

# DISTRIBUTED TRAINING WITH PYTORCH

#### **Umar Jamil**

Downloaded from: https://github.com/hkproj/pytorch-transformer-distributed
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#### **Outline**

- Introduction to distributed training
  - Why we need it
  - Data Parallel vs Model Parallel
- Review of neural networks
  - Loss function and gradient
  - Gradient accumulation
- Distributed Data Parallel training
  - How it works
  - Communication primitives
    - Broadcast operator
    - Reduce operator
    - All-Reduce operator
  - Managing failover
- Coding session
  - Infrastructure (Paperspace)
  - PyTorch code
- How PyTorch handles Distributed Data Parallel training
  - Bucketing
  - Computation-Communication overlap during backpropagation

#### What is distributed training?

Imagine you want to train a Language Model on a very big dataset, for example the entire content of Wikipedia. The dataset is quite big, because it is made up of millions of articles, each of them with thousands of tokens. To train this model on a single GPU may be possible, but it poses some challenges:

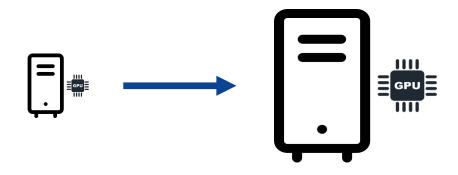
- 1. The model may not fit on a single GPU: this happens when the model has many parameters.
- 2. You are forced to use a small batch size because a bigger batch size leads to an Out Of Memory error on CUDA.
- 3. The model may take years to train because the dataset is huge.

If any of the above applies to you, then you need to scale your training setup. Scaling can be done vertically, or horizontally. Let's compare these two options.

Vertical scaling

money is all you need!

No code change

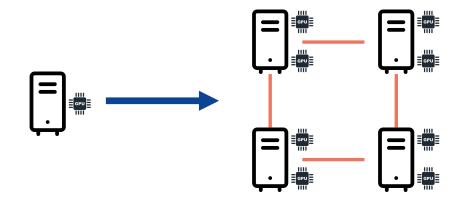


1x Server 8GB RAM 4GB GPU Memory 1x Server 64GB RAM 32GB GPU Memory

#### Horizontal scaling

Strategy is all you reed
Minimal code change (thanks to PyTorch)

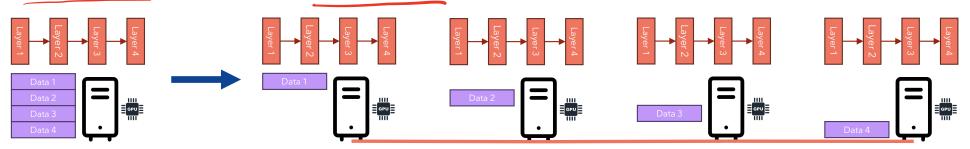
In this video we will explore horizontal scaling



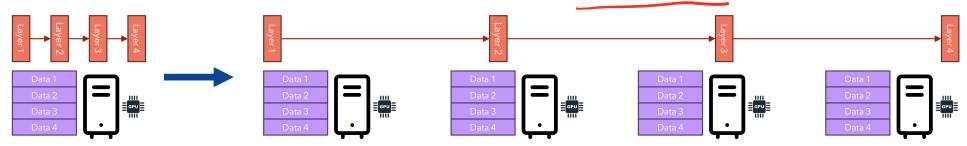
1x Server 8GB RAM 4GB GPU Memory **4x Servers** 8GB RAM 4GB GPU Memory (x2)

#### Data Parallelism vs Model Parallelism

If the model **can** fit within a single GPU, then we can distribute the training on multiple servers (each containing one or multiple GPUs), with each GPU processing a subset of the entire dataset in parallel and synchronizing the gradients during backpropagation. This option is known as **Data Parallelism**.



If the model **cannot** fit within a single GPU, then we need to "break" the model into smaller layers and let each GPU process a part of the forward/backward step during gradient descent. This option is known as **Model Parallelism**.



In this video, we will focus on Data Parallelism.

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#### A review of neural networks: a practical example

Imagine you want to train a neural network to predict the price  $(y_{pred})$  of a house given two variables: the number of bedrooms in the house  $(x_1)$  and the number of bathrooms in the house  $(x_2)$ . We think that the relationship between the output and the input variables is linear.

$$y_{pred} = x_1 w_1 + x_2 w_2 + b$$

Our goal is to use stochastic gradient descent to find the values of the parameters  $w_1$ ,  $w_2$  and b such that the MSE loss between the actual house price  $(y_{target})$  and the predicted  $(y_{pred})$  is minimized.

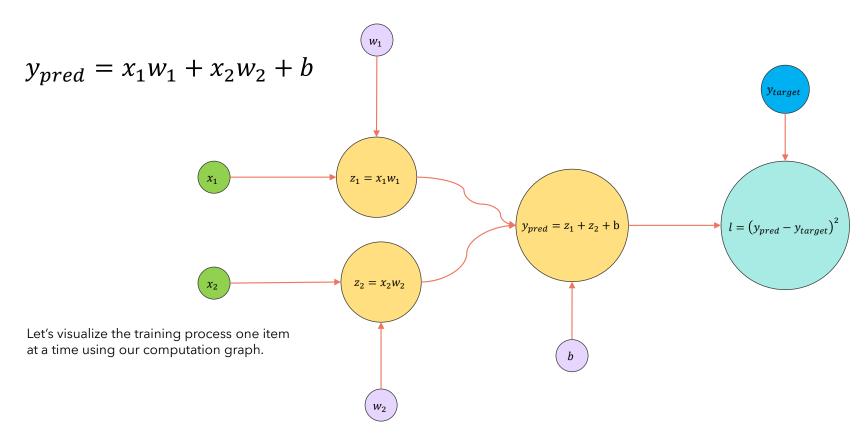
$$\underset{w1,w2,b}{\operatorname{argmin}} (y_{pred} - y_{target})^2$$

#### PyTorch's training loop (without accumulation)

```
def train no accumulate(params: ModelParameters, num epochs: int = 10, learning rate: float = 1e-3):
   for epoch in range(1, num epochs+1):
       for (x1, x2), y target in training data:
           # Calculate the output of the model
           z1 = x1 * params.w1
           z2 = x2 * params.w2
           y_pred = z1 + z2 + params.b
           loss = (v pred - v target) ** 2
           # Calculate the gradients of the loss w.r.t. the parameters
           loss.backward()
           # Update the parameters (at each iteration)
           with torch.no grad():
                # Equivalent to calling optimizer.step()
                params.w1 -= learning rate * params.w1.grad
               params.w2 -= learning_rate * params.w2.grad
                params.b -= learning rate * params.b.grad
               # Reset the gradients to zero
                # Equivalent to calling optimizer.zero_grad()
                params.w1.grad.zero_()
               params.w2.grad.zero_()
                params.b.grad.zero_()
```

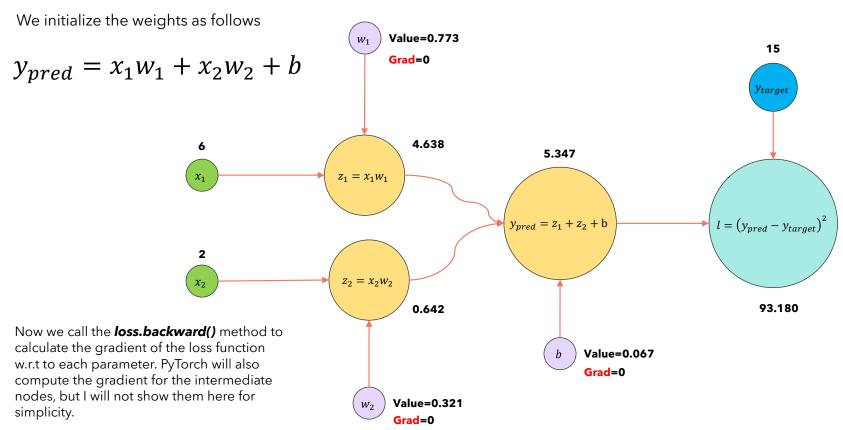
#### Computation graph

PyTorch will convert our neural network into a computational graph.

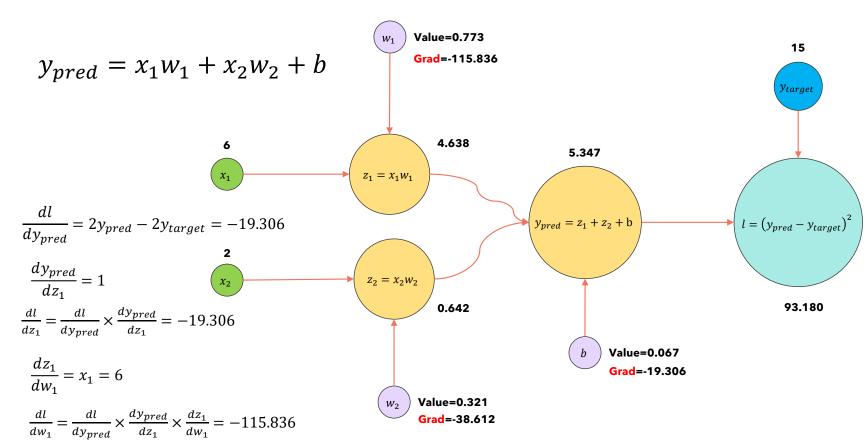


# Computational graph: step 1 (forward)

We run a forward step using the input  $x_1 = 6$ ,  $x_2 = 2$  and  $y_{target} = 15$ .

