

# Thesis Notepad of Ideas and Progress etc

June 22, 2025



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# Chapter 1

## Introduction

**TODO: Ask about whole story telling element and how i should frame the headless chicken with failing ideas approach i regreably followed**



# Chapter 2

## Understanding Adam

We discuss some of the underlying theory, review some important literature and run some experiments with the aim of understanding the dynamics of AdamW and the role which certain design choices play in Adam’s efficacy.

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**Algorithm 1** Adam Optimizer (Bias Correction Experimental Focus)

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**Require:**  $(\eta_t)_{t \geq 1} \subset \mathbb{R}_{\geq 0}, \beta_1, \beta_2 \in [0, 1), \epsilon \in \mathbb{R}_{>0}$  ▷ Standard hyperparameters  
**Require:**  $\theta_0$  ▷ Initialised model weights  
**Require:** `do_bias_correction`: bool ▷ **Experimental flag**

- 1: Initialize  $m_0, v_0, t \leftarrow 0$
- 2: **while**  $\theta_t$  not converged **do**
- 3:      $t \leftarrow t + 1$
- 4:      $g_t \leftarrow \nabla_{\theta} f_t(\theta_{t-1})$
- 5:      $m_t \leftarrow \beta_1 m_{t-1} + (1 - \beta_1) g_t$
- 6:      $v_t \leftarrow \beta_2 v_{t-1} + (1 - \beta_2) g_t^2$
- 7:     **if** `do_bias_correction` **then**
- 8:          $\hat{m}_t \leftarrow \frac{m_t}{1 - \beta_1^t}, v_t \leftarrow \frac{v_t}{1 - \beta_2^t}$  ▷ bias correction not by default
- 9:     **end if**
- 10:      $\theta_t \leftarrow \theta_{t-1} - \eta_t \frac{m_t}{\sqrt{v_t} + \epsilon}$
- 11: **end while**

---

Experiments

### 2.1 Why is AdamW the Top Dog?

The Adam optimizer [KB17] was first published just over a decade ago. With the integral addition of decoupled weight decay [LH19], it has been the de facto stochastic optimizer choice for nearly all SOTA deep learning training, particularly in language model pre-training. Extensive research into understanding Adam’s dynamics has been conducted however several key aspects remain not completely understood.

The optimizer itself is simply a synthesis of two already well-established optimization ideas - momentum and RMSProp. Adam enjoys the upsides of both methods and has proved to be a powerful and robust c

Despite extensive research into optimizers for deep learning in the past decade, AdamW remains the default choice for the vast majority of both industry and academic applications.

While a substantial number of publications claim great and consistent

## 2.2 Preliminaries and Related Work

**TODO: Talk about the papers about understanding Adam and then a bit about what is stil not understood/ bottlenecks/is it really the best we can do**

### 2.2.1 EAdam Paper probably

### 2.2.2 Relation to signSGD

### 2.2.3 Why transformers need Adam: hessian and class imbalance

**TODO: Talk about this latest batch size SGD paper from group that challenges these prior explanations**

### 2.2.4 The Secret Sauce paper

## 2.3 Some Experiments/Ablations (Mostly an experiment graveyard sadly)

Need a lot of sensitivity curves for models in vision and language setting.

- Nested moving average vs regular Adam: are they the same? Interesting since the choice of whichway to take the EMA is sort of arbitrary
- Adam2SGD: Could look at SGD2Adam since the paper from Teodora showed some advantage in doing so
- Epsilon scheduling experiments: could be very related to the sgd thing... Can also be bespoke to parameter (seems like a universal epsilon for all parameters could be foolish.
- Beta scheduling too with experiments about the angle between the two
- The BACKPack package experiment with class variances individual. Relationship to signSGD stuff from Hennig paper.



### 2.3.1

### 2.3.2 Nesting the Moving averages

#### Coupling of momentum and RMSProp

The momentum and RMS boil down to keeping track of a moving average ( $\text{EMA}_{\beta}$ ) parametrized by  $\beta \in \mathbb{R}_{\geq 0}$ .

