Distributed training with TensorFlow on Piz Daint

Synchronous Distributed Training with TensorFlow and Horovod

Rafael Sarmiento and Henrique Mendonça ETHZürich / CSCS CSCS/USI Summer School 2020

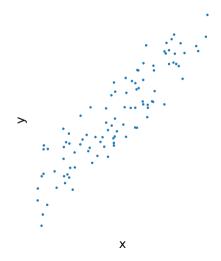


Outline

- Stochastic Gradient Descent
- [lab] Simple Stochastic Gradient Descent
- Synchronous Distributed Stochastic Gradient Descent
- Ring Allreduce
- Horovod
- [lab] Simple Stochastic Gradient Descent with Horovod



We want to train a model on this data



We choose a model and a cost function

$$y = mx + n$$

$$L = \frac{1}{N} \sum_{i}^{N} (\hat{y}_i - y_i)^2$$

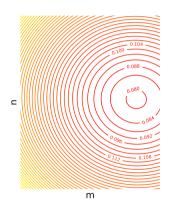
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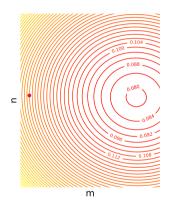
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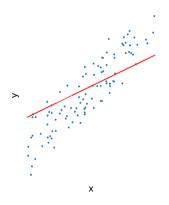


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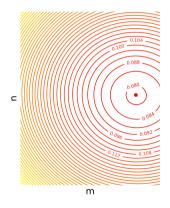
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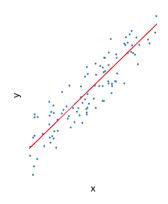
We need to choose an optimizer

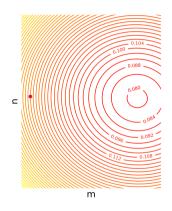




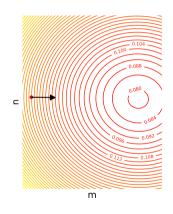
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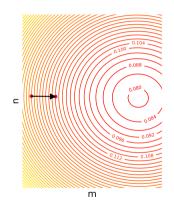




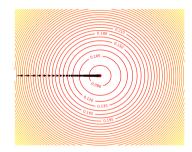
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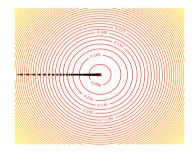


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- Update the parameters $W_t = W_{t-1} \eta \frac{\partial L}{\partial W} \big|_{\{x,y\}_{t-1}}$



Gradient
Descent
batch_size = training_set_size





Gradient
Descent
batch_size = training_set_size

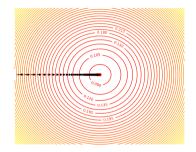
0.100 0.100 0.100 0.100 0.100 0.100

Stochastic Gradient

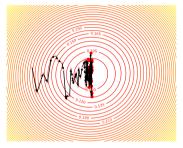
Descent

batch_size = 1

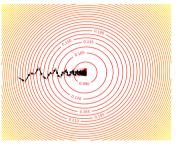




Gradient
Descent
batch_size = training_set_size



Stochastic Gradient
Descent
batch_size = 1



Minibatch Stochastic Gradient

Descent

1 < batch_size < training_set_size

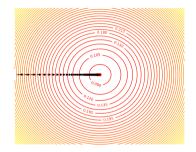


[lab] Simple Stochasting Gradient Descent

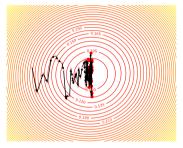
Let's run the notebook 1-linear_regression_SGD_TF2-simple.ipynb. There we use an unidimensional linear model to understand the trajectories of the SGD minimization.

Try different batch sizes and see how the trajectory changes.

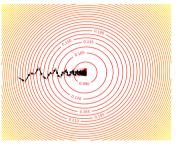




Gradient
Descent
batch_size = training_set_size



Stochastic Gradient
Descent
batch_size = 1



Minibatch Stochastic Gradient

Descent

1 < batch_size < training_set_size



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- Large batches may not fit on the GPU memory



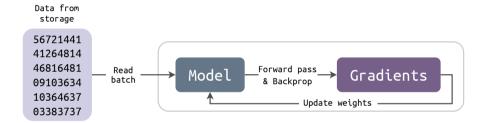
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- Splitting the training into multiple nodes/GPUs enables the use of large batches



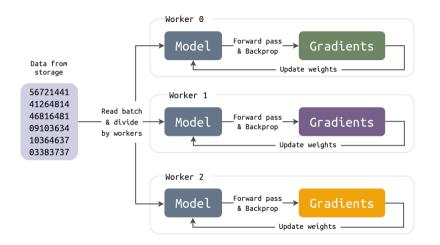
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- Large batches may not fit on the GPU memory
- Splitting the training into multiple nodes/GPUs enables the use of large batches
- Multiple nodes/GPUs does not necessarily mean more throughput or faster convergence