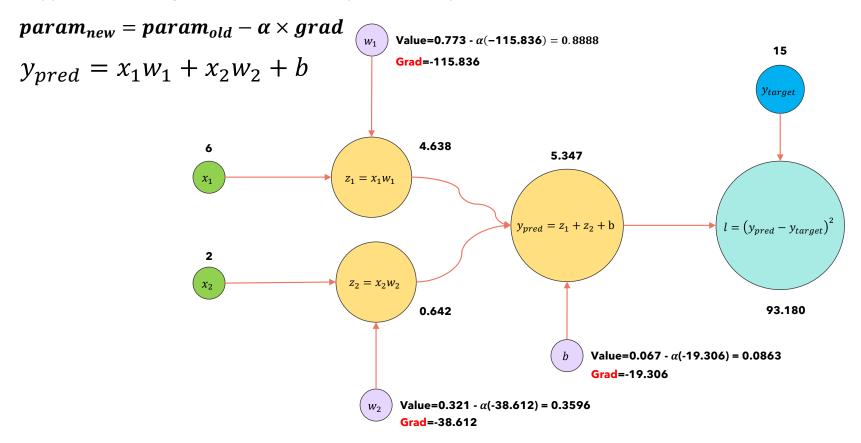


# Computational graph: step 1 (loss.backward)



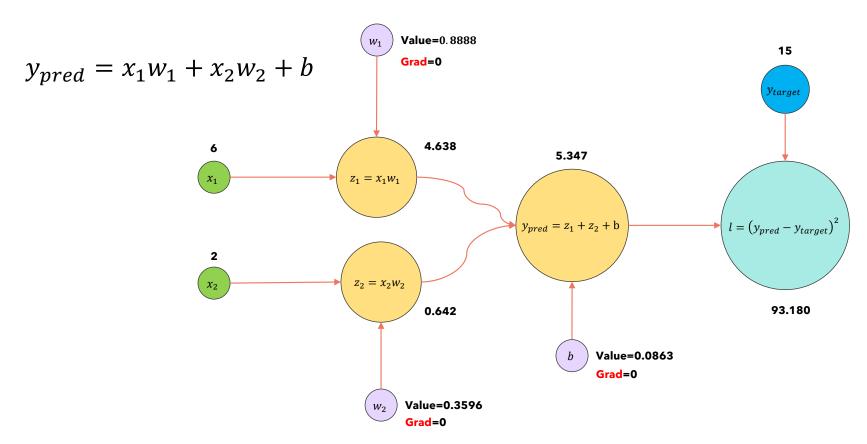
# Computational graph: step 1 (optimizer.step)

Suppose the **learning rate** is  $\alpha = 10^{-3}$ . Each parameter is updated as follows:



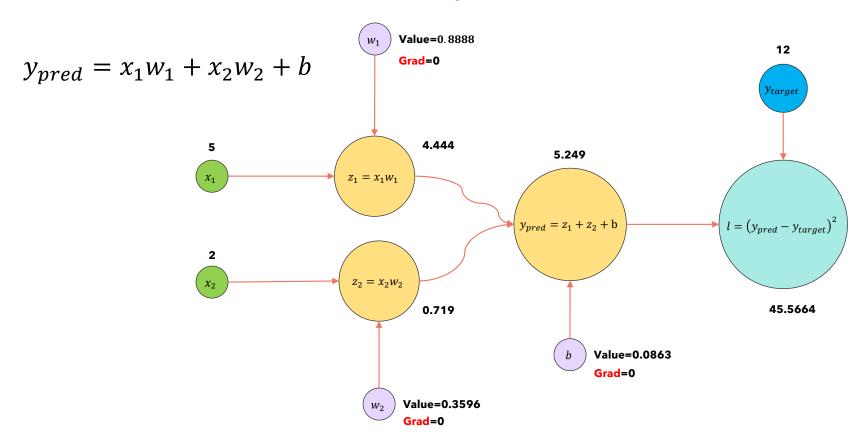
#### Computational graph: step 1 (optimizer.zero)

We reset the gradient of all the parameters to zero.

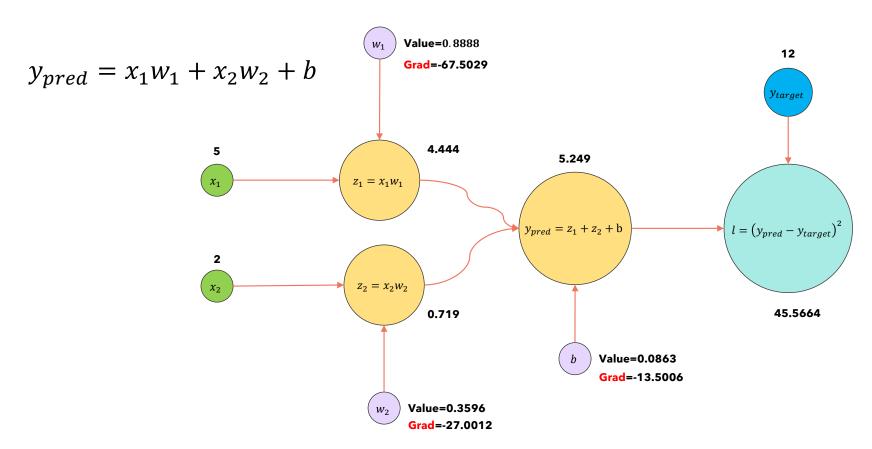


# Computational graph: step 2 (forward)

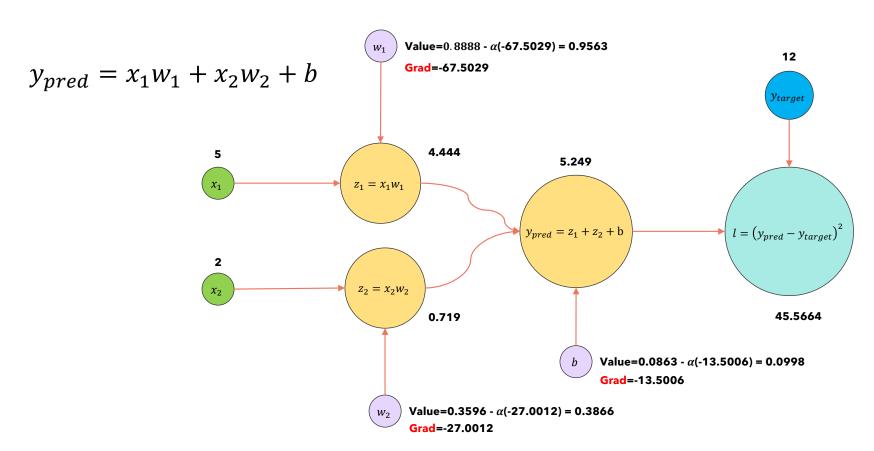
We run a forward step using the input  $x_1 = 5$ ,  $x_2 = 2$  and  $y_{target} = 12$ .



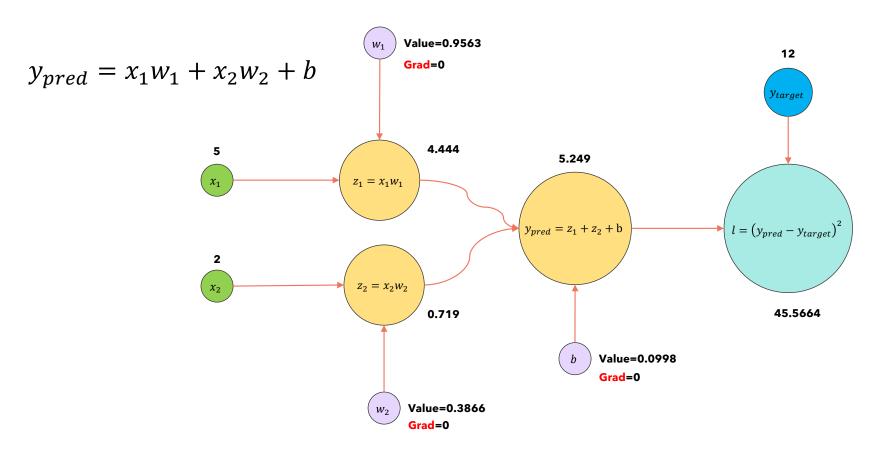
# Computational graph: step 2 (loss.backward)



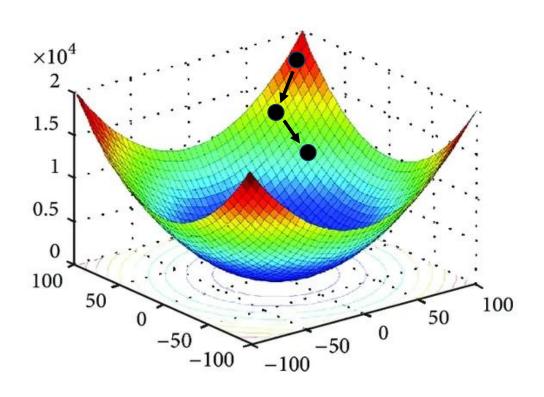
#### Computational graph: step 2 (optimizer.step)



# Computational graph: step 2 (optimizer.zero)



#### Gradient descent (without accumulation)



Initial weights

Data Item 1 (forward)

Data Item 1 (loss.backward)

Data Item 1 (optimizer.step)

Data Item 1 (optimizer.zero)

Data Item 2 (forward)

Data Item 2 (loss.backward)

Data Item 2 (optimizer.step)

Data Item 2 (optimizer.zero)

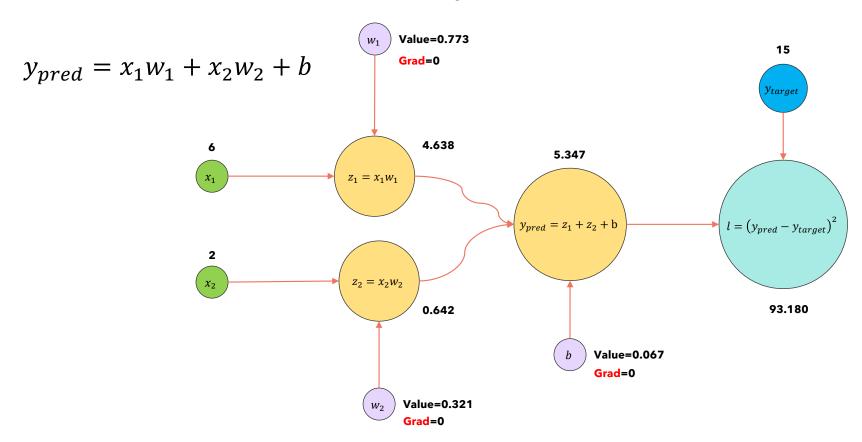
Without gradient accumulation, at every step (every data item), we update the parameters of the model.

#### PyTorch's training loop (with accumulation)

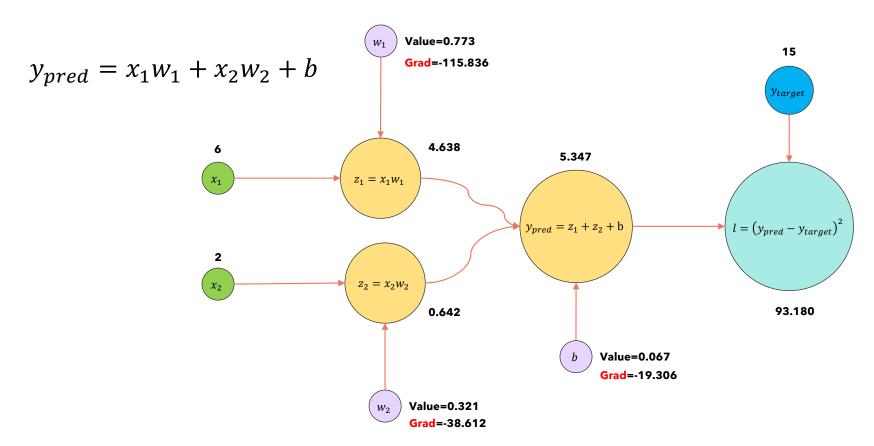
```
def train_accumulate(params: ModelParameters, num_epochs: int = 10, learning_rate: float = 1e-3, batch_size: int = 2):
    for epoch in range(1, num_epochs+1):
        for index, ((x1, x2), y target) in enumerate(training data):
            # Calculate the output of the model
            z1 = x1 * params.w1
            z2 = x2 * params.w2
           y_pred = z1 + z2 + params.b
            loss = (y pred - y target) ** 2
            # Calculate the gradients of the loss w.r.t. the parameters
            loss.backward()
            # Everytime we reach the batch size or the end of the dataset, update the parameters
            if (index + 1) % batch size == 0 or index == len(training data) - 1:
                with torch.no grad():
                    # Equivalent to calling optimizer.step()
                    params.w1 -= learning rate * params.w1.grad
                    params.w2 -= learning rate * params.w2.grad
                    params.b -= learning rate * params.b.grad
                    # Reset the gradients to zero
                    # Equivalent to calling optimizer.zero grad()
                    params.w1.grad.zero_()
                    params.w2.grad.zero_()
                    params.b.grad.zero_()
```

# Computational graph: step 1 (forward)

We run a forward step using the input  $x_1 = 6$ ,  $x_2 = 2$  and  $y_{target} = 15$ .