While regular adam is given by:

$$w^{t+1} = w^t - \eta \frac{\text{EMA}_{\beta_1}(g_t)}{\sqrt{\text{EMA}_{\beta_2}(g_t^2) + \varepsilon}}$$

one could also consider the double nesting formula

$$w^{t+1} = w^t - \eta \text{EMA}_{\beta_1} \left( \frac{g_t}{\sqrt{\text{EMA}_{\beta_2}(g_t^2) + \varepsilon}} \right)$$

Or expressed in two steps:

$$\begin{aligned} v^t &= \beta_2 v^{t-1} + (1 - \beta_2) g_t \odot g^t \\ K^t &= \beta_1 K^{t-1} + (1 - \beta_1) \frac{g_t}{\sqrt{v^t + \varepsilon}} \end{aligned}$$

Of course, the

When doing this, the bias correction term becomes less clear cut. Of course the second moment estimate  $v^t$  is the same ( $1 - \beta_1^t$  if you believe  $E[g_t]$  is similar on initialization). But after the nested average we have a sum of the form

$$K^t = (1 - \beta_1) \sum_{j=0}^{t-1} \beta_1^{t-i-1} \frac{g_j}{\sqrt{v_j + \varepsilon}}$$

Can we make the same kind of assumption for initial values of  $t$ ? Namely that  $E[\frac{g_j}{\sqrt{v_j + \varepsilon}}]$  is "close enough" for all small  $j$  so that we can reduce the ugly expression above to a geometric sum. If so, great the term is the same. If not, any other clever tricks? Indeed as we demonstrate in detail in 3 that bias correction is currently poorly understood and is generally not required if proper learning rate scheduling is implemented.

### 2.3.3 Adam2SGD and $\varepsilon$ schedule

**TODO: Ask Antonio if this is just fucked/should go in the appendix/nowhere**

### 2.3.4 How EMA smooths the update signal

**TODO: Add this angle between gradients and updates vs update and previous update experiment with plots and actually do the beta scheduling thing you keep thinking/talking about!**

### 2.3.5 BackPACK

**TODO: Talk about the shitty variance KL divergence experiment to see if  $v_t$  is somehow a proxy for variance. Also the whole optimal batch size scaling based on the variance of the mini-batch**

### 2.3.6 The $\varepsilon$ Hyperparameter

While regular adam is given by:

$$w^{t+1} = w^t - \eta \frac{\text{EMA}_{\beta_1}(g_t)}{\sqrt{\text{EMA}_{\beta_2}(g_t^2) + \varepsilon}}$$

with  $\varepsilon = 1e - 8$  as the default choice.

It has been observed in a number of papers ([YG20]), that language model pretraining can be improved by altering the epsilon hyperparameter. Indeed increasing  $\varepsilon$  to  $1e - 6$  can decrease final validation perplexity but at the cost of decreased training stability. Conversely, decreasing  $\varepsilon$  can improve stability without noticable decrease in performance. In every one of these experiments, the  $\varepsilon$  value is chosen to apply globally, however the We consider the distribution of

# Chapter 3

## Bias Correction in Adam: A Detailed Analysis

### 3.1 Bias Correction: An Unnecessary Evil?

The AdamW optimizer can be represented as follows: ( **TODO: Probably smneed to replace this with the algorithm itself. Then can do some kind of "if do bias correction" condition:**

$$\begin{aligned} m^{t+1} &= \beta_1 m^t + (1 - \beta_1) g^t & , \quad v^{t+1} &= \beta_2 v^t + (1 - \beta_2) g^t \odot g^t \\ \hat{m}^{t+1} &= \frac{1}{1 - \beta_1^{t+1}} m^{t+1} & , \quad \hat{v}^{t+1} &= \frac{1}{1 - \beta_2^{t+1}} v^{t+1} \\ w^{t+1} &= (1 - \eta_t \alpha) w^t - \eta_t \frac{\hat{m}^{t+1}}{\sqrt{\hat{v}^{t+1}} + \epsilon} \end{aligned}$$

Where  $\eta_t \in \mathbb{R}_{\geq 0}$  is the learning rate at time step  $t$ ,  $\alpha \in \mathbb{R}_{\geq 0}$  is the weight decay and  $\beta_1, \beta_2, \epsilon$  are the model's hyperparameters.

One observes in a large number of research papers (gotta find some nice examples), an extensive discussion into the role of bias correction. Indeed bias correction is generally viewed by the optimization community as an inexorable component of the Adam(W) optimizer

The purpose of this chapter is to discuss the role of bias correction on the performance of models trained using AdamW. For the purposes of all experiments conducted, we make use of a custom implementation of the AdamW optimizer.