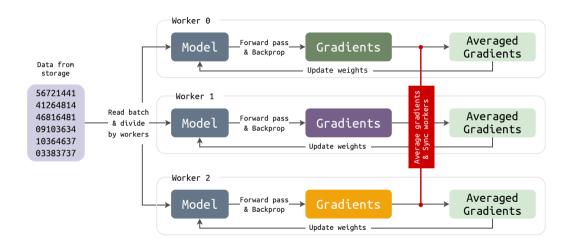
Distributing the training with data parallelism



Distributing the training with data parallelism



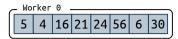
Distributing the training with data parallelism



The Allreduce operation

- The Allreduce name comes from the MPI standard.
- MPI defines the function MPI_Allreduce to reduce values from all ranks and broadcast the result of the reduction such that all processes have a copy of it at the end of the operation.
- Allreduce can be implemented in different ways depending on the problem.





65 18 20 21 40 11 50 5

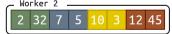
10 36 1 34 6 17 9 1

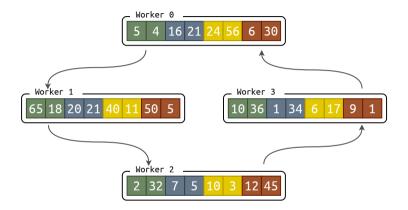
Worker 2 2 32 7 5 10 3 12 45

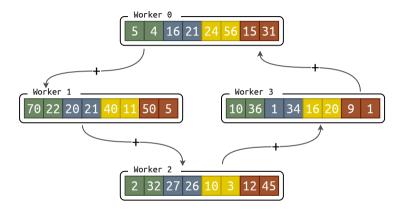


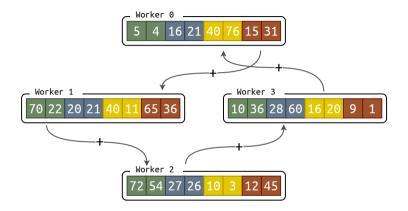


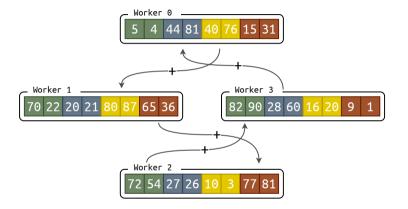


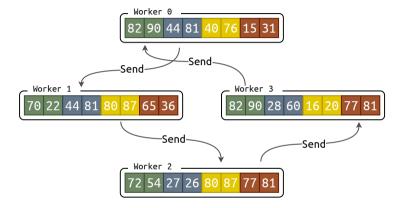


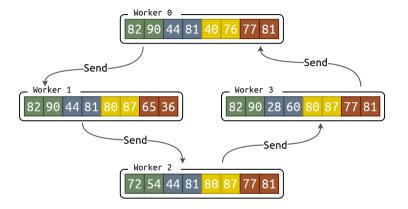


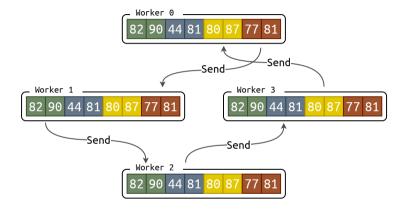












- ullet Each of the N workers communicates only with other two workers 2(N-1) times.
- The values of the reduction are obtained with the first N-1 communications.
- The second N-1 communications are performed to update the reduced values on all workers.
- The total amount of data sent by each worker $\left[2(N-1)\frac{\text{array_size}}{N}\right]$ is virtually independent of the number of workers.

Communication between Cray XC50 Nodes on Piz Daint

- Aries interconnect with the Dragonfly topology
- Direct communications between nodes on the same electrical group (2 cabinets / 384 nodes)
- Communications between nodes on different electrical groups passes by switches (submit with option #SBATCH --switches=1 to make your job wait for a single-group allocation)
- More info on CSCS user portal



Horovod



Horovod is an open-source distributed training framework for TensorFlow, Keras, PyTorch, and MXNet developed by Uber. The goal of Horovod is to make distributed Deep Learning fast and easy to use.

Horovod



- Minimal code modification required
- Uses bandwidth-optimal communication protocols
- Seamless integration with Cray-MPICH and use of the NVidia Collective Communications Library (NCCL-2)
- Actively developed
- Growing community

NVIDIA Collective Communications Library (NCCL)



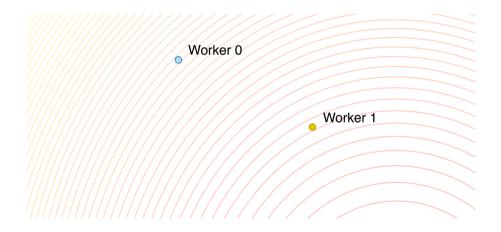
NCCL implements multi-GPU and multi-node collective communication primitives that are performance optimized for NVIDIA GPUs. NCCL provides routines such as Allgather, Allreduce and Broadcast, optimized to achieve high bandwidth over PCIe and NVLink high-speed interconnect.