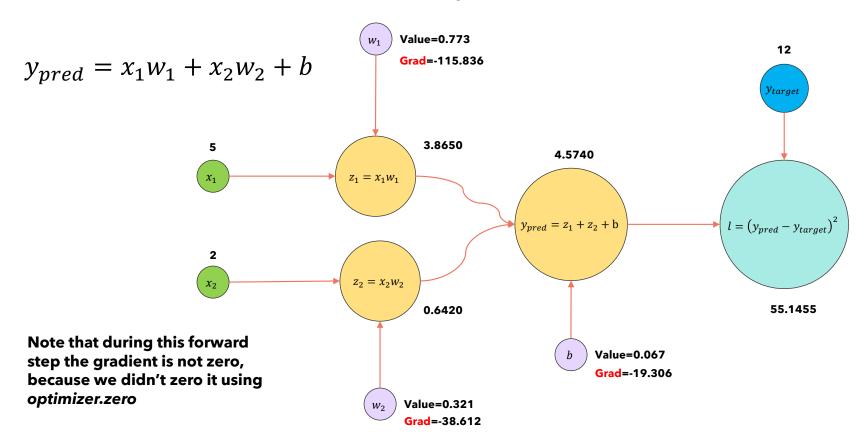


# Computational graph: step 1 (loss.backward)

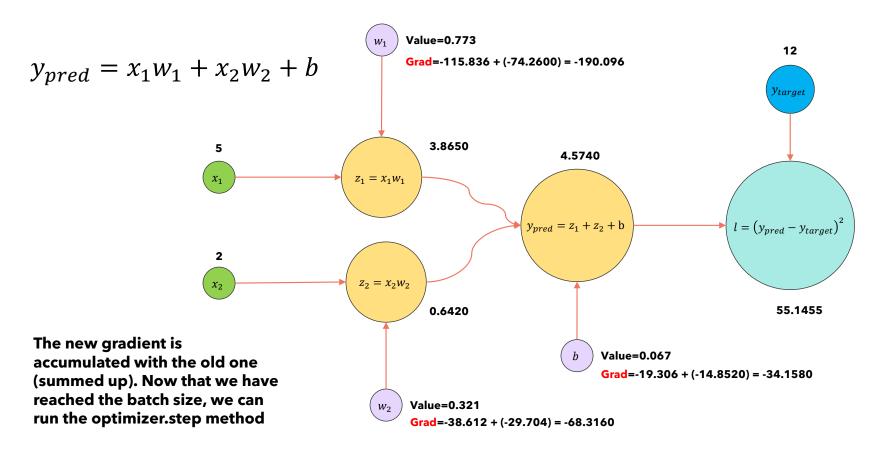


# Computational graph: step 2 (forward)

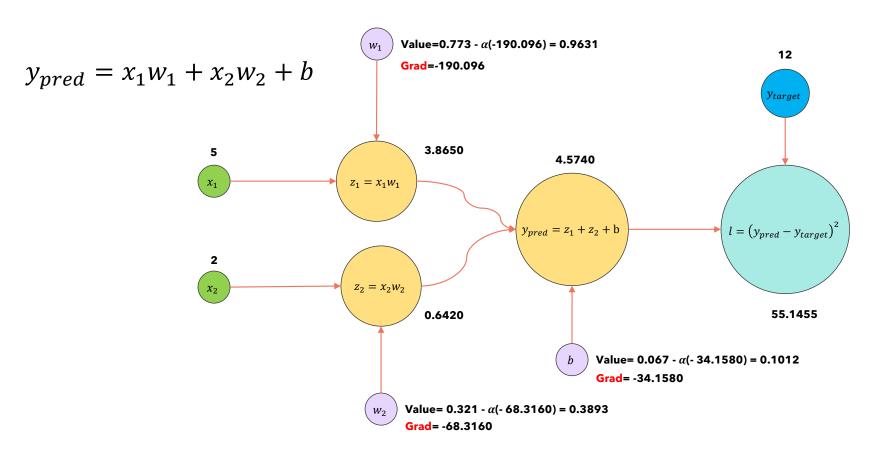
We run a forward step using the input  $x_1 = 5$ ,  $x_2 = 2$  and  $y_{target} = 12$ .



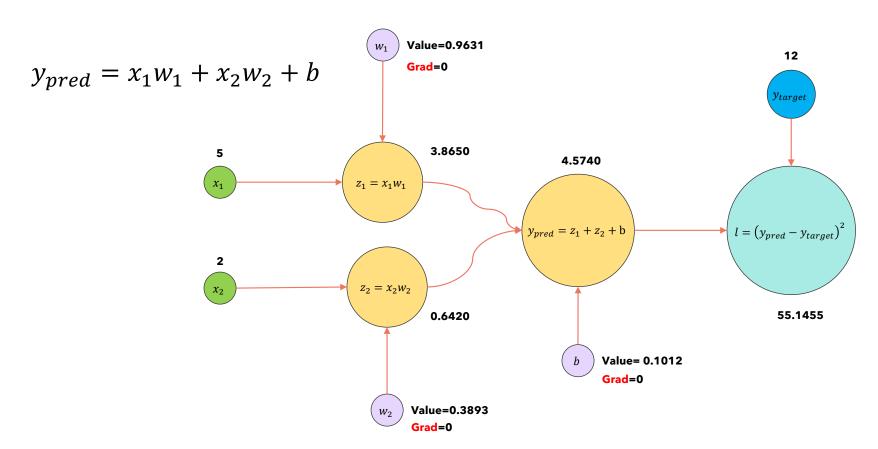
# Computational graph: step 2 (loss.backward)



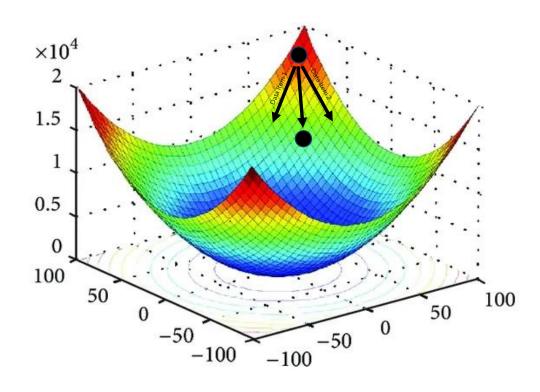
# Computational graph: step 2 (optimizer.step)



#### Computational graph: step 2 (optimizer.zero)



#### Gradient descent (with accumulation)



Initial weights

Data Item 1 (forward)

Data Item 1 (loss.backward)

Data Item 2 (forward)

Data Item 2 (loss.backward)

The two gradients are summed up

Data Item 2 (optimizer.step)

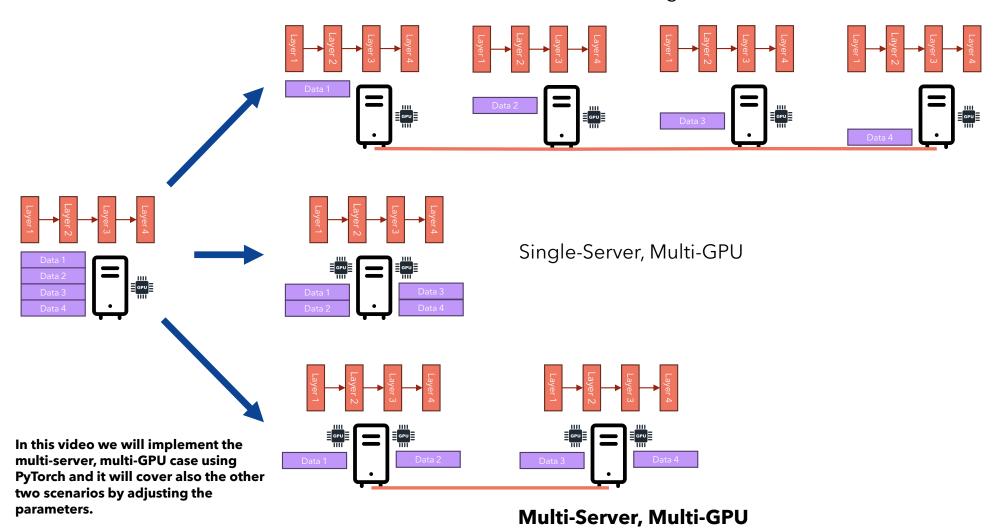
Data Item 2 (optimizer.zero)

With gradient accumulation, we update the parameters of the model only after we accumulated the gradient of a batch

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#### Multi-Server, Single-GPU

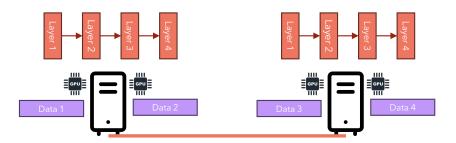


#### Distributed Data Parallel in detail

From now on, I will use the term "node" and "GPU" interchangeably. If a cluster is made up of 2 computers, each having 2 GPUs, then we have 4 nodes in total.

Distributed Data Parallel works in the following way:

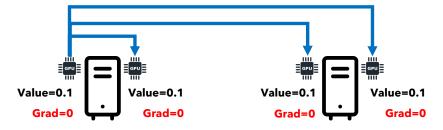
- 1. At the beginning of the training, the model's weights are initialized on one node and sent to all the other nodes (**Broadcast**)
- 2. Each node trains the same model (with the same initial weights) on a subset of the dataset.
- 3. Every few batches, the gradients of each node are accumulated on one node (summed up), and then sent back to all the other nodes (**All-Reduce**).
- 4. Each node updates the parameters of its local model with the gradients received using its own optimizer.
- 5. Go back to step 2



Model weights are initialized here (e.g., randomly)



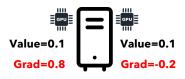
Initial weights are sent to all the other nodes (Broadcast)

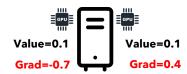


Each node runs a forward and backward step on one or more batch of data.

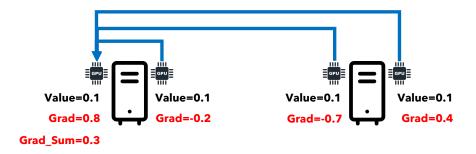
This will result in a local gradient.