#### 3.1.1 A Discussion of the "Proof" for Bias Correction

In a number of pedagogical resources (cite Goodfellow, Geiger lecture series), blogs and indeed seminal research papers - some variant of the following "proof" is given to justify

the inclusion of bias correction:

$$\begin{aligned}\mathbb{E}[m_t] &= \mathbb{E} \left[ (1 - \beta_1) \sum_{j=1}^t \beta_1^{t-j+1} g_j \right] \\ &= \mathbb{E} \left[ (1 - \beta_1) g_t \sum_{j=1}^t \beta_1^{t-j+1} \right] \text{ (assuming } \mathbb{E}[g_t] \approx \mathbb{E}[g_i] \forall i < t) \\ &= (1 - \beta_1^t) \mathbb{E}[g_t]\end{aligned}$$

It is therefore argued that dividing by  $1 - \beta_1^t$  removes the expected "bias" in the exponential moving average.

Where an analogous argument is used to justify the bias correction factor for  $v_t$ .

This assumption that  $\mathbb{E}[g_t] \approx \mathbb{E}[g_i]$  for  $i < t$  is blatantly false. We argue through a number of experiments with a custom implementation of AdamW that the bias correction step can be removed from AdamW with no adverse effects.

**TODO: some visualiation or way to make this look nice**

## 3.2 Experimental Outline

We conduct a number of experiment in both a vision and language setting.

In the vision setting, we consider:

- A multi-layered perceptron (MLP) and simple convolutional neural network (CNN) on the MNIST, Fashion MNIST and SVHN.
- A less overparametrized ResNet implementation on the CIFAR10 dataset (only  $\sim 1\text{M}$  parameters rather than ResNet 18 which has  $\sim 11\text{M}$ )
- Both a ResNet56<sup>1</sup>, Densenet121 [HLvdMW18] and a vision transformer on the CIFAR100 dataset
- Both a ResNet50 and a vision transformer on the Tiny Imagenet.

We consider both the setting with a warmup followed by cosine decay learning rate schedule and constant learning rate.

For both CIFAR-10/100, standard data augmentation techniques were implemented along with a tuned weight decay hyper-parameter.

In the case of pretraining language models, we restrict our attention to transformer based models. In particular we make use of the [following enhanced implementation](#) of nanoGPT ([Kar22]) including RMSNorm (instead of batch/layer normalization), SwiGLUi [Sha20] and Rotary Positional Embedding

For each model, we run a sweep over a range of learning rates and

---

<sup>1</sup>Pytorch's ResNet50 was specifically designed for Imagenet's  $224 \times 224$  images and applies aggressive downsampling in the early layers. When applied to CIFAR-100's  $32 \times 32$  images, important information is then lost and . We therefore make use of a commonly-used specialised ResNet architecture [Ide]

We note that the exact hyperparameter configuration used for each experiment is not of integral importance in distinguishing the performance of AdamW with and without bias correction (and indeed zero-initialisation). In each case, we run a sweep to ensure a model of near SOTA performance but this is mainly for vanity and we also consider results across a wide range of hyperparameters to prove the performance similarities are consistent across the space of all hyperparameter configurations.

## Bias Correction isn't the End of the Story

When initially implementing the bias correction-free version of AdamW, we assumed that initialising the moments  $m_t, v_t$  with the gradients  $g_0, g_0^2$  respectively would be an effective way to remove any bias - thus eliminating the need for bias correction. However, we discovered that this is not quite as simple as initially expected. While initialising the moments as zero certainly induces a bias, it is not the same bias that is corrected by bias correction.

We write down, in very explicit terms, the update step for the first and then all subsequent steps) in for each of the four possibilities (zero init, bias correction)  $\in$  (True, False) Consider the following closed form expression for the exponential moving averages:

$$m_t = \beta_1^t m_0 + (1 - \beta_1) \sum_{j=1}^t \beta_1^{t-j} g_j$$

We consider the four possible configurations of ZI and BC for the very first step of the optimizer:

$g_1$  is computed from the first batch passed in, then we have:

$$m_0 = \begin{cases} 0, & \text{if ZI} \\ g_1, & \text{else} \end{cases}$$

Then the EMA update occurs with the same grad ( $g_1$  is used in first update too)

We have  $m_1 = \beta_1 m_0 + (1 - \beta_1) g_1$

$$m_1 = \begin{cases} (1 - \beta_1) g_1, & \text{if ZI} \\ \beta_1 g_1 + (1 - \beta_1) g_1 = g_1, & \text{else} \end{cases}$$

