GPipe: Easy Scaling with Micro-Batch Pipeline Parallelism

Yanping Huang, et al.

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Presenter: Constant (Sang-Soo) Park sonicstage12@naver.com
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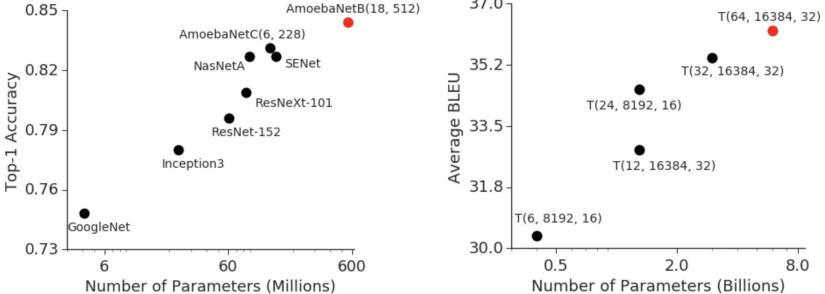
Contents of presentation

- Introduction
- Gpipe concept
 - Interface, Algorithm, Performance optimization
- Performance analysis

Bring remarkable quality improvements to several fields

- Trend: accuracy improvements on ImageNet with increase in model capacity
- Similar phenomenon be overserved in context of natural language processing

 $T(L,H,A) = transformer\ with\ L\ encoder\ and\ decoder$ (dimension of H and A attention heads) $\frac{AmoebaNetB(18,512)}{AmoebaNetC(6,228)}$



Practical challenges with larger models

- HW constraints (memory limitation, communication bandwidth on accelerator)
- **Dividing larger models** into partitions and assignment of different partitions to different accelerators^[1]

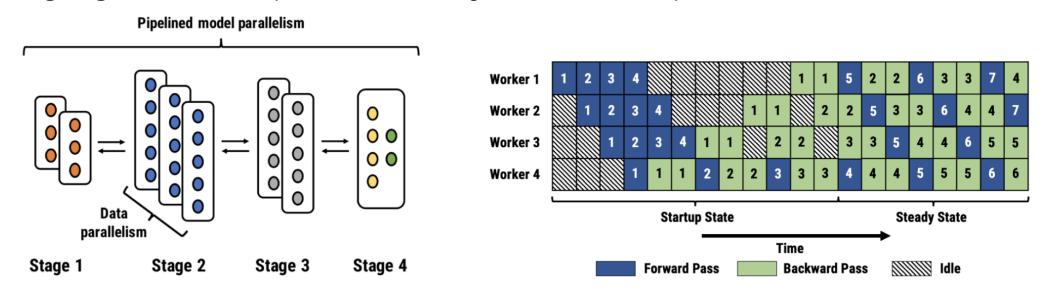
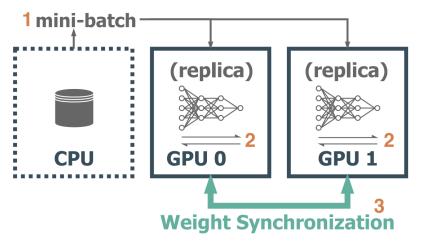


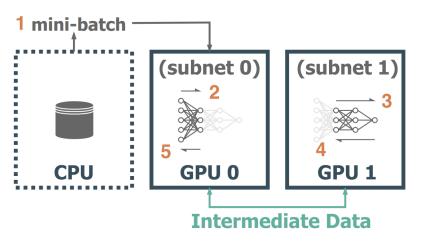
Figure 2: In the figure on the left, we show an example pipeline-parallel assignment with eight GPUs split across four stages. Only activations and gradients at stage boundaries are communicated. Stages 1, 2, and 3 have replicated stages to ensure a load-balanced pipeline. The figure on the right shows another pipeline with 4 workers, showing start-up and steady states. In this example, the backward pass takes twice as long as the forward pass.