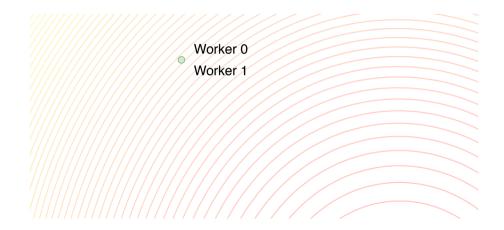
Horovod: 1. Import and initialize the library (tf.keras)

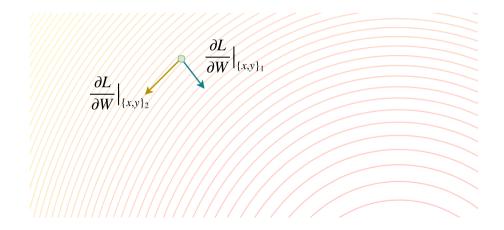
import horovod.tensorflow.keras as hvd

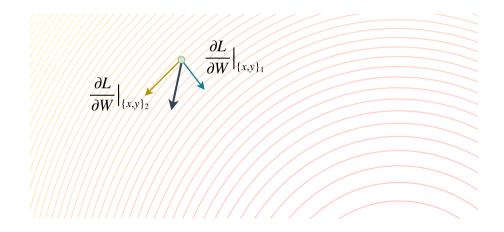
hvd.init()

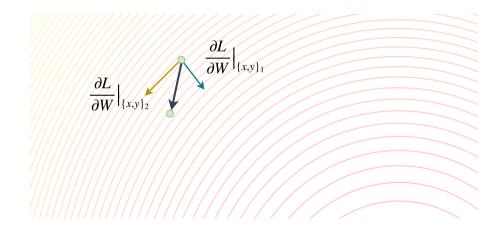


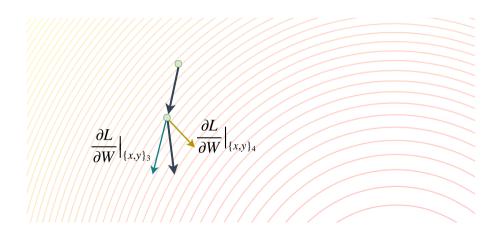












Horovod: 2. Sync the initial state of the workers (tf.keras)

```
initial_sync = hvd.callbacks.BroadcastGlobalVariablesCallback(0)
model.fit(dataset, ..., callbacks=[initial_sync])
```



Horovod: 3. Wrap the optimizer with Horovod's one (tf.keras)

```
opt = tf.keras.optimizers.SGD(learning_rate)
opt = hvd.DistributedOptimizer(opt)
```

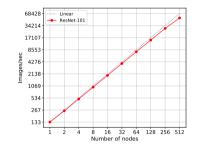


Horovod: 4. Checkpoints

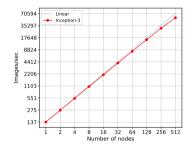
```
# Save checkpoints for the worker of rank 0.
# This will prevent all workers from corrupting a
# single checkpoint file.
if hvd.rank() == 0:
```



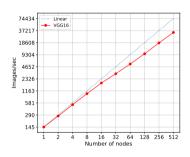
Benchmarks results on Piz Daint (CNNs on Imagenet)



num layers : 347 num weights: 44,601,832



num layers : 313 num weights: 23,817,352



num layers : 23 num weights: 138.357.544



Running TensorFlow + Horovod on Piz Daint

```
#!/bin/bash -l
#SBATCH --iob-name=tf hvd
#SBATCH --time=00:15:00
#SBATCH --nodes=2
#SBATCH --ntasks-per-core=1
#SBATCH --ntasks-per-node=1
#SBATCH --cpus-per-task=12
#SBATCH --constraint=qpu
module load daint-gpu
module load Horovod/0.19.1-CravGNU-19.10-tf-2.2.0
export OMP NUM THREADS=$SLURM CPUS PER TASK
export NCCL DEBUG=INFO
export NCCL IB HCA=ipoqif0
export NCCL IB CUDA SUPPORT=1
srun python my script.py
```



Some additional considerations

- Data must be split equally by workers to avoid load imbalance.
- If applicable, data can be split such that each worker does not need to read all files.
- Dataset splits resulting in non-homogeneous datasets may harm the convergence.
- Consider scaling the learning rate (learning_rate * hvd.size())

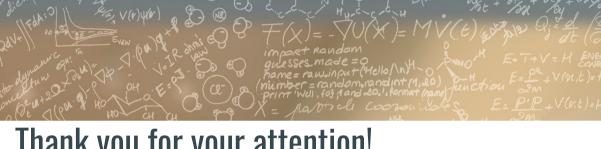


[lab] Simple Stochastic Gradient Descent with Horovod

The notebook 2-exercise-linear_regression_SGD_TF2-horovod.ipynb uses the same model that we saw before. We will addapt it to Horovod and we will run it with 2 workers.

Visualize the trajectories before and after adding each Horovod modification. Try to understand why each line of Horovod is needed.





Thank you for your attention!