Then bias correction is either applied or not. Therefore we have:

$$\hat{m}_1 = \begin{cases} g_1, & \text{if ZI and BC} \\ (1 - \beta_1) g_1, & \text{if ZI and no BC} \\ \frac{1}{1 - \beta_1} g_1, & \text{if no ZI and BC} \\ g_1, & \text{if no ZI and no BC} \end{cases}$$

By the very same logic, we have

$$\hat{v}_1 = \begin{cases} g_1^2, & \text{if ZI and BC} \\ (1 - \beta_2)g_1^2, & \text{if ZI and no BC} \\ \frac{1}{1 - \beta_2} g_1^2, & \text{if no ZI and BC} \\ g_1^2, & \text{if no ZI and no BC} \end{cases}$$

Thus the first optimizer step is given by (denote the unit vector in the direction of  $g_1$  by  $\hat{g}_1$  and these expressions are vectorized over all  $g_1^{(i)}$ ):

$$\Rightarrow \text{first step} = s_1 = \frac{\hat{m}_1}{\sqrt{\hat{v}_1}} = \begin{cases} \frac{g_1}{\sqrt{g_1^2 + \epsilon}} = \frac{g_1}{|g_1| + \epsilon} = \frac{1}{1 + \epsilon/|g_1|} \text{sign}(g_1), & \text{if ZI and BC} \\ \frac{(1 - \beta_1)g_1}{\sqrt{1 - \beta_2} |g_1| + \epsilon} = \frac{1 - \beta_1}{\sqrt{1 - \beta_2}} \frac{1}{1 + \frac{\epsilon}{|g_1| \sqrt{1 - \beta_2}}} \text{sign}(g_1) & \text{if ZI and no BC} \\ \frac{\frac{1}{1 - \beta_1} g_1}{\frac{1}{\sqrt{1 - \beta_2}} |g_1| + \epsilon} = \frac{\sqrt{1 - \beta_2}}{1 - \beta_1} \frac{1}{1 + \frac{\epsilon \sqrt{1 - \beta_2}}{|g_1|}} \text{sign}(g_1), & \text{if no ZI and BC} \\ \frac{1}{1 + \epsilon/|g_1|} \text{sign}(g_1), & \text{if no ZI and no BC} \end{cases}$$

The direction of the Adam update is then the same for all cases but the magnitude varies substantially. So the magnitude of the first gradient (which is basically random in some capacity since the weights are randomly computed and there is just one noisy batch of gradient data to go by) basically determines the direction of the first step.

What matter is the order of magnitude of  $|g_1|$  relative to For the case no ZI and BC, basically the factor  $\frac{\sqrt{1 - \beta_2}}{1 - \beta_2}$  is making the gradient substantially larger (depends somewhat on values of  $\beta_1, \beta_2$ ).

From the general formula:

$$m_t = \beta_1^t m_0 + (1 - \beta_1) \sum_{j=1}^t \beta_1^{t-j} g_j \quad \text{and} \quad \hat{m}_t = \frac{1}{1 - \beta_1^t} m_t \text{ if BC else } m_t$$

$$\text{and } (1 - \beta_1^t) = (1 - \beta_1) \sum_{j=1}^{t-1} \beta_1^j$$

we consider the four cases with ZI and BC (let  $B := \sum_{j=1}^{t-1} \beta_1^j$ :

$$\hat{m}_t = \begin{cases} \frac{1}{B} \sum_{j=1}^t \beta_1^{t-j} g_j & \text{if ZI and BC} \\ (1 - \beta_1) \sum_{j=1}^t \beta_1^{t-j} g_j & \text{if ZI and no BC} \\ \frac{\beta_1^t}{(1-\beta_1)^B} g_1 + \frac{1}{B} \sum_{j=1}^t \beta_1^{t-j} g_j & \text{if no ZI and BC} \\ \beta_1^t g_1 + (1 - \beta_1) \sum_{j=1}^t \beta_1^{t-j} g_j & \text{if no ZI and no BC} \end{cases}$$

why not just divide by  $1 - \beta_2$ ? rather than  $1 - \beta_2^t$ ? makes no difference up to scaling

### 3.3 Is Bias Correction just Hidden Learning rate scheduling?

In our experiments comparing adam with and without bias correction, we found that certain choices of the pair  $\beta_1, \beta_2$  caused a much greater discrepancy in performance, but only when no learning rate schedule was used. This lead to the following insight.