The local gradient may be the accumulation of one or more batches.





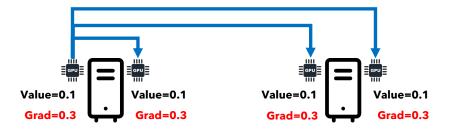
The sum of all the gradients is cumulated on one node (Reduce)



# Distributed Data Parallel: step 3

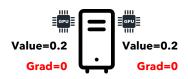
The cumulative gradient is sent to all the other nodes (Broadcast).

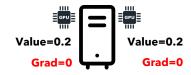
The sequence of Reduce and Broadcast are implemented as a single operation (All-Reduce).



# Distributed Data Parallel: step 4

Each node updates the parameters of its local model using the gradient received. After the update, the gradients are reset to zero and we can start another loop.





#### Collective Communication Primitives

In distributed computing environments, a node may need to communicate with other nodes. If the communication pattern is similar to a client and a server, then we talk about point-to-point communication, because one client connects to one server in a request-response chain of events.

However, there are cases in which one node needs to communicate to multiple receivers *at once*: this is the typical case of data parallel training in deep learning: one node needs to send the initial weights to all the other nodes. Moreover, all the other nodes, need to send their gradients to one single node and receive back the cumulative gradient. **Collective communication allows to model the communication pattern between groups of nodes**.

Let's visualize the difference between the two modes of communication.

### Point-To-Point

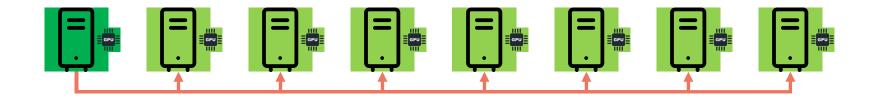
Imagine you need to send a file to 7 friends. With point-to-point communication, you'd send the file iteratively to each of the friend one by one. Suppose the internet speed is 1 MB/s and the file is 5 MB in size.



**Total time: 5s** 

### Point-To-Point

Since the internet communication is 1 MB/s and the file is 5 MB in size, your connection would be split among the 7 friends (each friend would be receiving the file at ~ 143 KB/s). **The total time is still 35s.** 

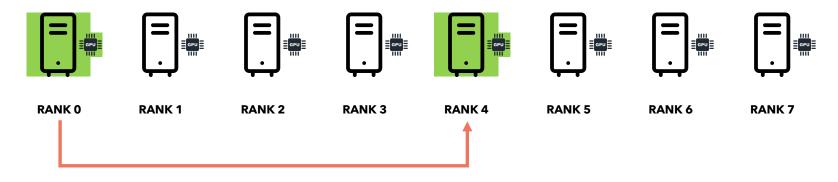


**Total time: 35s** 

Let's see how collective communication would manage this!

### Collective Communication: Broadcast

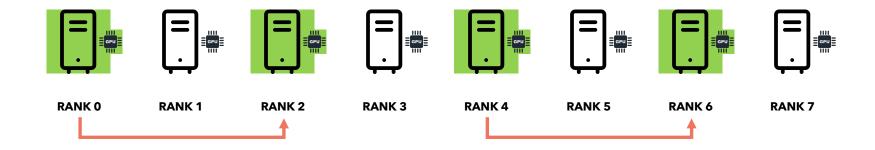
The operation of sending a data to all the other nodes is known as the **Broadcast** operator. Collective Communication libraries (e.g. NCCL) assign a unique ID to each node, known as **RANK**. Suppose we want to send 5 MB with an internet speed of 1 MB/s.



**Total time: 5s** 

### Collective Communication: Broadcast

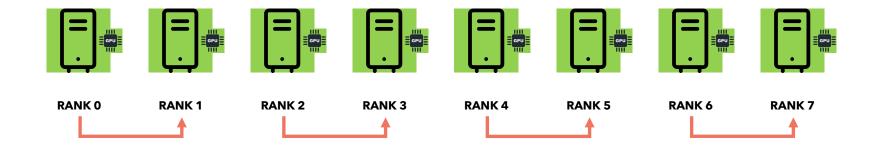
The operation of sending a data to all the other nodes is known as the Broadcast operator. Collective Communication libraries (e.g. NCCL) assign a unique ID to each node, known as **RANK**. Suppose we want to send 5 MB with an interned speed of 1 MB/s.



**Total time: 10s** 

#### Collective Communication: Broadcast

The operation of sending a data to all the other nodes is known as the Broadcast operator. Collective Communication libraries (e.g. NCCL) assign a unique ID to each node, known as **RANK**. Suppose we want to send 5 MB with an interned speed of 1 MB/s.



**Total time: 15s** 

This approach is known as Divide-and-Conquer. With collective communication, we exploit the interconnectivity between nodes to avoid idle times and reduce the total communication time.

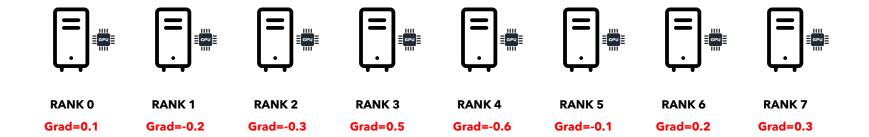
## Reduce operation

The **Broadcast** operator is used to send the initial weights to all the other nodes when we start the training loop.

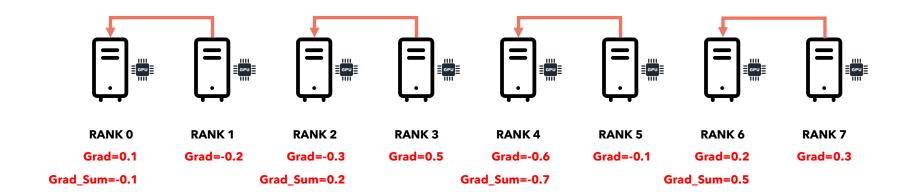
At every few batches of data processed by each node, the gradients of all nodes need to be sent to one node and accumulated (summed up). This operation is known as **Reduce**.

Let's visualize how it works.

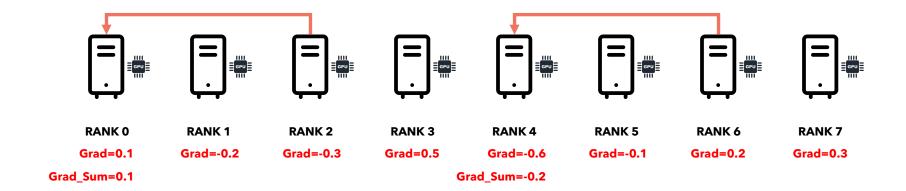
Initially, each node has its own gradient.



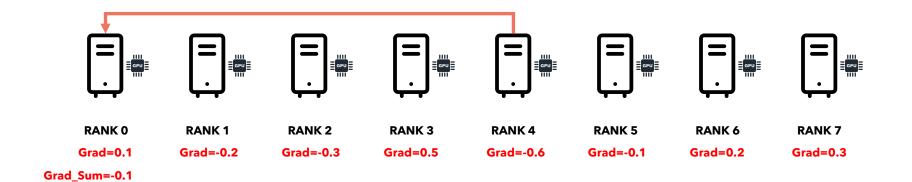
Each node sends the gradient to its adjacent node, who will sum it with its own gradient.



Step = 1



Step = 2



Step = 3

With only 3 steps we accumulated the gradient of all nodes into one node. It can be proven that the communication time is logarithmic w.r.t the number of nodes.

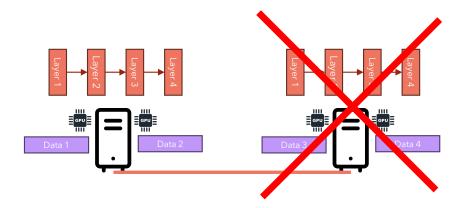
Having accumulated the gradients of all the nodes into a single node, we need to send the cumulative gradient to all the nodes. This operation can be done using a **Broadcast** operator.

The sequence of **Reduce-Broadcast** is implemented by another operator known as **All-Reduce**, whose runtime is generally lower than the sequence of **Reduce** followed by a **Broadcast**.

I will not show the algorithm behind **All-Reduce**, but you can think of it as a sequence of **Reduce** followed by a **Broadcast** operation.

## Failover: what happens if one node crashes?

Imagine you're training in a distributed scenario like the one shown below and one of the nodes suddenly crashes. In these case, 2 GPUs out of 4 become unreachable. How should the system react?

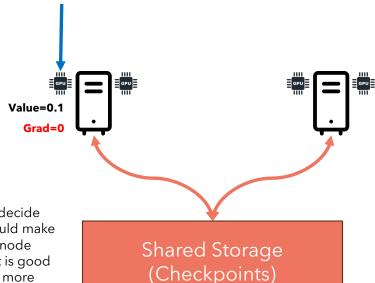


One way, would be to restart the entire cluster and that's easy. However, by restarting the cluster, the training would restart from zero, and we would lose all the parameters and computation done so far. **A better approach is to use checkpointing.** 

Checkpointing means saving the weights of the model on a shared disk every few iterations (for example every epoch) and resume the training from the last checkpoint in case there's a crash.

# Failover: using checkpointing

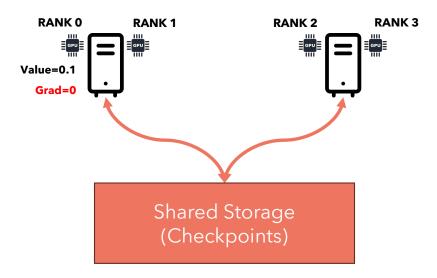
Model weights are initialized here (using the latest checkpoint)



We need a shared storage because PyTorch will decide which node will initialize the weights and we should make no assumption on which one will it be. So, every node should have access to the shared storage. Plus, it is good rule in distributed systems to not have one node more important than others, because every node can fail at any time.

# Failover: who should save the checkpoint?

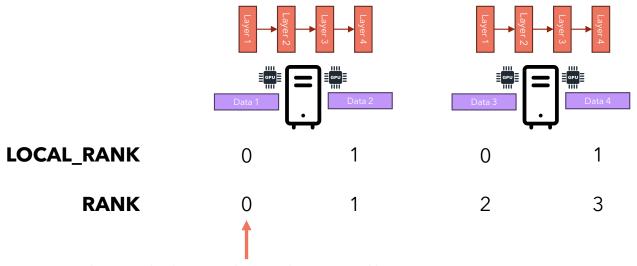
When we start the cluster, PyTorch will assign a unique ID (**RANK**) to each GPU. We will write our code in such a way that whichever node is assigned the **RANK 0** will be responsible for saving the checkpoint, so that the other nodes do not overwrite each other's files. So only one node will be responsible for writing the checkpoints and all the other files we need for training.



## LOCAL\_RANK vs RANK

The environment variable **LOCAL\_RANK** indicates the ID of the GPU on the local computer, while the **RANK** variable indicates the a globally unique ID among all the nodes in the cluster.

Please note that ranks are not *stable*, meaning that if you restart the entire cluster, a different node may be assigned the rank number 0.



