



Outline

Part IB - Practical Guidelines for Running a Project:

Choosing Hyperparameters – a bit of exploration and exploitation

Job workflow - make it efficient and easy to organize

CPUs vs GPUs

Parallelize Models and Multinode Execution, with an exercise

Choosing Hyperparameters

- Hyperparameters are found by searching, not by the network algorithm
- Generally, hyperparameters related to:

 architecture (layers, units, activation, filters, ...)
 algorithm (learning rate, optimizer, epochs, ...)
 efficient learning (batch size, normalization, initialization, ...)
- Some options are determined by task: loss function, CNN vs MLP, ...
- Use what works, from related work or the latest recommendations,

Hyperparameters Search

- Can take a long time, hard to find global optimal
- Start with small data, short runs to get sense of range of good parameter values
- Easy but possibly time-consuming method: grid search over uniformly spaced values
- Do "exploration" then "exploitation", ie search wide then search deep Keras Tuner functions can help with the wide search

Keras Hyperparameter Search Tool

Keras Hypertuner class implements several search strategies:

Hyperband is like a tournament of hyperparameter configurations, with incremental training, to weed out worse ones

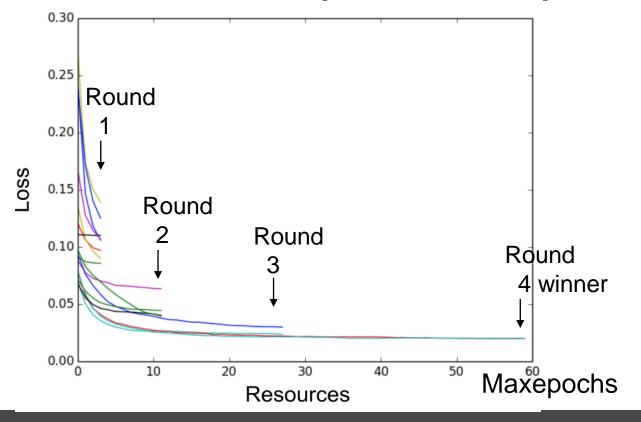
RandomSearch will search randomly through the space of configurations and try to find better regions

Bayesian optimization is like function approximation to pick out next configuration



Hyperband Bracket

Each round runs several network configurations for small number of epochs Several rounds with increasing epochs make up a bracket Several brackets are run to end up with several possible overall winners.



Note, you could run a small grid search around hyperband winners to confirm performance

Workflow and Organizing Jobs

Job Level: What makes sense to include in each job?

Model Level: run & test model for each parameter configuration

Data Level: loop through cross validation datasets (if applicable)

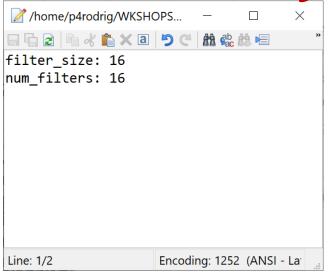
- Consider how long each a model runs for 1 configuration of hyperparameters for 1 dataset
- Organize jobs into reasonable chunks of work
- For large models consider model-checkpoints
- Tensorboard is available but needs to be secure (ask for details)



Organizing Configurations – one way

Code snippet: using 'YAML' file to set up hyperparameter configuration

Create text file with "Parameter: Value" pairs



Read file as python dictionary

```
import yaml
with open("./modelrun_args.yaml", "r") as f:
    my_yaml=yaml.safe_load(f) #this returns a python dictionary

filter_size=my_yaml.get("filter_size")
num_filters=my_yaml.get("num_filters")
print('arguments, filter_size:',filter_size,' num_filters',num_filters)
```