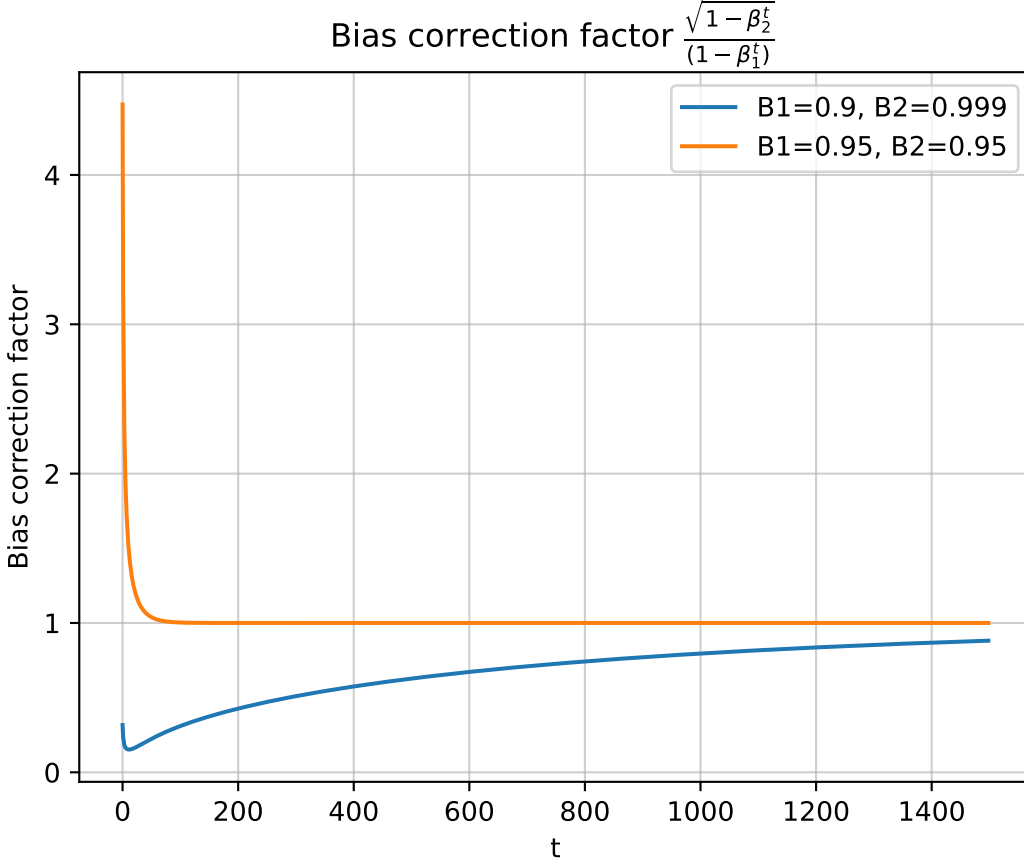
Consider the following factorization (we will ignore the  $\varepsilon$  in the denominator for simplicity)<sup>2</sup>

$$\frac{\hat{m}_t}{\hat{v}_t} = \frac{\frac{1}{1-\beta_1^t} m_t}{\sqrt{\frac{1}{1-\beta_2^t} v_t}} = \frac{\sqrt{1-\beta_2^t}}{1-\beta_1^t} \frac{m_t}{\sqrt{v_t}} = \rho(t; \beta_1, \beta_2) \frac{m_t}{\sqrt{v_t}} \quad (3.1)$$

The behaviour of  $\rho(t; \beta_1, \beta_2) := \frac{\sqrt{1-\beta_2^t}}{1-\beta_1^t}$  which we refer to as the *bias-correction factor* depends highly dependent on the choice of  $\beta_1, \beta_2 \in [0, 1)$   
Consider the following plot:

---

<sup>2</sup>We note that if the  $\varepsilon$  term is included, 3.1 evaluates to  $\frac{\sqrt{1-\beta_2^t}}{1-\beta_1^t} \frac{m_t}{\sqrt{v_t + \varepsilon \sqrt{1-\beta_2^t}}}$ . Since  $\sqrt{1-\beta_2^t} \in [0, 1] \forall t \in \mathbb{N} \forall \beta_2 \in [0, 1)$ , bias correction should have minimal effect



For the pair  $(\beta_1, \beta_2) = (0.9, 0.999)$ , the factor resembles a very steady warmup whereas for the pair  $(\beta_1, \beta_2) = (0.95, 0.95)$ , the factor is merely a large spike which quickly converges to 1

Consider the derivative of  $\rho$  with respect to  $t$ :

$$\frac{d\rho}{dt} = \frac{-\frac{\beta_2^t \log \beta_2 (1 - \beta_1^t)}{2\sqrt{1 - \beta_2^t}} + \sqrt{1 - \beta_2^t} \beta_1^t \log \beta_1}{(1 - \beta_1^t)^2}$$

where  $\log$  denotes the natural logarithm.

