



Training for Brilliant Minds

# Machine Learning at CSC

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# Overview

What CSC service to use?

Puhti supercomputer

Data storage

GPU utilization

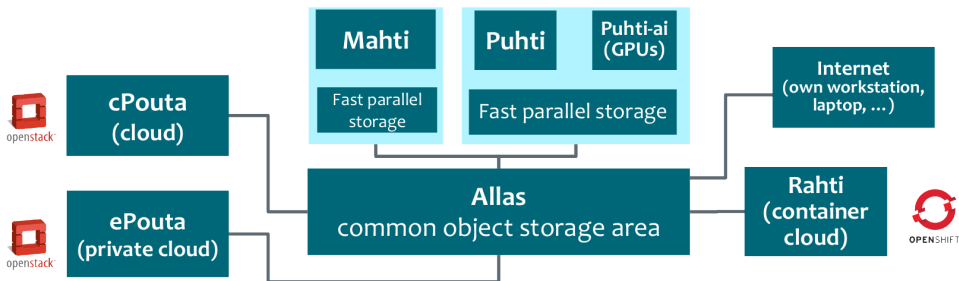
Multi-GPU and multi-node jobs

Singularity containers

What CSC service to use?

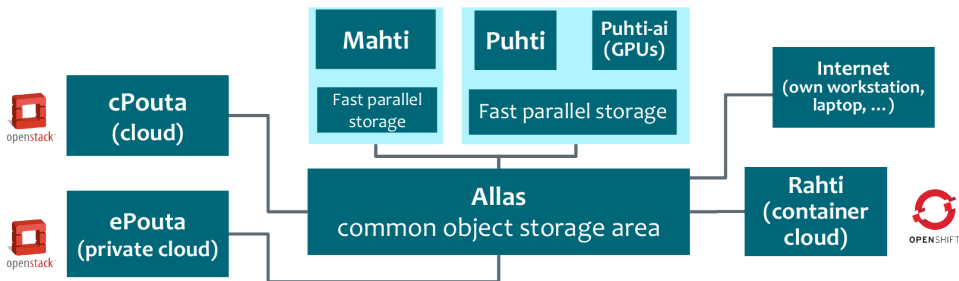
# What CSC service to use?

- CSC's supercomputer **Puhti**
- Virtual server on **Pouta**
- Container cloud **Rahti**



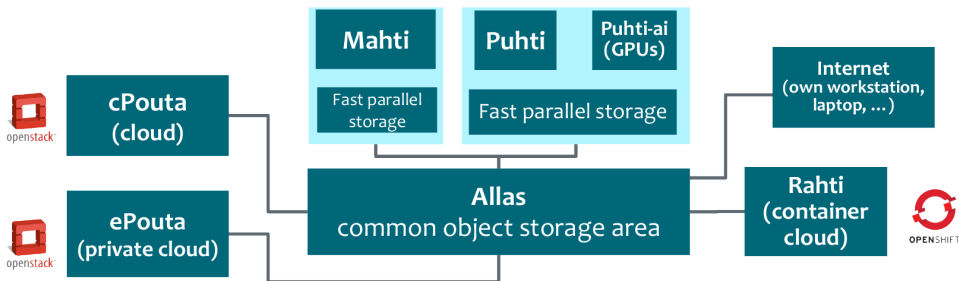
# What CSC service to use?

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  - Cluster with GPU-accelerated nodes
  - Multi-user environment
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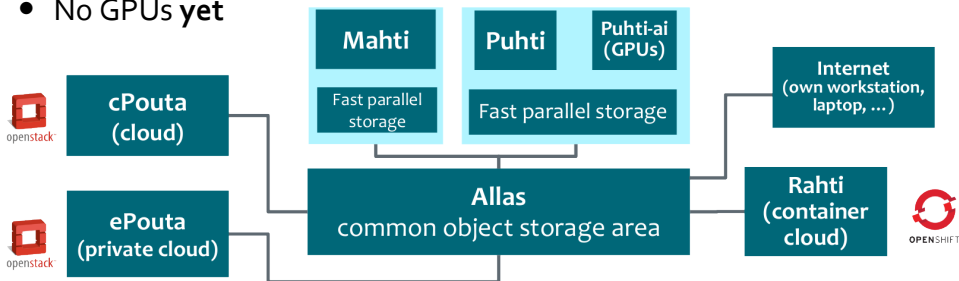
# What CSC service to use?