Example slurm job script and execution for Expanse

You could also modify or set up parameters; and save yaml files for each run

```
#!/usr/bin/env bash
#SBATCH --job-name =mnist0522
#SBATCH --account=sds164
#SBATCH --partition=compute
#SBATCH --nodes=1
#SBATCH --ntasks-per-node=128
#SBATCH --time=00:10:00
#SBATCH --output=myjoboutput.o%j.%N.out
module purge
module load singularitypro
module list
echo "filter_size: 3 " > modelrun_args.yaml
echo "num_filters: 16 " >> modelrun_args.yaml
singularity exec --bind /expanse,/scratch --nv \
  /cm/shared/apps/containers/singularity/tensorflow/tensorflow-latest.s
  python3 Intro mnist cnn2 forbatch.py > mymnist stdoutput.txt
```

note on using GPU

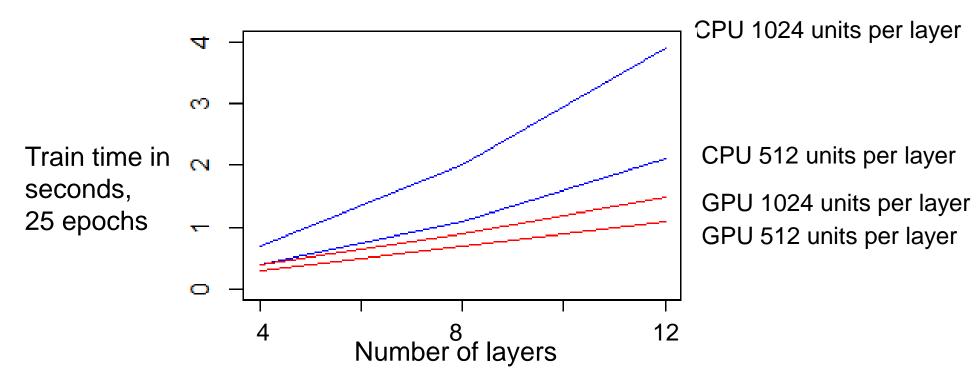
- GPU node has multiple GPU devices
- By default tensforflow will run on 0th gpu device if GPU is available, otherwise it will use all CPU cores

note on using GPU

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Code snippet to check for GPU devices

GPU shared (V100) vs CPU (128 cores) For MLP with Dense Layers, 80000x200 data matrix



GPUs faster, but you might have to wait more in job queue; also some memory limits compared to CPU, may need to use smaller batch size



Parallel DL models with multiple nodes/devices

- The main approach to parallelize training: Data Parallel:
 - 1. Split up data (in Keras see 'tf.data.Datasets' API or use numpy)
 - 2. Launch your script on each device
 - 3. Each device trains a copy of the model with a part of the data
 - 4. Aggregate parameter updates across model instances

Parallel DL models with multiple nodes/devices

- The main approach to parallelize training: Data Parallel:
 - 1. Split up data
 - 2. Launch your script on each device
 - 3. Each device trains a copy of the model with a part of the data
 - 4. Aggregate parameter updates
- Main tools: Keras/Tensorflow 'strategy' or use Horovod MPI wrappers
- Other approaches include Model Parallel (e.g. few layers per device), or Model and Data parallel
- Also, using mixed precision can reduce memory footprint

Keras/Tensorflow strategy single GPU node

Set up a 'mirror' strategy

```
mirrored_strategy = tf.distribute.MirroredStrategy(["GPU:0", "GPU:1", "GPU:2", "GPU:3"])
```

You also need the strategy scope around the model definition so that it can make copies

```
if (n_gpus>0):
    with mirrored_strategy.scope():
       multi_dev_model=build_model()
```

Then train as normal (use batch size multiple of 32)

Keras/Tensorflow strategy multiple GPU node

Keras also has a 'multiworker' strategy but it requires setting up config files with IP addresses

But, on HPC systems resources are shared so IP addresses are dynamic

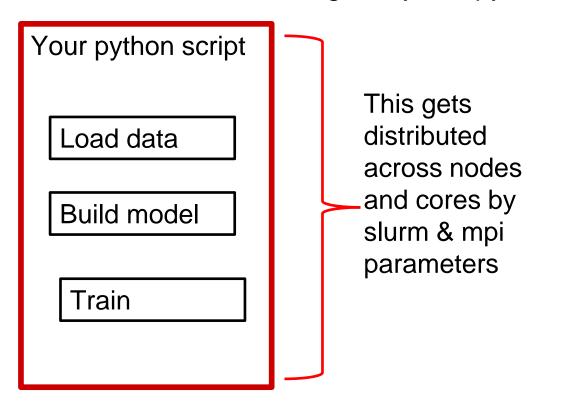
Thus, it is better to use Horovod with MPI and slurm batch job