Used to save checkpoint and/or initialize services (like W&B)

#### How to integrate DistributedDataParallel into your project?

```
def train():
    if global_rank == 0:
        initialize services() # W&B, etc.
    data_loader = DataLoader(train_dataset, shuffle=False, sampler=DistributedSampler(train_dataset, shuffle=True))
    model = MyModel()
    if os.path.exists('latest_checkpoint.pth'): # Load latest checkpoint
        # Also load optimizer state and other variables needed to restore the training state
        model.load state dict(torch.load('latest checkpoint.pth'))
    model = DistributedDataParallel(model, device_ids=[local_rank])
    optimizer = torch.optim.Adam(model.parameters(), lr=10e-4, eps=1e-9)
    loss_fn = torch.nn.CrossEntropyLoss()
    for epoch in range(num epochs):
        for data, labels in data_loader:
           loss = loss fn(model(data), labels) # Forward step
           loss.backward() # Backward step + gradient synchronization
           optimizer.step() # Update weights
           optimizer.zero_grad() # Reset gradients to zero
           if global_rank == 0:
                collect statistics() # W&B, etc.
        if global_rank == 0: # Only save on rank 0
           # Also save the optimizer state and other variables needed to restore the training state
           torch.save(model.state_dict(), 'latest_checkpoint.pth')
if name == ' main ':
    local_rank = int(os.environ['LOCAL_RANK'])
    global_rank = int(os.environ['RANK'])
    init_process_group(backend='nccl')
    torch.cuda.set device(local rank) # Set the device to local rank
    train()
    destroy_process_group()
```

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# **Prerequisites**

- Basic understanding of neural networks and PyTorch
- (Optional) watch my previous video on how to code a Transformer model from scratch

# When does PyTorch synchronize gradients?

PyTorch will synchronize the gradients every time we call the method **loss.backward**. This will lead to:

- 1. Each node calculating its local gradients (derivative of the loss function w.r.t each node of the computational graph)
- 2. Each node will send its local gradient to one single node and receives back the cumulative gradient (All-Reduce)
- 3. Each node will update its weights using the cumulative gradient and its local optimizer.

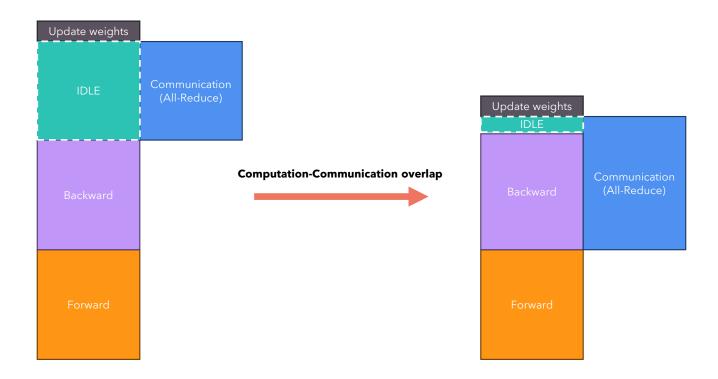
We can avoid PyTorch synchronizing the gradient at every backward step and instead, let it accumulate the gradient for a few steps by using the **no\_sync()** context. Let's see how it works.

# When does PyTorch synchronize gradients?

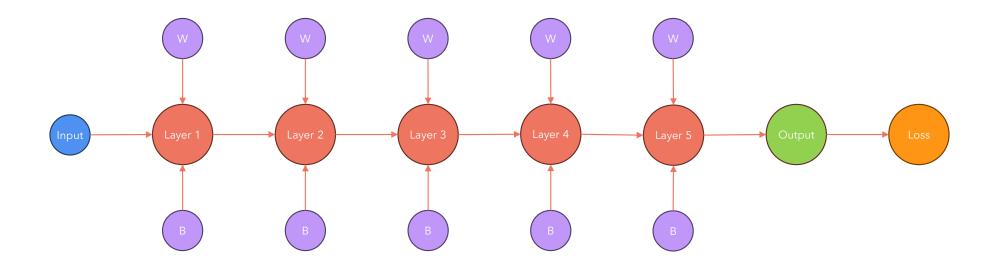
```
def train():
    if global rank == 0:
        initialize_services() # W&B, etc.
    data loader = DataLoader(train dataset, shuffle=False, sampler=DistributedSampler(train dataset, shuffle=True))
    model = MvModel()
    if os.path.exists('latest_checkpoint.pth'): # Load latest checkpoint
        # Also load optimizer state and other variables needed to restore the training state
       model.load state dict(torch.load('latest checkpoint.pth'))
    model = DistributedDataParallel(model, device_ids=[local_rank])
    optimizer = torch.optim.Adam(model.parameters(), lr=10e-4, eps=1e-9)
    loss_fn = torch.nn.CrossEntropyLoss()
    for epoch in range(num epochs):
        for data, labels in data loader:
            if (step number + 1) % 100 != 0 and not last step: # Accumulate gradients for 100 steps
                with model.no sync(): # Disable gradient synchronization
                    loss = loss fn(model(data), labels) # Forward step
                    loss.backward() # Backward step + gradient ACCUMULATION
            else:
                loss = loss fn(model(data), labels) # Forward step
                loss.backward() # Backward step + gradient SYNCHRONIZATION
                optimizer.step() # Update weights
               optimizer.zero_grad() # Reset gradients to zero
            if global rank == 0:
                collect_statistics() # W&B, etc.
       if global rank == 0: # Only save on rank 0
            # Also save the optimizer state and other variables needed to restore the training state
            torch.save(model.state dict(), 'latest checkpoint.pth')
if __name__ == '__main__':
    local rank = int(os.environ['LOCAL RANK'])
    global rank = int(os.environ['RANK'])
    init_process_group(backend='nccl')
    torch.cuda.set_device(local_rank) # Set the device to local rank
    train()
    destroy process group()
```

## PyTorch tricks: Computation-Communication overlap

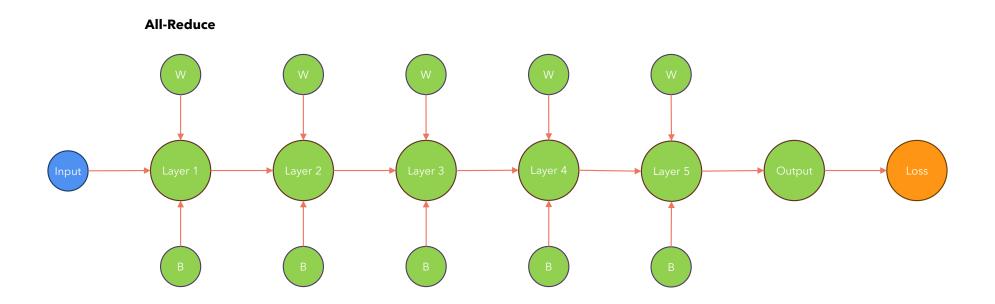
Since each GPU needs to send its gradient to a central node for accumulation, this can lead to an idle time in which the GPUs are not working, but only communicating with each other. PyTorch handles this communication delay in a smart way. Let's see how it works.



## Computation-Communication overlap: details

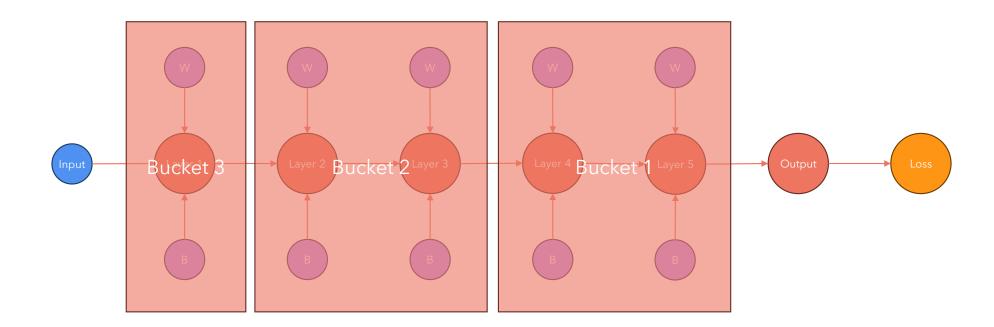


## Computation-Communication overlap: details



## Computation-Communication overlap: bucketing

Instead of sending each gradient one by one, which would result in a large communication overhead, gradients are packed together into buckets of equal size. PyTorch recommends 25MB as the size of the bucket.



# **Parallelism Fundamentals**

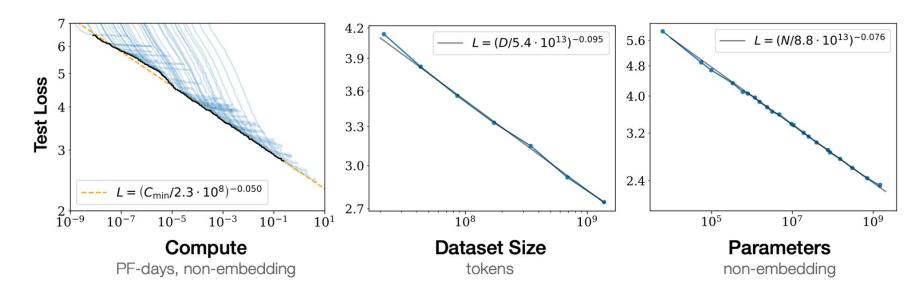
CS 229S Fall 2023



#### Today's Lecture

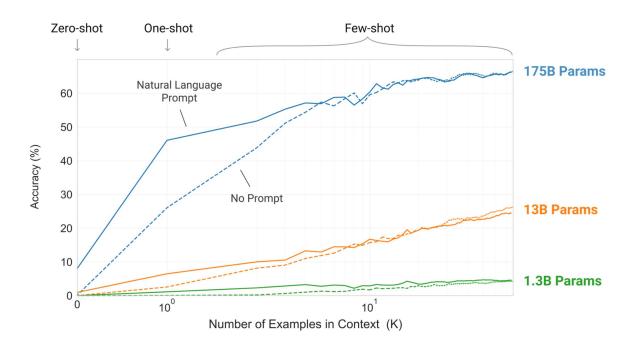
- Data parallelism
- Model Parallelism
- Pipeline Parallelism
- Practical challenges and solution for deploying parallelism in the real world

#### Model performance improves from all forms of scale



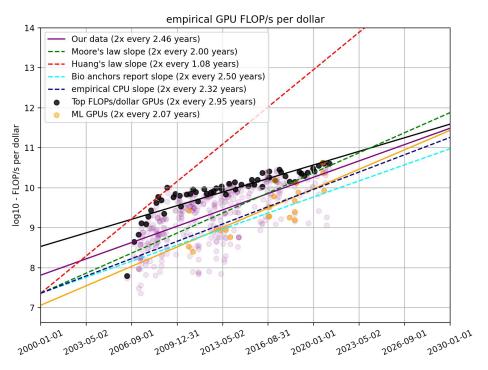
LLM performance improves from scaling up the total compute time, the dataset size, and the model size

#### LLMs exhibit emergent properties at scale



Scaling up to very large model and context sizes dramatically increases the capability of LLMs!

