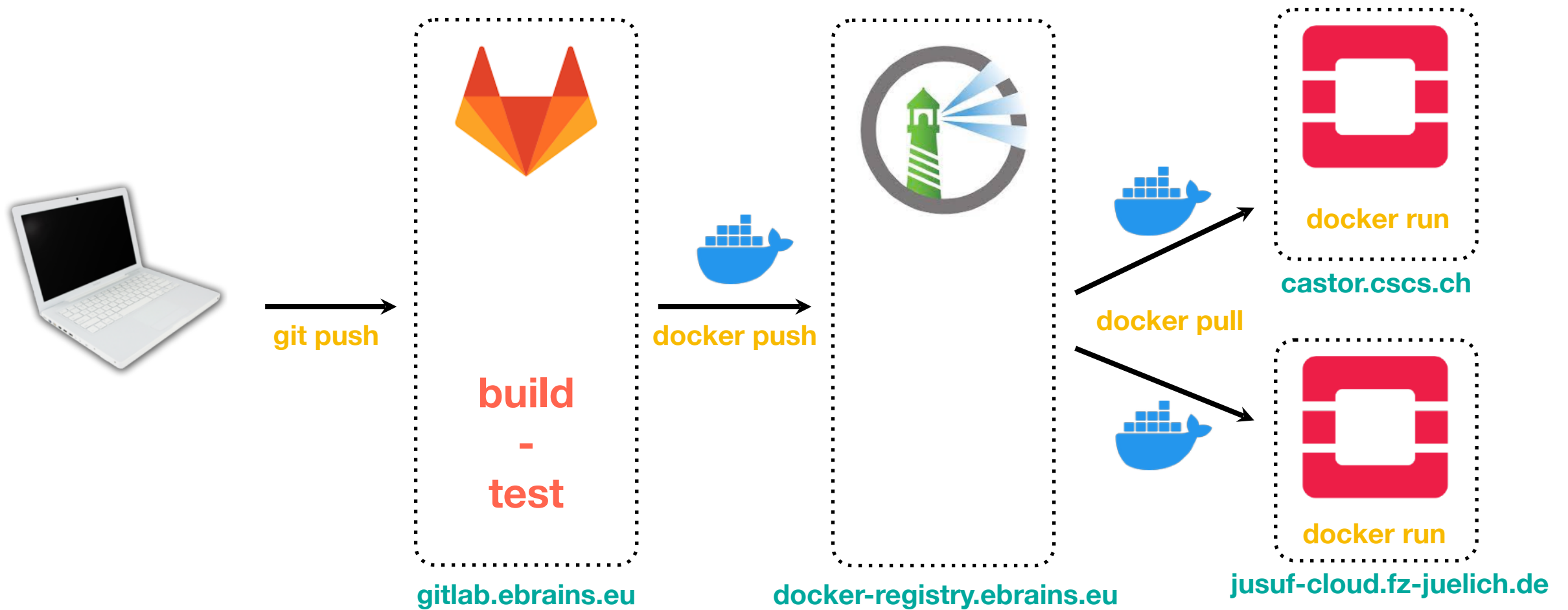
Architecture



Architecture

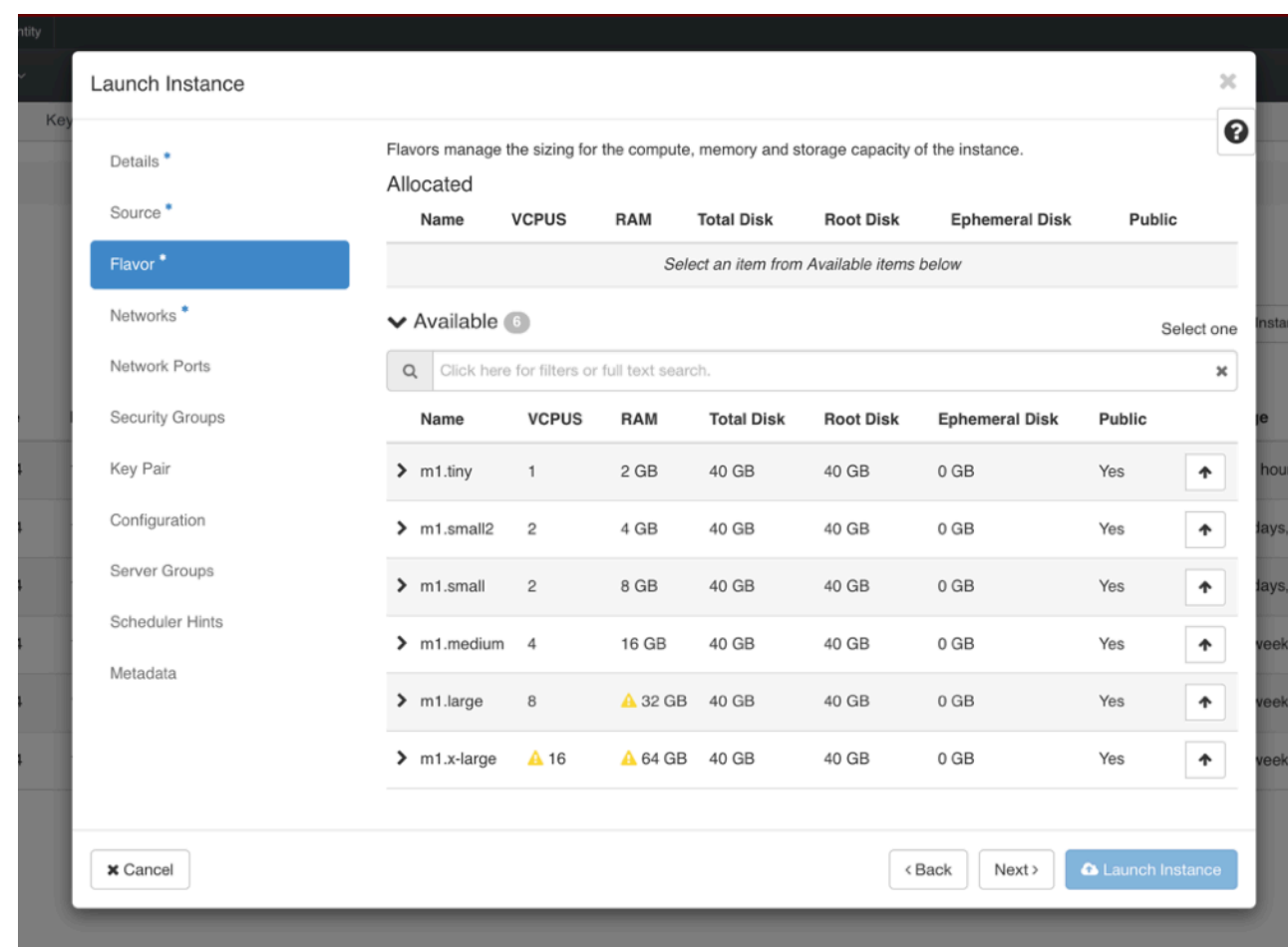


Deployment



Castor, Pollux and Jusuf Cloud

- Production and staging servers running on Fenix VM Services at CSCS (Castor/Pollux)
- Backup production servers running on Fenix VM Services at Jülich (JUSUF Cloud)
- Resources obtained via ICEI project
- VMs and volume storage are launched and configured manually via OpenStack dashboard and ssh



Future plans / in progress

- **increase automation of VM creation and Docker deployment using OpenStack API/ command line and Ansible**
- **automated failover between Castor and JUSUF**
- **centralised logging / monitoring**

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