- CSC's supercomputer **Puhti**
  - Cluster with GPU-accelerated nodes
  - Multi-user environment
- Virtual server on **Pouta**
  - Your "own" server
  - Less powerful than Puhti
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# What CSC service to use?

- CSC's supercomputer **Puhti**
  - Cluster with GPU-accelerated nodes
  - Multi-user environment
- Virtual server on **Pouta**
  - Your "own" server
  - Less powerful than Puhti
- Container cloud **Rahti**
  - Easy to run containers
  - No GPUs **yet**

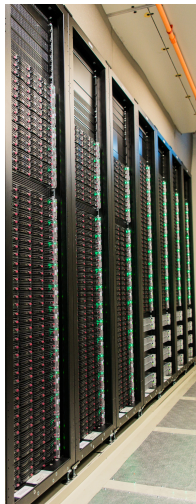


Puhti supercomputer



# Puhti supercomputer

- *Puhti-AI*, cluster with 80 nodes with 4 GPUs each  $\rightarrow$  320 GPUs in total
- Latest generation Nvidia V100 GPUs (Volta) with 32 GB of memory
- Fast network:  $2 \times 100$  Gbps links to each node
- Each node has a fast 3.2 TB local NVME disk



# Getting access to Puhti

<https://docs.csc.fi/computing/overview/>

To use Puhti you need to:

- Have a CSC account
- Be member of a CSC project, either by
  - creating a new project, or
  - joining an existing project (ask the PI to add you!)
- Finally, the project needs to have Puhti access

→ MyCSC portal: <https://my.csc.fi/>

# Accessing Puhti

- Using an ssh client such as OpenSSH or PuTTY
- Basic Linux skills are required!
- More info: <https://docs.csc.fi/computing/connecting/>

```
$ ssh <csc_username>@puhti.csc.fi
```

```
$ ssh <csc_username>@puhti-login2.csc.fi
```

# Supported frameworks

We currently support:

- **Python Data** – collection of Python libraries for data analytics and machine learning
- **TensorFlow** – deep learning library for Python
- **PyTorch** – machine learning framework for Python
- **MXNet** – deep learning library for Python
- **RAPIDS** – suite of libraries for data analytics and machine learning on GPUs

<https://docs.csc.fi/apps/#data-analytics-and-machine-learning>

# Example: TensorFlow

- First check the application page for instructions:  
<https://docs.csc.fi/apps/tensorflow/>
- Load the default version:  
`module load tensorflow`
- or specific version:  
`module load tensorflow/2.0.0`
- **Note:** some modules are *Singularity-based*!

# What if some package is missing?

If you are using our module, but a trivial package is missing ...

- install it yourself, e.g.,  
`pip install --user <packagename>`
- ...or if it might be generally useful, send an email to [servicedesk@csc.fi](mailto:servicedesk@csc.fi) – we can install it for you!

# What if some package is missing?

If you need a specific setup, and our modules are not right for you ...