### Hardware improvements lag behind LLM scaling



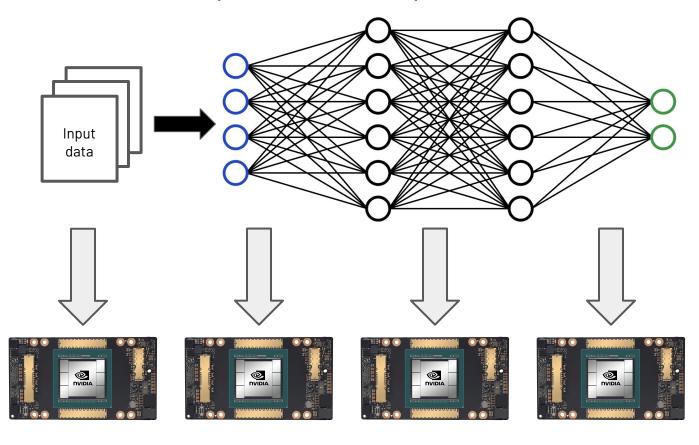
1000 parameters) GPT-3 Megatron-Turing (175B)NLG (530B) 100 Megatron-LM Turing-NLG Model Size (in billions of (8.3B)(17.2B)10 (11B) GPT-2 (1.5B)BERT-Large (340M) 0.1 **ELMo** (94M) 0.01 2021 2022 2018 2019 2020

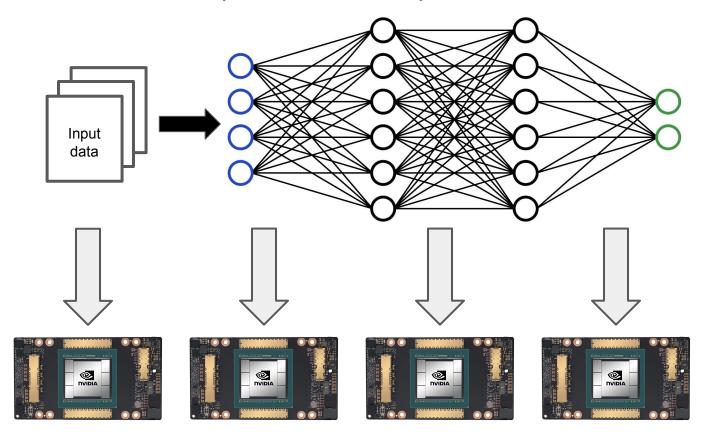
GPU FLOPS performance doubles every ~2.5 years

...but LLMs are getting ~**10x bigger every year**!

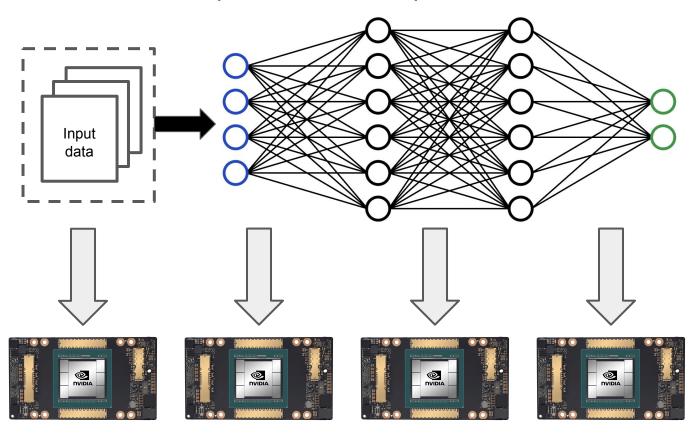
#### Parallelism is necessary to train and serve LLMs at scale

- We need to parallelize computation across multiple GPUs to handle the rapidly increasing scale of LLMs
- What are the key challenges when we start parallelizing LLMs?
  - Minimizing communication overhead: We want to minimize the time spent sending data between GPUs
  - Minimizing synchronization overhead: We want to minimize the dependencies between each GPU so that each GPU can proceed with its own computation independently
- Addressing these challenges are fundamental when it comes to designing parallel computing and distributed systems! (CS 149 and CS 244B for more detailed treatments)



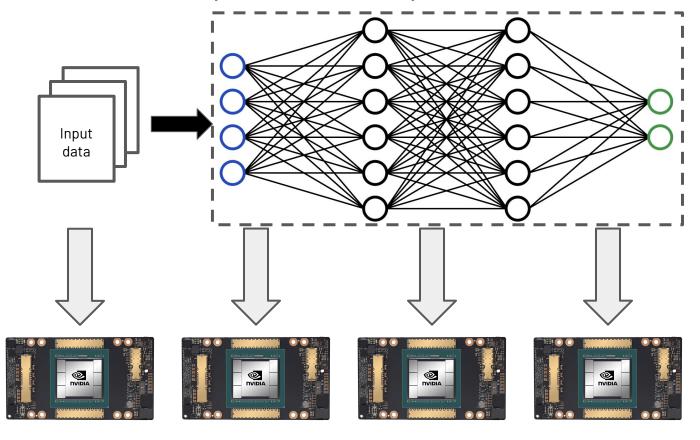


We have two options for partitioning computation across devices



We have two options for partitioning computation across devices

1. Partition the input data



We have two options for partitioning computation across devices

- 1. Partition the input data
- 2. Partition the model parameters

• There are three widely adopted techniques for parallelizing LLMs:

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  - **Data parallelism**: Partition the input data, and replicate the model weights

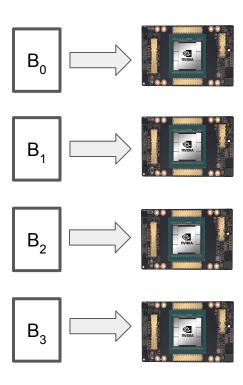
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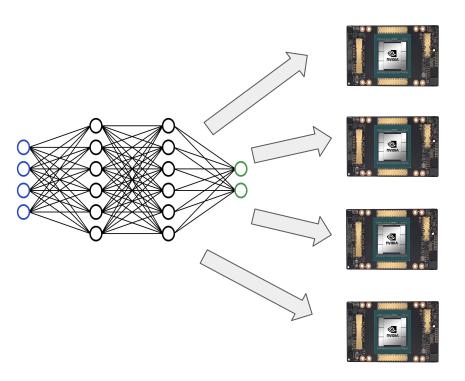
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- Each of these techniques has tradeoffs which we will discuss

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- These techniques are not mutually exclusive they can be combined to build extremely efficient systems!

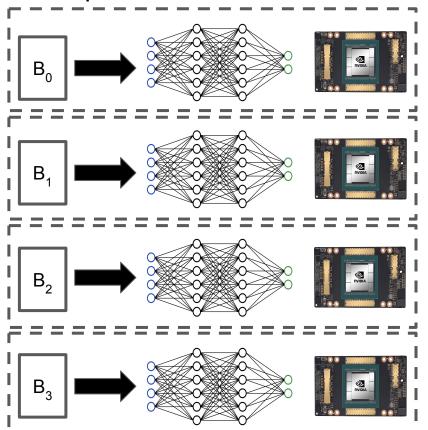
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- Each of these techniques has tradeoffs which we will discuss
- These techniques are not mutually exclusive they can be combined to build extremely efficient systems!
- These techniques are not exhaustive this is an active area of research!



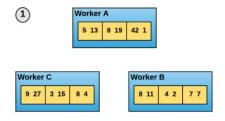
- We partition the input data on the batch dimension across the different GPUs
- For example, if we had 4 GPUs and a batch size of 64, then each GPU would receive an input sub-batch of size 16

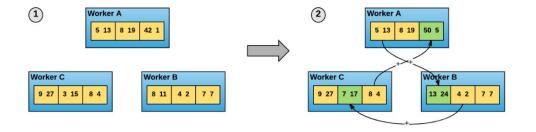


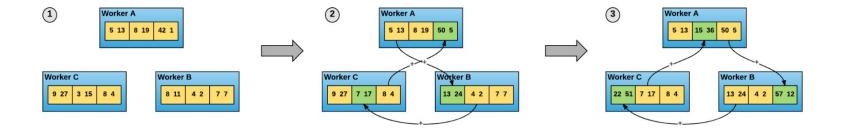
- We partition the input data on the batch dimension across the different GPUs
- For example, if we had 4 GPUs and a batch size of 64, then each GPU would receive an input sub-batch of size 16
- We replicate the model weights across the GPUs, meaning every GPU has a complete copy of all the weights

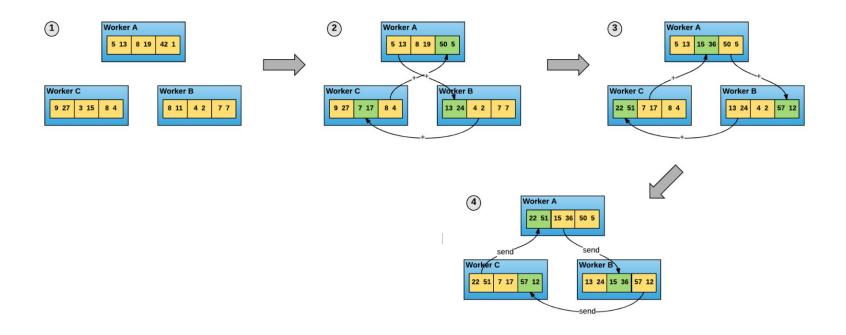


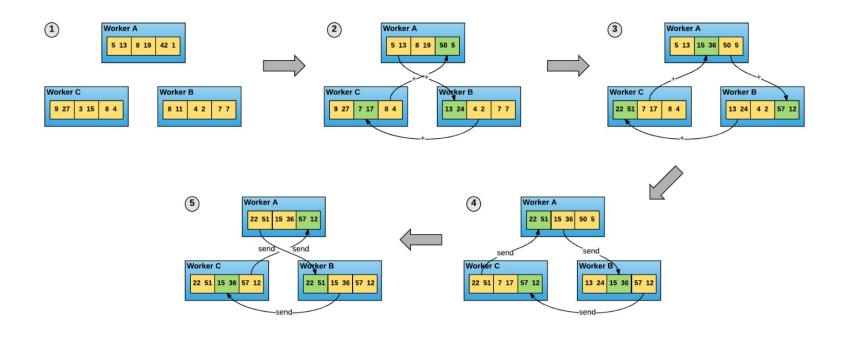
- Forward and backward pass computation on each GPU proceeds independently
- Each GPU produces its own set of gradients for its own sub-batch
- At the end of the backward pass, the gradients are aggregated across all GPUs and averaged
- The GPUs share their gradients via an all-reduce operation