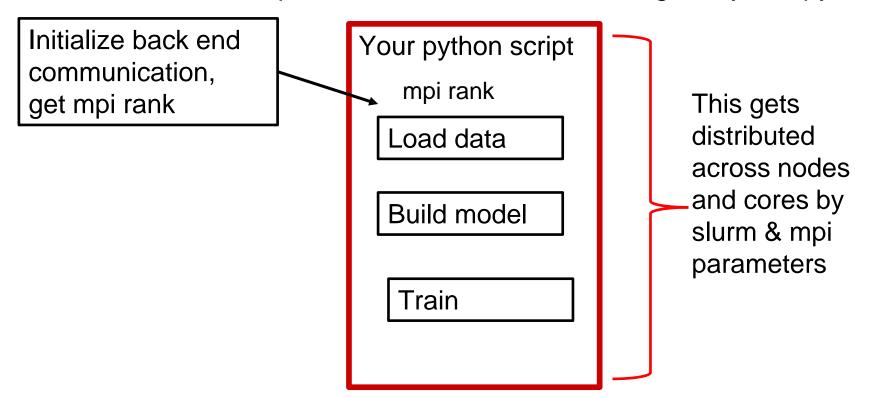
For example, single node, single device execution

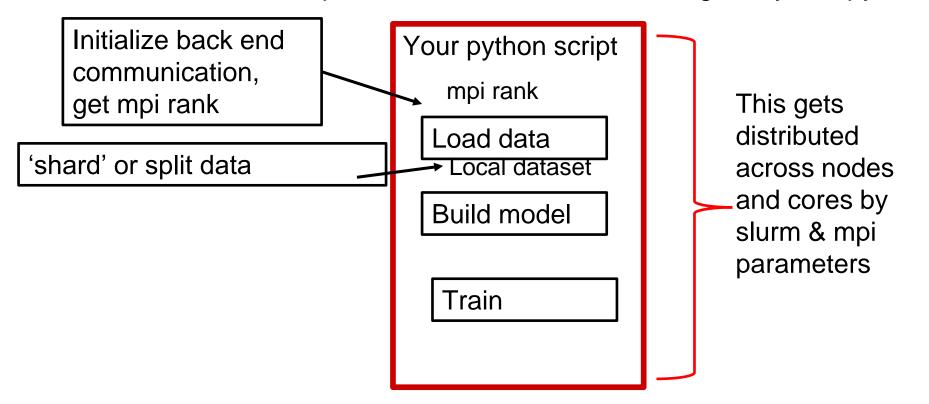
In slurm batch script:

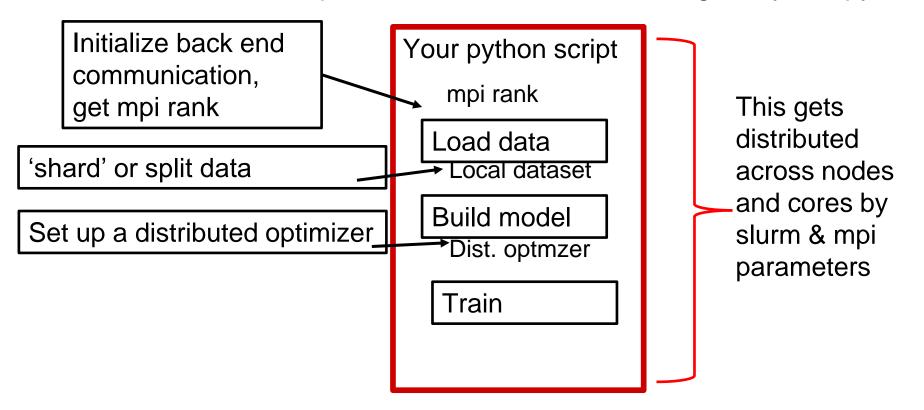
singularity → python

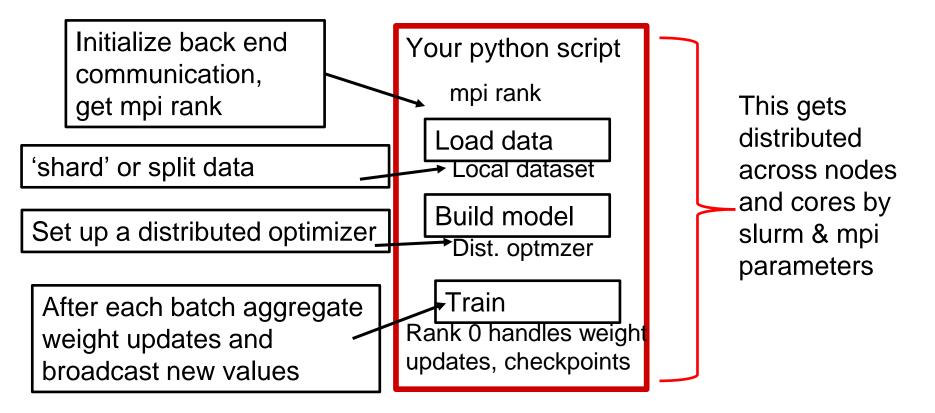
Your python script
Load data
Build model
Train











mpi launches one instance per processor

mpi launches one instance per processor

In slurm batch script:

mpirun –n **number of tasks** singularity → python

device =GPU:0

. . .

device =GPU:0

device =GPU:0

Your python script

mpi rank

Load data

Local dataset

Build model

Dist. optmzer

Train

Rank 0 handles updates

Your python script

mpi rank

Load data

Local dataset

Build model

Dist. optmzer

Train

Your python script

mpi rank

Load data

Local dataset

Build model

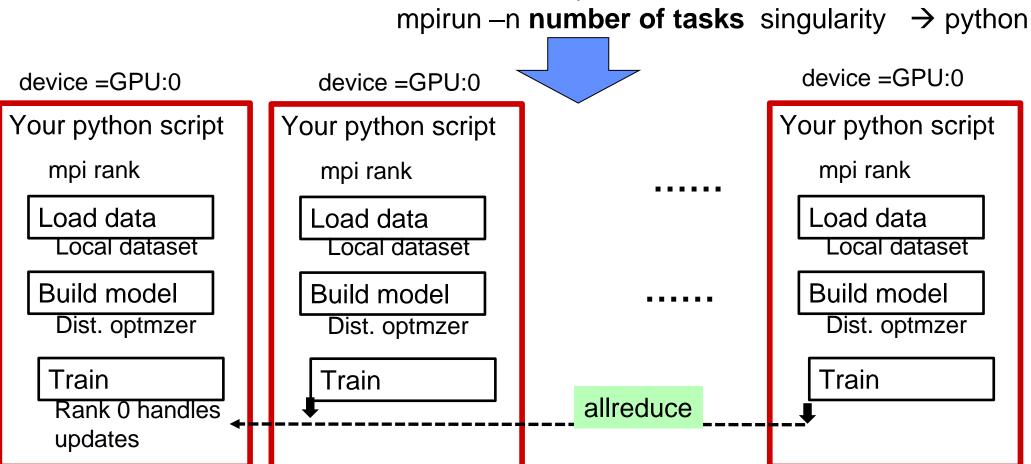
Dist. optmzer

Train



For each input batch: Horovod will aggregate & share weights updates

In slurm batch script:



For each input batch: Horovod will aggregate & share weights updates

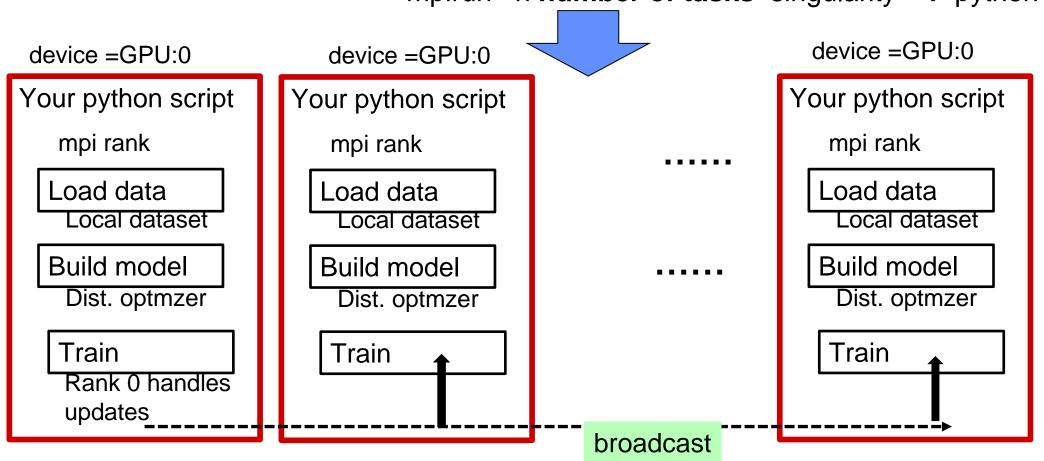
In slurm batch script:

mpirun –n **number of tasks** singularity \rightarrow python device =GPU:0 device =GPU:0 device =GPU:0 Your python script Your python script Your python script mpi rank mpi rank mpi rank Load data Load data Load data Local dataset Local dataset Local dataset Build model Build model Build model Dist. optmzer Dist. optmzer Dist. optmzer Train Train Train Rank 0 handles updates broadcast

For each input batch: Horovod will aggregate & share weights updates

In slurm batch script:

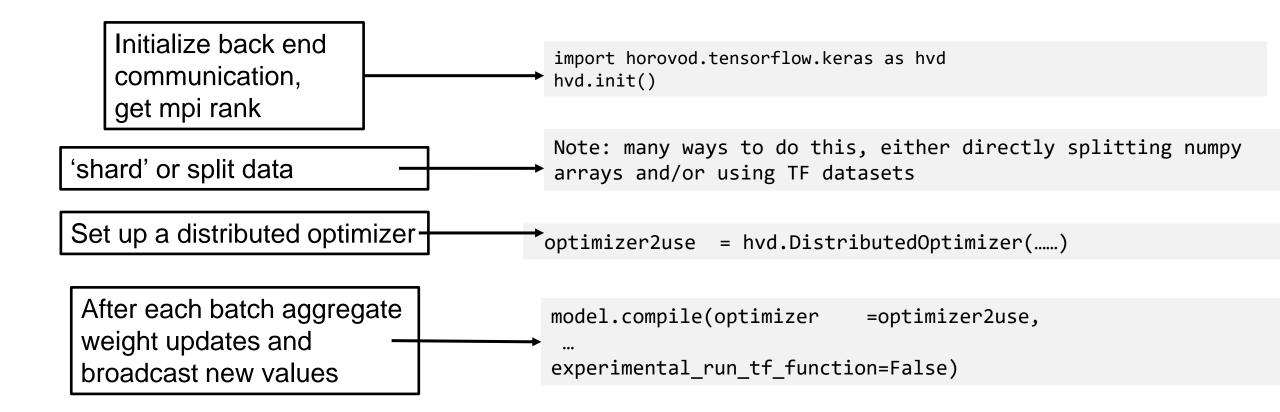
mpirun –n **number of tasks** singularity → python



Bigger batch size helps, but it uses more memory

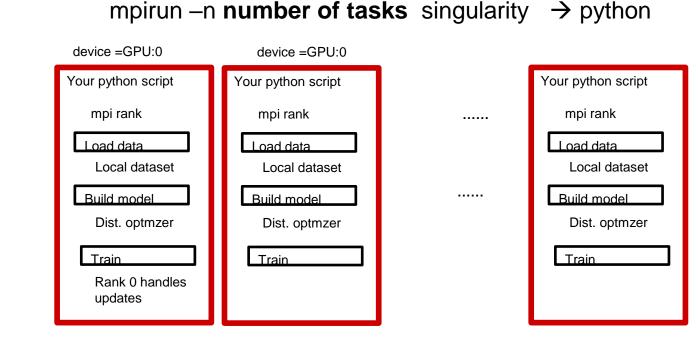
Code snippets – Horovod functions

Not many lines of code, but becareful with sharding, batch size, See https://horovod.readthedocs.io/en/latest/keras.html



Exercise, multinode MNIST programming and execution

- Goal: Get familiar with Keras and Horovod coding for multinode execution
- Goal: Get familiar with slurm batch script multinode parameters
- Let's login and start a notebook (see next pages for quick overview)

