

Machine Learning on Puhti

Part 2: Scaling up and using resources efficiently

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Webinar: Machine Learning on Puhti

Wed, June 3

Part 1: Getting started

- CSC's services
- Puhti supercomputer
- Available software
- Running jobs on Puhti
- Data storage

TODAY

Part 2: Scaling up and using resources efficiently

- Efficient data storage
- GPU utilization
- Multi-GPU and multi-node jobs
- Singularity containers

https://github.com/csc-training/ml-webinar/

Efficient data storage



Data storage on Puhti

- Disk space and number of files are limited on Puhti!
 - \rightarrow 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></user-name>	10 GiB	100 000 files	No
projappl	Project	/projappl/ <project></project>	50 GiB	100 000 files	No
scratch	Project	/scratch/ <project></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=cet>
#SBATCH --partition=qpu
#SBATCH --ntasks=1
#SBATCH --cpus-per-task=10
#SBATCH --mem=64G
#SBATCH --time=1:00:00
#SBATCH --qres=qpu: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 squeue to find out on what node (computer) it is running

• ssh into that node, e.g., ssh r01g01

• run nvidia-smi





GPU utilization

For a finished job:

- run gpuseff <jobid>
- shows GPU utilization 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

- If you need more than 4 GPUs, we recommend Horovod
- Supported for TensorFlow and PyTorch on Puhti
- Uses MPI and NCCL2 for interprocess communication
- Also requires some changes in the code, see e.g.: https://github.com/horovod/horovod/tree/master/examples
- In Puhti modules with -hvd suffix are Horovod-enabled

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=roject>
#SBATCH --partition=qpu
#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 (preferrably not in login nodes!):

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 koira." | singularity_wrapper run \
    stream fi_tdt parse_plaintext
```



Further resources

- Accessing and using Puhti: https://docs.csc.fi/computing/overview/
- MyCSC portal: https://my.csc.fi
- Machine learning software on Puhti:

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

• GPU-accelerated machine learning guide:

```
https://docs.csc.fi/support/tutorials/gpu-ml/
```

Slides and code from this webinar:

```
https://github.com/csc-training/ml-webinar/
```

 Deploying machine learning on Rahti: https://github.com/cscfi/rahti-ml-examples

Thank you!