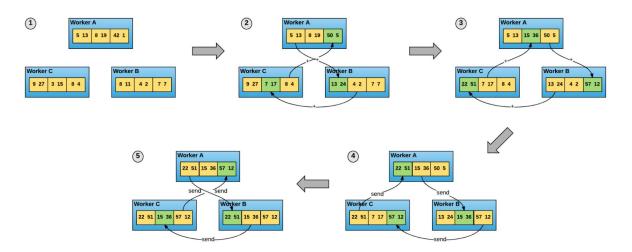






### All-reduce Algorithm

- Ring all-reduce proceeds in 2 \* (N-1) iterations for a group of N GPUs
- In each iteration, each node sends fragments of its data to 2 of its peers
  - First N-1 iterations: each GPU adds received values to its own data
  - Next N-1 iterations: each GPU replaces its own data with received values



# Ring All-reduce Efficiency Analysis

- Total volume of data sent (N is the total number of GPUs):
  - Assume each GPU has data of size X bytes
  - $\circ$  Then each node sends (2 \* (N-1)) \* X/N bytes

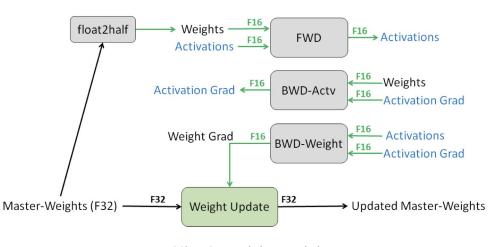
# Ring All-reduce Efficiency Analysis

- Total volume of data sent (N is the total number of GPUs):
  - Assume each GPU has data of size X bytes
  - $\circ$  Then each node sends 2 \* (N-1) \* X/N bytes
- Total time to complete all-reduce operation:
  - Assume network bandwidth of B bytes / second
  - $\circ$  Then the communication runtime is 2 \* (N-1) \* X /(N\*B) seconds

- What are the pros of data parallelism?
  - Most direct approach for increasing throughput since global batch size increases with each additional GPU
  - Relatively easy to implement (e.g. built-in support from PyTorch)

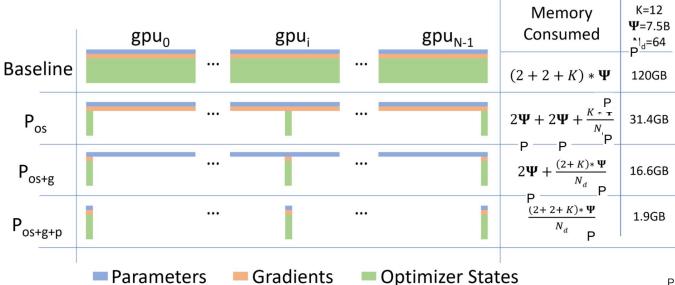
- What are the pros of data parallelism?
  - Most direct approach for increasing throughput since global batch size increases with each additional GPU
  - Relatively easy to implement (e.g. built-in support from PyTorch)
- What are the cons of data parallelism?
  - Requires the model weights and activations to fit in GPU memory (e.g. impossible to run GPT-3 with pure data parallelism on an 80 GB GPU)
  - Communication-intensive every weight must be synchronized across all GPUs

- The main limitation of data parallelism is the high memory requirement
- If we train model with P parameters using mixed-precision training:
  - o fp16 model parameters: 2P bytes
  - o fp16 gradients: 2P bytes
  - Adam optimizer states
    - fp32 model parameters: 4P bytes
    - fp32 momentum: 4P bytes
    - fp32 variance: 4P bytes
  - $\circ$  Total: 2P + 2P + 4P + 4P + 4P = 16P bytes just for parameters and optimizer states



Mixed precision training

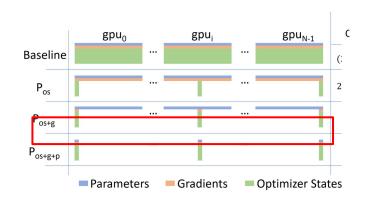
 We can significantly reduce memory usage by using the ZeRO techniques (Zero-Redundancy Optimizer)

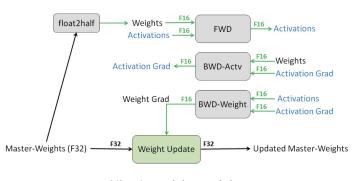


P: number of parameters

K: storage multiplier for the optimizer state

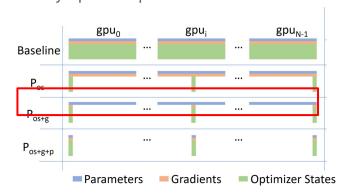
- Stage 1: Optimizer state partitioning
  - Group the optimizer states into  $N_d$  equal partitions, such that the *i*-th (data parallel) process only updates the state for the *i*-th partition



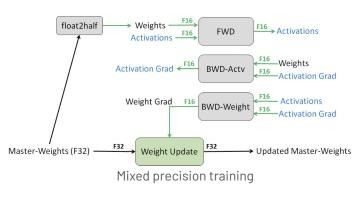


Mixed precision training

- Stage 1: Optimizer state partitioning
  - Group the optimizer states into N<sub>d</sub> equal partitions, such that the i-th (data parallel) process only updates the state for the i-th partition
  - Each process performs an all-gather across the N<sub>d</sub> processes at the end of each training step to get the fully updated parameters



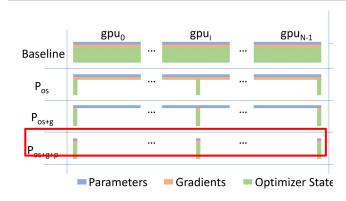
ZeRO Stage c	Memory Usage (bytes)
Baseline	16P
Stage 1: P <sub>os</sub>	4P + 12P / N <sub>d</sub>



- Stage 2: Optimizer state + gradient partitioning
  - As each data parallel process only updates its own parameter partition, it only needs the reduced gradients for the corresponding parameters
  - This reduces the memory footprint required to hold the gradients on each data parallel process from 2P bytes to 2P / N<sub>d</sub>
  - Memory savings: By removing both gradient and optimizer state redundancy, we reduce the memory footprint further down to:

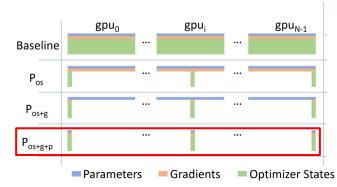
2P + 14P /	$N_{\rm d} \approx 2P$
<b>†</b>	
Weights in fp16	Gradients in fp16 and optimizer state

ZeRO Stage c	Memory Usage (bytes)
Baseline	16P
Stage 1: P <sub>os</sub>	4P + 12P / N <sub>d</sub>
Stage 2: P <sub>os+g</sub>	2P + 14 P / N <sub>d</sub>



- Stage 3: Optimizer state + gradient + parameter partitioning
  - Instead of storing all parameters on all data parallel processes, we can just fetch (i.e. gather) the parameters we need to compute the forward and backward pass
  - $\circ$  This approach increases the total communication volume by 1.5x\*, but reduces memory proportional to  $N_{\rm d}$
  - $\circ$  Memory savings: With parameter partitioning, we reduce the memory consumption 16P to 16P /  $N_d$

ZeRO Stage	Memory Usage (bytes)
Baseline	16P
Stage 1: P <sub>os</sub>	4P + 12P / N <sub>d</sub>
Stage 2: P <sub>os+g</sub>	2P + 14 P / N <sub>d</sub>
Stage 3: P <sub>os+g+p</sub>	16P / N <sub>d</sub>

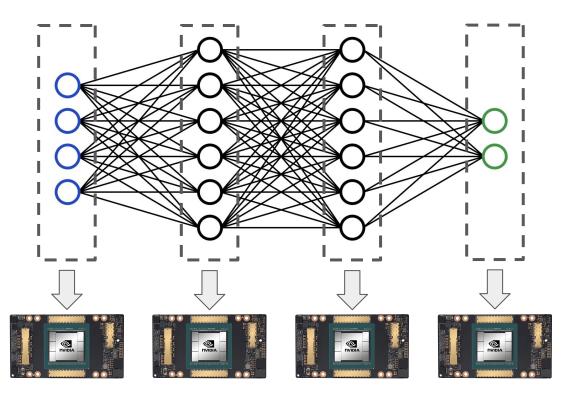


## PyTorch FSDP

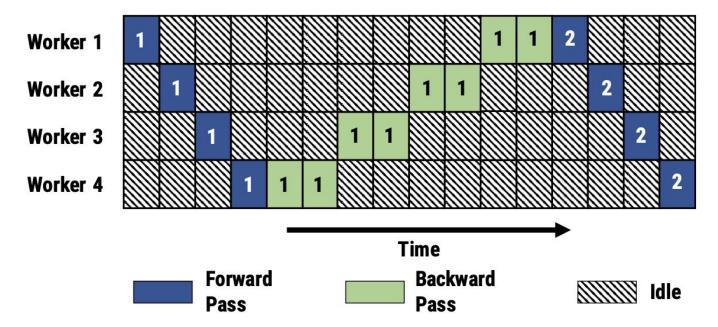
- The PyTorch FSDP (Fully Sharded Data Parallel) API implements data parallelism with ZeRO optimizations
- Example usage:

```
import torch
from torch.distributed.fsdp import FullyShardedDataParallel as FSDP
torch.cuda.set_device(device_id)
module = ... # Module definition
sharded_module = FSDP(module)
optim = torch.optim.Adam(sharded_module.parameters(), Ir=0.0001)
x = ... # Input data
y = sharded_module(x)
loss = y.sum()
loss.backward()
optim.step()
```

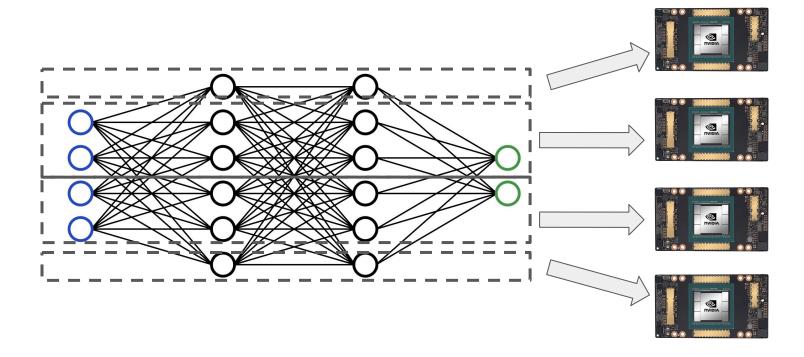
- Another way to reduce memory usage is model parallelism: replicate the input data, but partition the model weights
- Two general approaches for model parallelism:
  - Slice the model "vertically": place subsets of layers on different GPUs
  - Slice the model "horizontally": shard the model weights across different GPUs
- We will discuss the pros and cons of each these approaches



Vertically slicing the model gives each GPU its own subset of layers

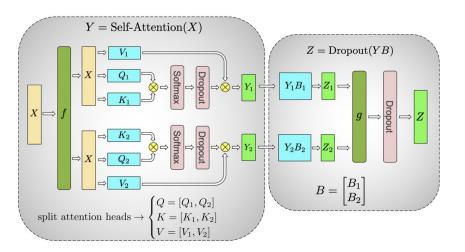


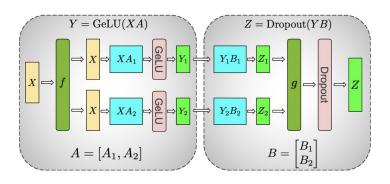
Vertical slicing severely lowers hardware utilization because the devices are frequently idle