- use a virtualenv:

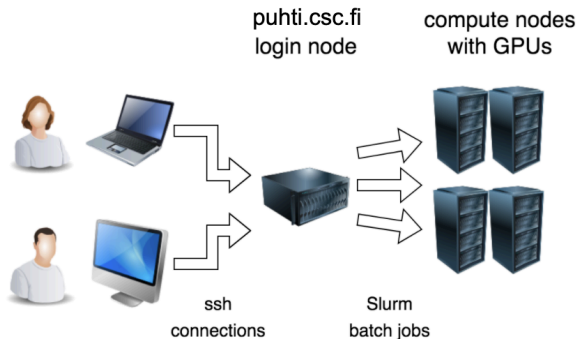
```
$ python3 -m venv myenv  
$ source myenv/bin/activate  
$ pip install ...
```

- use conda: <https://docs.csc.fi/support/tutorials/conda/>
- use singularity containers:  
<https://docs.csc.fi/computing/containers/run-existing/>
- or if generally useful, send an email to [servicedesk@csc.fi](mailto:servicedesk@csc.fi)

# Running a job on Puhti

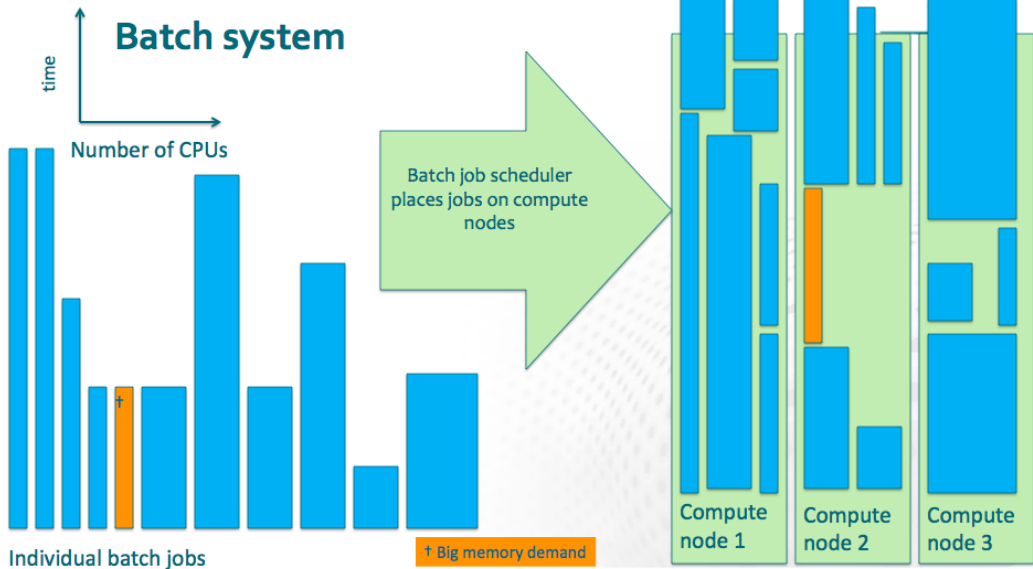
Don't run heavy computing jobs in the login nodes!

- Puhti uses the *Slurm* batch job system
- Jobs do not run instantly but are put in a *queue*
- Resources (runtime, memory, number of cores) need to be specified





# Running a job on Puhti



# Running a job on Puhti

Create a job script, for example `run.sh`:

```
#!/bin/bash
#SBATCH --account=<project>
#SBATCH --partition=gpu
#SBATCH --ntasks=1
#SBATCH --cpus-per-task=10
#SBATCH --mem=64G
#SBATCH --time=1:00:00
#SBATCH --gres=gpu:v100:1

module load tensorflow/2.0.0
srun python3 myprog.py <options>
```

<https://docs.csc.fi/computing/running/creating-job-scripts/>

# Running a job on Puhti

Example job script for Singularity-based modules:

```
#!/bin/bash  
#SBATCH --account=<project>  
#SBATCH --partition=gpu  
#SBATCH --ntasks=1  
#SBATCH --cpus-per-task=10  
#SBATCH --mem=64G  
#SBATCH --time=1:00:00  
#SBATCH --gres=gpu:v100:1
```

```
module load tensorflow/nvidia-20.03-tf2-py3
```

```
srun singularity_wrapper exec python3 myprog.py <options>
```