Model parallelism

- Hard to design and implement efficient model parallelism algorithm^[2]
- Using multiple GPUs, Each GPU is responsible for weight updates of assigned model layers
- Architecture and task-specific and Increasing demand for reliable and flexible infrastructure



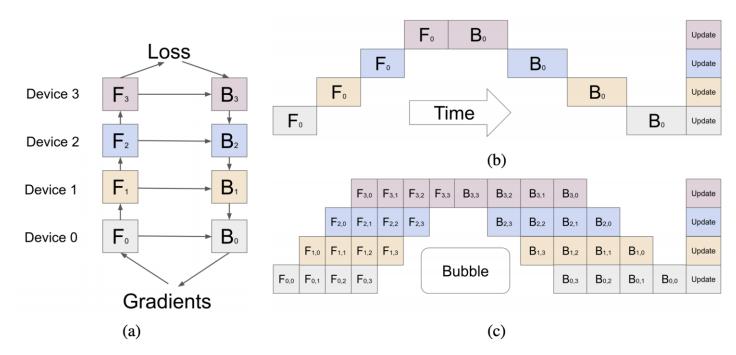




(b) Model Parallelism.

Flexible library for efficient training of large neural networks

- Scaling arbitrary DNN architecture by partitioning models, supporting re-materialization on accelerators
- Layers can be partitioned into cells and each cell is then placed on separate accelerator
- Pipeline parallelism algorithm with batch splitting (mini-batch, micro-batches)



GPipe: Interface

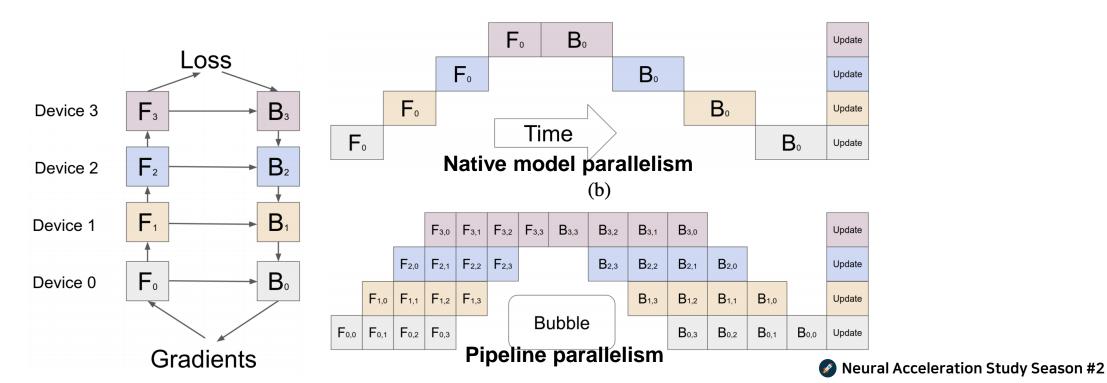
Background

- Deep neural network can be defined as sequence of *L* layers
- Each layer L_i is composed of forward computation (f_i) with set of parameters (w_i)
- Sequence of *L* layers can be partitioned into *K* composite layers, or cells
- Set of parameter (p_k) consist of consecutive layers between layers i and j.
- $p_k = [w_i, w_{i+1}, ..., w_j], F_k = f_j \circ \cdots \circ f_{i+1} \circ f_i, B_k$ can be computed from F_k , cost estimation function (c_i)
- Number of model partitions K, number of micro-batches M, sequence and definitions of L layers

GPipe: Algorithm

Network into L cells and placing cells on k^hth accelerator

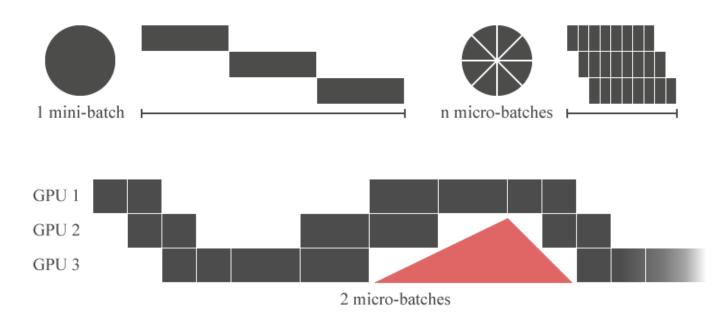
- Forward: dividing every mini-batch of size N into M equal micro-batches (pipelined through K accelerators)
- Backward, gradients for each micro-batch are computed on same model parameters used for forward
- At end of each mini-batch, gradients from *M* micro-batches are accumulated/applied to update parameters