Since  $\log \beta_i < 0$  for any  $\beta_i \in (0, 1)$ , we note that  $\rho$  has positive  $t$ -derivative if and only if

$$\frac{\beta_2^t \log(\beta_2)}{2(1 - \beta_2^t)} < \frac{\beta_1^t \log(\beta_1)}{1 - \beta_1^t}$$

As we can see, in the case of  $(\beta_1, \beta_2) = (0.95, 0.95)$ , the derivative is *never* positive. It appears that what is far more important than the inclusion of bias correction is rather the initialization of the moments. One would naively assume that initializing the gradients to what they actually are should outperform setting them to zero. However there are minor improvements to initialising at zero



Table 3.1: *ZI* denotes Zero init, *BC* denotes Bias Correction. Not doing *ZI* means we initialize  $m$  and  $v$  at  $g_0$  and  $g_0^2$  respectively. Default for AdamW is *ZI* and *BC*. Performing bias correction is not as important as initialization in Adam. Averaged results over 4 random seeds

HPs lr:0.008,  $\beta_1 : 0.95$ ,  $\beta_2 : 0.95$ ,  $wd : 0.1$

|         | AdamW            | AdamW no BC, ZI  | AdamW BC, no ZI  | AdamW no BC, no ZI |
|---------|------------------|------------------|------------------|--------------------|
| Val ppl | $21.87 \pm 0.11$ | $21.93 \pm 0.04$ | $22.83 \pm 0.15$ | $22.64 \pm 0.13$   |



# Chapter 4

## Beyond Adam: Is Muon the future?

While first-order optimization methods such as AdamW are currently the de-facto choice for training deep neural networks, In recent years, there have been a number of papers which attempt to improve upon Adam(W) by leveraging information about the structure of individual layers.

Newton’s method, which requires explicit computation of the inverse-Hessian matrix, is rarely sufficiently efficient, or even tractable, in deep learning. Similarly, Natural Gradient Descent [Ama98] requires computation of the inverse-Fischer information matrix

While there has been a myriad of literature Let  $f : \mathcal{X} \times \Theta \rightarrow \mathcal{Y}$  with  $\Theta \subset \mathbb{R}^p$  denote an arbitrary deep neural network and let  $L : \mathcal{X} \times \Theta \times \mathcal{Y} \rightarrow \mathbb{R}_{\geq 0}$  denote a (continuously differentiable) loss function.

The textbook definition of most traditional stochastic optimization algorithms involves initialising some  $\theta_0 \in \Theta$  and then updating with some rule

$$\theta_{t+1} = \theta_t - \eta_t s_t$$

for some  $s_t \in \mathbb{R}^d$  which is typically a function of the previous stochastic gradients  $(g_j)_{j=1}^t$  where  $g_j := \nabla_{\theta_j} L(x_j, \theta_j, y_j)$  and  $(x_j, y_j) \subset \mathcal{X} \times \mathcal{Y}$  is the sampled mini-batch for each  $j \in \{1, \dots, t\}$ .

Indeed, the vast majority of commonly-used first-order optimizers (SGD, RMSProp, Adadelta, AdamW, Lion) ignore the structure of each individual layer of weights of a deep neural network and treat the parameters simply as a long concatenated vector made up of the flattened parameters of each layer.

An idea which has regained popularity in recent years is to exploit the individual structure of each layer of the matrix, leveraging this information to compute update directions tailored to each layer. For illustration’s purpose, consider an MLP defined by

$$f($$

‘ The library [Modula](#) is built precisely on this notion, explicitly constructing a mathematical object called a *module* (not to be confused with the traditional interpretation in commutative algebra). These

A current SOTA language model is [LSY<sup>+</sup>25], which uses a mixture of experts (MoE) transformer model architecture. The model explicitly uses Muon to optimizer linear (non-embedding) layers. Namely

## 4.1 Related Work

## 4.2 Understanding Muon

The following exposition builds on the work of [BN24a], [BN24b], [Ber25], [JJB<sup>+</sup>24], [PXA<sup>+</sup>25] supplemented with some additional pedagogy and proofs. **TODO: give a nice Muon explanation**

### 4.2.1 Metrized Deep Learning

### 4.2.2 Dualized Gradients and Metrized Deep Learning

The paper [BN24a] revitalize an old principle in optimization theory. Namely, the insistence on viewing the gradient as a vector living in the dual space to the corresponding weight space.

