

SVD-LoRA: Singular Value Decomposition Compression of Low Rank Adaptations

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Introduction

- Model sizes have escalated in recent years; bigger models often perform better.
- Scaling both training data and model parameters improves performance.
- Larger models limit AI research for those without massive computational resources.
- Existing strategies for scaling include quantization, pruning, and distillation.

Related Work

- *Quantization*: Reduces weight precision, but might decrease model performance.
- *Pruning*: Removes redundant connections; requires fine-tuning.
- Low Rank Adaptation: Represents weights in lower-dimensional space.
- *Distillation*: Trains a smaller model to mimic a larger one's outputs.

Methodology

Conclusion

Introduced SVDLoRA method for model compression.

Demonstrated its potential for model robustness and usability.

Future Work

- Further studies are required.
- Potential application with Switch Transformers and Mixture of Experts models.
- Compare SVDLoRA with other compression methods.
- Theoretical explorations into the lower bound of compression for neural network weights.

- Use SVD to represent original neural network weights with low-rank approximations.
- Represent weight matrix with two separate low-rank matrices.
- Construct a difference matrix between original matrix and SVD. Use SVD to compress the difference matrix.
- The compressed difference matrices are added to the existing SVD low-rank matrices. (Inspired by the model saving practices of the Stable Diffusion community)
- This step is repeated until a required number of submatrices is generated, with each new difference matrix is the result of the reconstructed matrix (using all SVD steps) with the original matrix
- Approach reduces storage and computational complexity, while preserving model quality.
- Methodology can be generalized to use multiple matrices.

Experiments

- Tested on CIFAR-10 and NMT datasets.
- Different matrix counts and ranks were evaluated.
- Results vary but show potential.

| Submatrices | Eval Accuracy | Eval Runtime |
|-------------|----------------------|---------------------|
| Baseline | 0.9769 | 131.2141 |
| 1 | 0.9571 | 129.2873 |
| 2 | 0.962 | 128.7298 |
| 3 | 0.9617 | <u>129.0121</u> |
| 4 | <u>0.9622</u> | 129.2541 |
| 5 | 0.9619 | 129.3435 |
| 6 | 0.9611 | 129.3593 |
| 7 | 0.9588 | 129.4201 |
| 8 | 0.9613 | 129.3076 |
| 9 | 0.9612 | 130.1609 |

Table 1. CIFAR-10 accuracy across 1-9 sub-matrices with a fixed number of parameters

Parameter % Eval Accuracy Eval Runtime

| Baseline | 0.9769 | 131.2141 |
|----------|---------------|-----------------|
| 10% | 0.9617 | 130.4624 |
| 20% | 0.9746 | 128.9813 |
| 30% | 0.9766 | <u>129.0198</u> |
| 40% | 0.9763 | 129.067 |
| 50% | 0.9767 | 129.135 |
| 60% | 0.9767 | 129.3274 |
| 70% | <u>0.9768</u> | 129.5422 |
| 80% | 0.9764 | 129.5246 |
| 90% | 0.9759 | 129.4248 |
| 100% | 0.9759 | 129.7098 |

Table 2. CIFAR-10 Accuracy across 10-100% parameters at 3 sub-matrices

| Factor | Mean | | |
|--------------|---------|---------|--|
| | BLEU | chrf | |
| Baseline | 30.0678 | 58.495 | |
| 30% (r=102) | 19.8637 | 41.0294 | |
| 40% (r=136) | 31.0818 | 55.9282 | |
| 50% (r=170) | 30.437 | 58.7765 | |
| 60% (r=204) | 29.8488 | 58.4441 | |
| 70% (r=238) | 28.7292 | 57.9683 | |
| 80% (r=272) | 28.392 | 57.7862 | |
| 90% (r=306) | 29.4163 | 58.3949 | |
| 100% (r=340) | 29.6074 | 58.3529 | |

Table 3. Mean (5 salt language) Evaluation Metrics at different parameter counts

Unsuccessful Directions

• Attempts to directly parameterize a randomly initalized lower rank matrices with backpropagation or

evolutionary optimization were unsuccessful.

• A loss of mean squared error between reconstructed matrix and original matrix as well as difference between norms and difference between singular values did not lead to convergence