GPipe: Algorithm

Pipeline parallelism

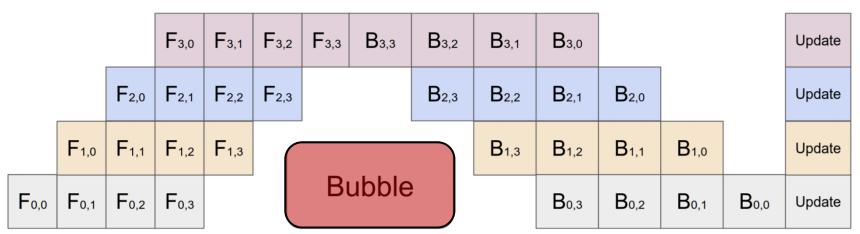
- Amount of overlapped processing increases, allowing same amount of data to be processed in less time^[3]
- As size of Micro-batch gets smaller, next GPU can start working faster (idle time gradually decreases)
- However, size is too small, efficiency is reduced (Not enough to compute workload for GPU)



GPipe: Performance optimization

Re-materialization: to reduce activation memory requirement

- Forward, each accelerator only stores output activations a partition boundaries
- Backward, k^{th} accelerator recompute composite forward function (F_k)
- **Peak activation memory** is reduced to $O(N + \frac{L}{K} * \frac{N}{M})$, $\frac{N}{M}$ (micro-batch), $\frac{L}{K}$ (# of layers per partition)
- Bubble overhead: some idle time per accelerator

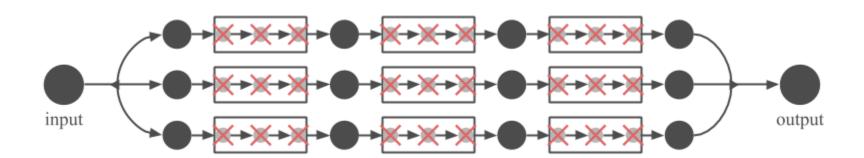


Bubble time $O\left(N + \frac{L}{K} * \frac{N}{M}\right)$ amortized over number of micro-steps M (to be negligible when $M \ge 4*K$)

GPipe: Performance optimization

Re-materialization with check-pointing

- Checkpoint: point at division of model
- Only hidden layers connecting separate models to each other is placed in memory
- In forward propagation, discard result of hidden layers inside separated models [3]
- **Peak activation memory** is reduced to $O(N + \frac{L}{K} * \frac{N}{M})$, w/o O(N * L)



Performance with two different types of models: AmoebaNet

- AmoebaNet (convolutional model), Transformer (sequence-to-sequence model)
- Both re-materialization and pipeline parallelism to benefit memory utilization
- Biggest model size GPipe can support under reasonably large input size
- 25x more memory than what is possible w/o GPipe

Table 1: Maximum model size of AmoebaNet supported by GPipe under different scenarios. Naive-1 refers to the sequential version without GPipe. Pipeline-k means k partitions with GPipe on k accelerators. AmoebaNet-D (L, D): AmoebaNet model with k normal cell layers and filter size k Transformer-L: Transformer model with k layers, 2048 model and 8192 hidden dimensions. Each model parameter needs 12 bytes since we applied RMSProp during training.

NVIDIA GPUs (8GB each)	Naive-1	Pipeline-1	Pipeline-2	Pipeline-4	Pipeline-8	_
AmoebaNet-D (L, D)	(18, 208)	(18, 416)	(18, 544)	(36, 544)	(72, 512)	
# of Model Parameters	82M	318M	542M	1.05B	1.8B	last along a distribution
Total Model Parameter Memory	1.05GB	3.8GB	6.45GB	12.53GB	24.62GB	imbalanced distribution
Peak Activation Memory	6.26GB	3.46GB	8.11GB	15.21GB	26.24GB	(model parameters)
Cloud TPUv3 (16GB each)	Naive-1	Pipeline-1	Pipeline-8	Pipeline-32	Pipeline-128	
Transformer-L	3	13	103	415	1663	_
# of Model Parameters	282.2M	785.8M	5.3B	21.0B	83.9B	
Total Model Parameter Memory	11.7G	8.8G	59.5G	235.1G	937.9G	
Peak Activation Memory	3.15G	6.4G	50.9G	199.9G	796.1G	_

Performance with two different types of models: Transformer

- AmoebaNet (convolutional model), Transformer (sequence-to-sequence model)
- Both re-materialization and pipeline parallelism to benefit memory utilization
- Biggest model size GPipe can support under reasonably large input size
- 298x more memory than what is possible w/o GPipe