Consider a differentiable loss function  $L : \Theta \rightarrow \mathbb{R}_{\geq 0}$  with respect to the weight space  $\Theta \subseteq \mathbb{R}^d$ . One can consider the first order Taylor approximation of  $L$  about arbitrary

$$L($$

### 4.2.3 Proof of Orthogonal Update step

### 4.2.4 Muon's Relationship to Shampoo

The Shampoo optimizer ([GKS18]) is intimately connected to the Muon update step. Indeed

### 4.2.5 The Newton Schulz Algorithm

Under the framework of steepest descent under the RMS  $\rightarrow$  RMS norm, the crucial step size for linear layers is seen as

## 4.3 Ablations on Simple Problems

To gain a better understanding of the dynamics of Muon, we study its behavior in simplified settings using both synthetic data and common toy datasets. Our goal is to examine

how Muon compares to SGD and AdamW in these controlled environments, and to identify the minimal conditions under which Muon offers practical advantages.

The experiments were all run with the [following codebase](#)

### 4.3.1 Linear Regression

To whet our appetite, we begin by studying the case of linear regression.

Since Muon explicitly depends on the weight space that admits a matrix structure, we propose the following multi-output linear regression problem as the simplest for which Muon can be applied (methodology for synthetic data generation will then be discussed in detail [later](#)):

$$Y = XW + \varepsilon \quad \text{where:} \quad \begin{cases} X \in \mathbb{R}^{N \times d} \text{ is our feature matrix} \\ W \in \mathbb{R}^{d \times D} \text{ is our weight matrix }^1 \\ Y \in \mathbb{R}^{N \times D} \text{ is our target space} \\ \varepsilon \in \mathbb{R}^{N \times D} \text{ noise where } \varepsilon_{i,j} \sim \mathcal{N}(0, \sigma_j^2) \text{ for } \sigma_1^2, \dots, \sigma_D^2 \in \mathbb{R}_{>0} \end{cases}$$

Indeed, excluding noise, this is an analogue of a hidden linear layer of a neural network. The objective (or loss) function is then given by

$$L(W; X, Y) := \|Y - XW\|_F^2 \quad (*)$$

where  $\|A\|_F := \sqrt{\sum_{i,j \in [m] \times [n]} |a_{ij}|^2}$  denotes the Frobenius norm of a matrix.

This objective function is manifestly convex in  $W$ . Indeed, we have

$$L(W) = \|Y - XW\|_F^2 = \|\text{vec}(Y - XW)\|_2^2 = \|\text{vec}(Y) - (\mathbb{1}_D \otimes X)\text{vec}(W)\|_2^2$$

where  $\otimes$  denotes the standard Kronecker Product and  $\text{vec}()$  denotes the vectorization of a rank  $k$  tensor into a rank 1 tensor.)

Clearly  $(*)$  is simply a quadratic function of the form  $f(z) = \|b - Az\|_2^2$  with positive-definite Hessian  $A^T A$

Using elementary properties of the tensor product (namely that  $(A \otimes B)^T = A^T \otimes B^T$  and  $(A \otimes B)(P \otimes Q) := AP \otimes BQ$  whenever such a product makes sense), one can easily see that the identity:

$$(\mathbb{1}_D \otimes X)^T (\mathbb{1}_D \otimes X) = \mathbb{1}_D \otimes (X^T X)$$

The corresponding Hessian is therefore uniquely determined by the structure of the feature matrix  $X$ . While such a simple problem admits an analytic solution (indeed  $W^* = (X^T X)^{-1} X^T Y$  provided  $X^T X$  is invertible), the study of gradient-based optimizers on this problem can lead to valuable insights.