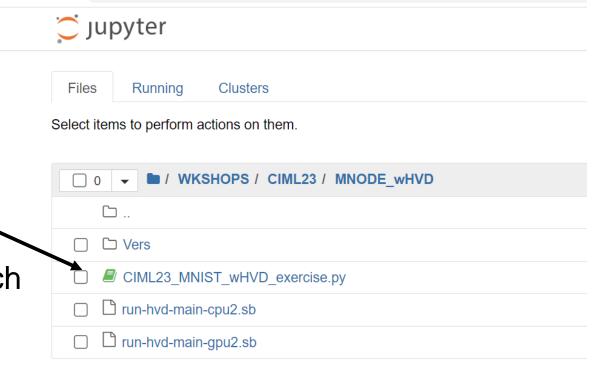


In terminal]\$ jupyter-compute-tensorflow

In jupyter notebook session open the MNIST_wHVD_exercise notebook

Open the *run-hvd-main-cpu2.sb* slurm batch script

Follow instructions in the notebook



Code highlights

Initialize back end communication, get mpi rank

split data using numpy arrays

```
# ------
per_worker_batch_size = 32  #Pick factors of 32 (especially for GPU)
num_workers = hvd.size()
```

Set up a distributed optimizer to manage weight updates

Also run sbatch command for this slurm script

Try reviewing stdout output file

```
Jupyter run-hvd-main-cpu2.sb✓ 15 minutes ago

File Edit View Language Plain Text

#SBATCH --job-name=tfhvd-cpu

#SBATCH --account=use300

#SBATCH --partition=compute

#SBATCH --nodes=2

#SBATCH --ntasks-per-node=16 #<<<<<----- change this to 16 and observe changes in training time

#SBATCH --mem=2436

#SBATCH --mem=2436

#SBATCH --time=00:15:00

#SBATCH --output=slurm.cpu2.%x.o%j.out
```

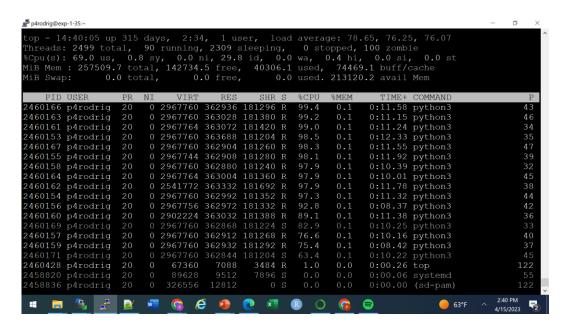
```
p4rodrig@login01 MNODE_wHVD]$
p4rodrig@login01 MNODE_wHVD]$ grep 'rank' stdout_cpu2_mnist_32.txt |grep ' 0'
NFO, global rank: 0 localrank 0
NFO, global rank: 16 localrank 0
NFO,cpus available rank: 0 [PhysicalDevice(name='/physical_device:CPU:0', dev
ce_type='CPU')]
This is rank 0 instance model evaluation
p4rodrig@login01 MNODE_wHVD]$
```

```
[p4rodrig@login01 MNODE_wHVD]$
[p4rodrig@login01 MNODE_wHVD]$ grep 'done, rk: 15' stdout_*
stdout_cpu2_mnist_32.txt:INFO,done, rk: 15 train time: 2.48225 secs
stdout_mainhvd_cpu2.txt:INFO,done, rk: 15 train time: 2.31222 secs
[p4rodrig@login01 MNODE_wHVD]$
[p4rodrig@login01 MNODE_wHVD]$
```

optional extra to try

You can run:

- \$ squeue —u userid to see nodes running your job
- \$ ssh exp-XX-YY to login to node
- \$ top -u userid to see processing on a CPU job



Or run: \$ nvidia-smi to see usage on GPU devices

🧬 p4rodrig	@login02:/exp	panse/lustre/	/projects/sds164/	p4rodrig/TFwl	-IVDtests				_		>
p4rod	rig@loo				squeue -u NAME		TIME	NODES	S NODEL	IST (F	REF
p4rod p4rod	rig@log	gin02 5-10-5	TFwHVDte 7 ~]\$ nv	sts]\$	vd-gp p4ro ssh exp-10 mi		0:01	1	. exp-1	0-57	
NVID					Version:						+
GPU Fan	Name		Persist	ence-M	Bus-Id 	Disp.A	Vola	tile (Jncorr. Comput	ECC	
					00000000 1005M:			11%	Def	0 ault N/A	-
					00000000 1005M:			6%	Def	0 ault N/A	

End