Horizontally slicing the model shards the layers across the devices

### Tensor Model Parallelism for LLMs





- For Transformer-based LLMs specifically, we can shard the self-attention and subsequent MLP weights (Megatron-style or tensor model parallelism)
- This requires adding an all-reduce after every attention and MLP computation to synchronize the weights

#### Tensor Model Parallelism

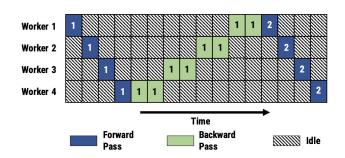
- What are the pros of tensor model parallelism (in particular Megatron)?
  - Reduces the amount of memory required per GPU
  - Keeps GPU utilization high compared to vertical slicing

#### Tensor Model Parallelism

- What are the pros of tensor model parallelism (in particular Megatron)?
  - Reduces the amount of memory required per GPU
  - Keeps GPU utilization high compared to vertical slicing
- What are the cons of tensor model parallelism (in particular Megatron)?
  - Very frequent synchronization (all-reduces) means we need extremely fast network connections to maintain high throughput
  - Not as easy to implement as data parallelism need to add the synchronization ops manually to your Attention module

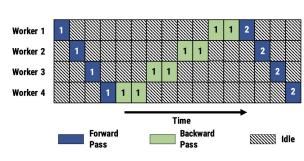
#### Tensor Model Parallelism

- Model parallelism solved our memory usage problem, but it requires very fast networking hardware due to the frequent all-reduces
- What do we do if we don't have high-memory GPUs or high-bandwidth network interconnect?
- Let's revisit the vertical slicing we saw earlier
  - This approach also reduces per-GPU memory usage
  - No all-reduces necessary just send activations or gradients from one GPU to the next
  - o Downside is poor GPU utilization can we fix this?

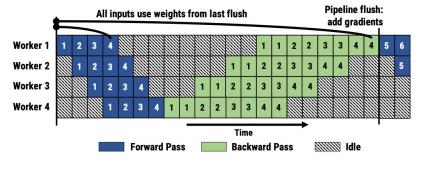


### Pipeline Parallelism

- We can apply the well-known technique of pipelining to improve GPU utilization
- Pipeline parallelism splits each batch into microbatches and injects multiple microbatches into the pipeline
- This can significantly reduce the amount of idle time on each GPU



Without pipelined computation



With pipelined computation

### Pipeline Parallelism

- Pipeline parallelism introduces two new considerations:
  - How to set the microbatch size
    - Larger microbatches = higher arithmetic intensity, smaller microbatches = smaller pipeline bubbles
  - How to decide the schedule of pipelined computation
    - We can dynamically choose which microbatches or layers to execute at each step; this will impact the pipeline bubble size

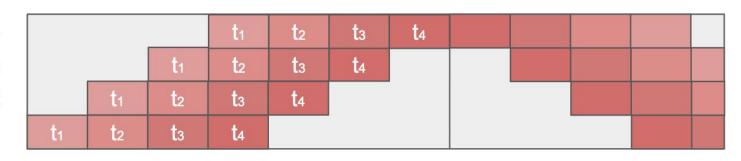




## Pipeline Parallelism for LLMs

- We can extend pipeline parallelism to LLMs by pipelining specifically on the sequence dimension
- In particular, we can split the input sequence into subsequences and have each stage process the tokens for each subsequence incrementally

GPU 4 GPU 3 GPU 2 GPU 1



#### Time

## Pipeline Parallelism

- What are the pros of pipeline parallelism?
  - Reduces the amount of memory required per GPU
  - Minimizes communication across GPUs (only point-to-point send operations instead of all-reduces)

### Pipeline Parallelism

- What are the pros of pipeline parallelism?
  - Reduces the amount of memory required per GPU
  - Minimizes communication across GPUs (only point-to-point send operations instead of all-reduces)
- What are the cons of pipeline parallelism?
  - Still suffers from low GPU utilization due to pipeline bubbles
  - Difficult to implement because it requires scheduling the different microbatches to be executed concurrently

# Summary of Different Parallelism Approaches

- In summary:
  - Data parallelism is effective if the model weights and activations fit into
     GPU memory
  - Tensor model parallelism is effective if the model weights and activations do not fit into GPU memory but we have a single server with very fast networking hardware
  - Pipeline parallelism is effective if the model weights and activations do not fit into GPU memory and we have multiple servers or a single server without fast networking hardware
- ...but we do not have to choose just one parallelism strategy!

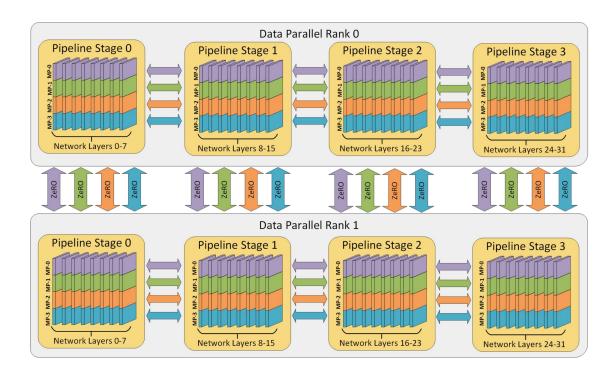
End of Lecture 8



# 3D / Pipeline-Tensor-Data (PTD) Parallelism

- We can use data

   parallelism, tensor
   model parallelism, and
   pipeline parallelism
   together (3D or PTD
   parallelism) to scale to
   thousands of GPUs
- This will be the focus of the Lecture 10

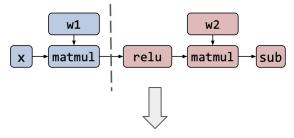


# Practical Challenges When Deploying Parallelism

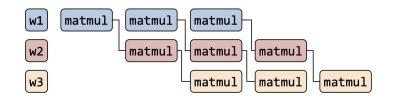
- Practical challenges when trying to deploy parallelism:
  - The space of possible strategies grows combinatorially large
  - Experimenting with different parallelization strategies is slow / expensive at the scale of hundreds or thousands of GPUs
- Can we automate parallelization for a given model and cluster?

- Alpa is a system for automatically selecting and executing the optimal parallelism strategy for a given model and cluster
- Alpa separately considers inter-operator parallelism (i.e. pipeline parallelism) and intra-operator parallelism (i.e. data and tensor model parallelism)

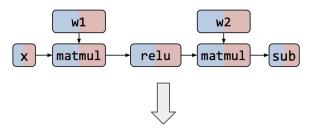
#### **Strategy 1: Inter-operator Parallelism**



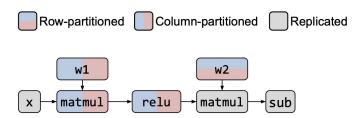
#### Pipeline the execution for inter-op parallelism



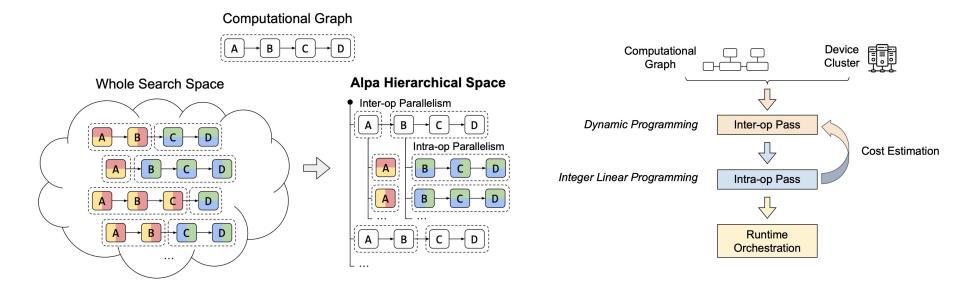
#### **Strategy 2: Intra-operator Parallelism**



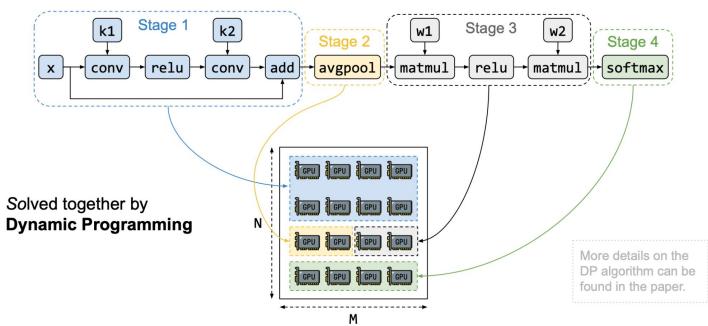
#### Multiple intra-op strategies for a single node



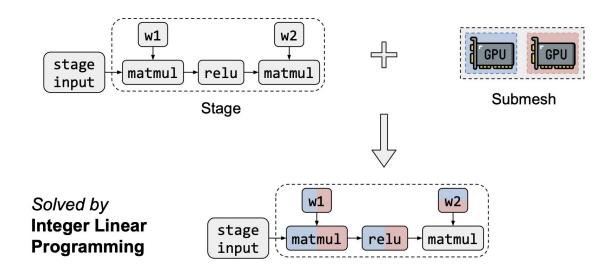
 Alpa will automatically search over the space of possible parallelism strategies for a given graph and choose the optimal combination of inter-op and intra-op parallelism



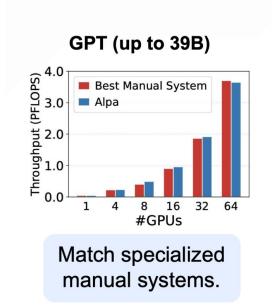
### Inter-op Pass

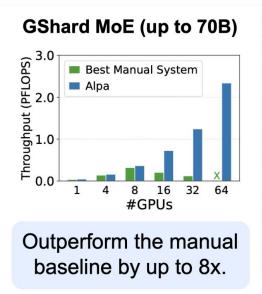


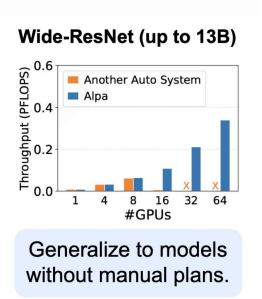
### Intra-op Pass



Stage with intra-operator parallelization







Weak scaling results where the model size grow with #GPUs. Evaluated on 8 AWS EC2 p3.16xlarge nodes with 8 16GB V100s each (64 GPUs in total).

### Conclusion

- We need to parallelize data and models to train and serve LLMs at scale
- The most widely used forms of parallelism are data parallelism, tensor model parallelism, and pipeline parallelism
- Each of these parallelization strategies has trade-offs depending on the amount of memory usage and communication overhead
- Techniques such as Alpa can help automate the decision of which parallelization strategy or strategies to use