# Running a job on Puhti

Submit the job:

```
sbatch run.sh
```

Check the queue:

```
squeue -l -u $USER
```

Cancel a job:

```
scancel <jobid>
```

<https://docs.csc.fi/computing/running/submitting-jobs/>

Data storage

# Data storage on Puhti

- Disk space and *number of files* are limited on Puhti!  
→ We want to ensure that the shared (Lustre) filesystem works efficiently for everyone!
- Useful command: `csc-workspaces`

	Owner	Path	Capacity	Number of files	Cleaning
home	Personal	/users/<user-name>	10 GiB	100 000 files	No
projappl	Project	/projappl/<project>	50 GiB	100 000 files	No
scratch	Project	/scratch/<project>	1 TiB	1 000 000 files	Yes - 90 days

Data quotas can be increased via MyCSC!

<https://docs.csc.fi/computing/disk/>

# Using Allas

- store big datasets in Allas, CSC's object storage
- download them to project scratch prior to computation
- you can also upload trained models (or keep in projappl)

```
$ module load allas
$ allas-conf
$ cd /scratch/<your-project>
$ swift download <bucket-name> your-dataset.tar
```

# Large number of files

- Many datasets contain a large number of small files
- Shared filesystem (Lustre) performs poorly in this scenario  
→ noticable slowdowns for all Puhti users!

Consider alternatives:

- packaging your dataset into larger files
- use NVME fast local storage on GPU nodes



# Using more efficient data formats

Instead of many small files, use one or a few bigger files.

Examples:

- TensorFlow's TFRecord format
- HDF5
- LMDB
- ZIP, for example via Python's `zipfile` library

# Fast local NVME drive

- All GPU nodes have a local NVME drive
- Just add `nvme:<number-of-GB>` to sbatch `--gres` flag

```
#!/bin/bash
#SBATCH --account=<project>
#SBATCH --partition=gpu
#SBATCH --ntasks=1
#SBATCH --cpus-per-task=10
#SBATCH --mem=64G
#SBATCH --time=1:00:00
#SBATCH --gres=gpu:v100:1,nvme:100
```

```
tar xf /scratch/<your-project>/your-dataset.tar -C $LOCAL_SCRATCH
```

```
srun python3 myprog.py --data_dir=$LOCAL_SCRATCH <options>
```

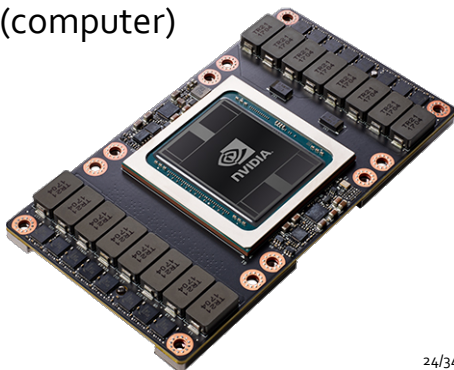
GPU utilization

# GPU utilization

GPUs are an expensive resource compared to CPUs ( $\times 60$  BUs!)  
 → GPU should be maximally utilized!

For a running job:

- use `queue` to find out on what node (computer) it is running
- `ssh` into that node, e.g., `ssh r01g01`
- find the process id of your job with `ps -u $USER`
- run `nvidia-smi`



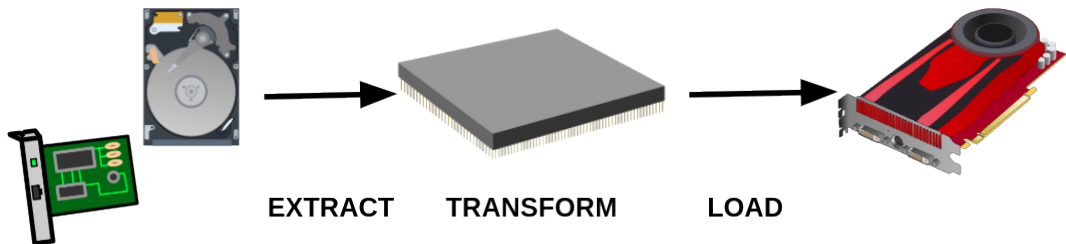
# GPU utilization

For a finished job:

- `run gpuseff <jobid>`
- shows GPU utilisation statistics for the whole running time
- **note:** gpuseff is currently in testing usage, and still under development

You can always contact our service desk if you need advice on how to improve your GPU utilization!

# Using multiple CPUs for ETL



- Common bottle-neck: CPU cannot keep up with GPU
- GPU has to wait for more data ...
- Solution: use more CPUs (they are “cheaper” than GPUs!)

A good **rule of thumb in Puhti** is to reserve 10 CPUs per GPU (as there are 4 GPUs and 40 CPUs per node).

# Using multiple CPUs for ETL

In Slurm scripts:

```
#SBATCH --cpus-per-task=10
```

**Note:** using multiple CPUs *not* automatic, code needs to support it!

For example in TensorFlow:

```
dataset = dataset.map(..., num_parallel_calls=10)
dataset = dataset.prefetch(buffer_size)
```

PyTorch:

```
train_loader = torch.utils.data.DataLoader(..., num_workers=10)
```

Multi-GPU and multi-node jobs



# Multi-GPU

Many frameworks support multi-GPU within a single node.

Slurm script:

```
#SBATCH --gres=gpu:v100:4
```

TensorFlow:

```
mirrored_strategy = \
    tf.distribute.MirroredStrategy()
with mirrored_strategy.scope():
    model = Sequential(...)
    model.add(...)
    model.add(...)
    model.compile(...)
```

PyTorch:

```
model = MyModel(...)
if torch.cuda.device_count() > 1:
    model = nn.DataParallel(model)
model.to(device)
```

# Multi-GPU and multi-node

- A single node has 4 GPUs
- If you need more than 4 GPUs, we recommend [Horovod](#)
- Supported for TensorFlow and PyTorch on Puhti
- Uses MPI and NCCL for interprocess communication
- Modules with `-hvd` suffix

Try:

```
module avail hvd
module load tensorflow/2.0.0-hvd
```

# Slurm example for Horovod

Example slurm script that uses **8 GPUs** across **two computers**

- MPI terminology: 8 tasks, 2 nodes
- Each task is 1 GPU and 10 CPUs

```
#!/bin/bash
#SBATCH --account=<project>
#SBATCH --partition=gpu
#SBATCH --ntasks=8
#SBATCH --nodes=2
#SBATCH --cpus-per-task=10
#SBATCH --mem=32G
#SBATCH --time=1:00:00
#SBATCH --gres=gpu:v100:4
```

```
srun python3 myprog.py <options>
```

# Singularity containers

# Singularity on Puhti

- Puhti supports Singularity-based containers
- Some of our modules use it – just remember to prefix commands with `singularity_wrapper exec`
- You can also convert your own Docker containers, see:  
<https://docs.csc.fi/computing/containers/run-existing/>

Conversion (preferably not in login nodes!):

```
$ singularity build pytorch_20.03-py3.sif \  
    docker://nvcv.io/nvidia/pytorch:20.03-py3
```

Extract from Slurm script:

```
srun singularity_wrapper exec --nv pytorch_20.03-py3.sif \  
    python3 myprog <options>
```

# Special Singularity-based applications

Specialized Singularity-based applications not shown with default modules

**Example:** Turku neural parser

```
$ module use /appl/soft/ai/singularity/modulefiles/  
$ module load turku-neural-parser/fi-en-sv-gpu  
$ echo "Minulla on koirra." | singularity_wrapper run \  
  stream fi_tdt parse_plaintext
```