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Normalized training throughput of AmoebaNet and Transformer

- # of micro-batches M is al least 5x # of partitions, bubble overhand is almost negligible
- Transformer, 3.5x speedup when it is partitioned across four times more accelerators
- AmoebaNet, sub-linear speedup due to its imbalanced computation distribution
- When M is relatively small, bubble overhead can no longer be negligible (M = 1, no pipeline parallelism)

Table 2: Normalized training throughput using GPipe with different # of partitions K and different # of micro-batches M on TPUs. Performance increases with more micro-batches. There is an almost linear speedup with the number of accelerators for Transformer model when $M \gg K$. Batch size was adjusted to fit memory if necessary.

TPU	AmoebaNet			Tr	ansform	ner
K =	2	4	8	2	4	8
M = 1	1	1.13	1.38	1	1.07	1.3
M = 4	1.07	1.26	1.72	1.7	3.2	4.8
M = 32	1.21	1.84	3.48	1.8	3.4	6.3

Table 3: Normalized training throughput using GPipe on GPUs without high-speed interconnect.

GPU	AmoebaNet			Tı	ansfor	mer
K =	2	4	8	2	4	8
M = 32	1	1.7	2.7	1	1.8	3.3

Image classification

- AmoebaNet with scaled input image size (480x480)
- Single model: 4 partitions, 84.4\$ (top-1), 97% (top-5)
- Effectiveness of giant convolution networks on other image datasets through transfer learning

Table 4: Image classification accuracy using AmoebaNet-B (18, 512) first trained on ImageNet 2012 then fine-tuned on others. Please refer to the supplementary material for a detailed description of our training setup. Our fine-tuned results were averaged across 5 fine-tuning runs. Baseline results from Real *et al.* [12] and Cubuk *et al.* [26] were directly trained from scratch. *Mahajan *et al.*'s model [27] achieved 85.4% top-1 accuracy but it was pretrained on non-public Instagram data. Ngiam *et al.* [28] achieved better results by pre-training with data from a private dataset (JFT-300M).

Dataset	# Train	# Test	# Classes	Accuracy (%)	Previous Best (%)
ImageNet-2012	1,281,167	50,000	1000	84.4	83.9 [12] (85.4*[27])
CIFAR-10	50,000	10,000	10	99.0	98.5 [26]
CIFAR-100	50,000	10,000	100	91.3	89.3 [26]
Stanford Cars	8,144	8,041	196	94.6	94.8 * [26]
Oxford Pets	3,680	3,369	37	95.9	93.8* [29]
Food-101	75,750	25,250	101	93.0	90.4* [30]
FGVC Aircraft	6,667	3,333	100	92.7	92.9 * [31]
Birdsnap	47,386	2,443	500	83.6	80.2* [32]

Massive Massively Multilingual Machine Translation

- Scaled transformer: (1) depth by increasing number of layers, (2) width by increasing hidden dimensions
- 1.3B wide model with T(12,16384,32), 1.3B deep model T(24,8192,16)
- Quality of both model is similar, deeper model outperforms by huge margins on low-resource languages (suggesting that increasing model depth might be better for generalization)

Figure 3: Translation quality across all languages with increasing multilingual model capacity. Languages are arranged in the order of decreasing training dataset size from left to right. T(L,H,A), depicts the performance of a Transformer with L encoder and L decoder layers, a feed-forward hidden dimension of H and A attention heads. We notice that increasing the model capacity, from 400M params (T(6,8192,16)) to 1.3B (T(24,8192,16)), and further, to 6B (T(64,16384,32)), leads to significant quality improvements across all languages. We also notice huge quality improvements for low-resource languages (right side of the plot), when compared against bilingual baselines, highlighting the significant transfer gains resulting from training a multilingual model.

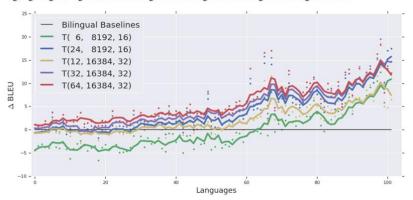


Table 5: The Effect of Batch Size

Batch Size	260K	1 M	4M
BLEU	30.92	31.86	32.71
Loss (NLL)	2.58	2.51	2.46

Thank you