While such a simple problem admits an analytic solution (indeed  $W^* = (X^T X)^{-1} X^T Y$  provided  $X^T X$  is invertible), we study the case where the number of datapoints is large

We consider both the case of full gradient methods and stochastic gradient methods. The gradient of  $(*)$  is given by

$$\begin{aligned}\nabla_W L &= \nabla_W \text{tr} \left( (Y - XW)^T (Y - XW) \right) \\ &= \nabla_W [\text{tr}(Y^T Y) - 2\text{tr}(W^T X^T Y) + \text{tr}(W^T X^T X W)] \\ &= 2X^T (XW - Y)\end{aligned}$$

The stochastic gradient for minibatch  $(X_t, Y_t) \subset \mathbb{R}^{N \times d} \times \mathbb{R}^{N \times D}$   $t \in \mathbb{N}$ , which we denote by  $G_t$ , is given by

$$G_t = \nabla_{W_t} L(W_t) = 2X_t^T (X_t W_t - Y_t)$$

### 4.3.2 A Quadratic Problem

Another commonly studied and generally well understood problem in optimization is the quadratic problem.

In the simplest possible case, this problem is formulated as

$$\text{minimize} \left( \varphi(x) := \frac{1}{2} x^T A x \right) \quad (4.1)$$

Where  $x \in \mathbb{R}^{m \times n}$  and  $A \in \mathbb{R}^{m \times m}$  is usually a symmetric positive semidefinite.

Again, since Muon insists upon a weight space with a matrix structure, we reformulate this problem into its simplest matrix analogue:

$$\text{minimize} \left( q(W) := \frac{1}{2} \text{trace}(W^T Q W) \equiv \|Q^{\frac{1}{2}} W\|_F^2 \right) \quad (4.2)$$

where  $W \in \mathbb{R}^{m \times n}$  and  $Q \in \mathbb{R}^{m \times m}$

We note that again vectorization reduces (4.2) to a problem of the form of (4.1):

$$\begin{aligned}q(W) &= \frac{1}{2} \sum_{i=1}^n (W^T Q W)_{ii} \\ &= \frac{1}{2} \sum_{i=1}^n w_i^T Q w_i \quad (\text{where } w_i := \text{col}_i(W)) \\ &= \frac{1}{2} \text{vec}(W)^T (\mathbb{1}_n \otimes Q) \text{vec}(W)\end{aligned}$$

**TODO: Discuss a bit with antonio about the whole connection of these two problems, is parameter structure even a proper thing? Also the homogenous vs heterogeneous Hessian situation.**

Provided the matrix

### 4.3.3 Logistic Regression

Similarly to the linear regression case, we consider multi-class classification with a softmax function. Again, because we are interested in the Muon optimizer, this is the simplest case for which the weights admit a natural matrix structure.

One can view Logistic regression as a neural network of just one layer so it is certainly worthy of scrutiny.

- Generate  $X$  and  $W^*$  as in the linear regression case
- Let  $Z = XW^*$  and  $P = \text{softmax}(Z) + \varepsilon$  for noise  $\varepsilon$
- Define  $Y_{ik} := \begin{cases} 1, & \text{if } k = \arg\max_{\ell \in \{1, \dots, K\}} P_{i\ell} \\ 0, & \text{else} \end{cases}$  for each  $i \in \{1, \dots, N\}$  (or in plain english: each row of  $Y$  is a one hot encoded vector in  $\mathbb{R}^K$ )
- Initialise a random  $W \in \mathbb{R}^{(d+1) \times D}$  and compare different optimizer's performance on minimizing loss.

Let  $f(X; W) := \text{softmax}(XW)$  As is standard for classification, we consider the cross-entropy loss:

$$L(W) = -\frac{1}{N} \sum_{i=1}^N \sum_{k=1}^K Y_{ik} \log(P_{ik}) \quad \text{where } P = \text{softmax}(XW)$$

Let us compute the gradient. We denote  $P = \text{softmax}(Z)$  where  $Z := XW$ . We invoke the chain rule to simplify matters:

$$\nabla_W L = \nabla_Z L \cdot \nabla_W Z = \frac{1}{N} X^T (\text{softmax}(XW) - Y)$$

### 4.3.4 Small MLPs

## 4.4 SVD structural analysis

The Muon optimizer makes use of the Newton Schulz algorithm

### 4.4.1 CIFAR-10 ResNet

The CIFAR speedrun is a competitive benchmark which aims to achieve a specified test set accuracy in the shortest possible training time (standardized in NVIDIA A100-minutes). Researchers submit their optimizations, which can include network architecture changes, alterations to the training pipeline and test time tricks, and the submission is then benchmarked by averaging over a number of random seeds. The goal is to achieve the shortest average training time while exceeding a pre defined test set accuracy.

Remarkably, with enhancements using Muon (**TODO: cite the right thing**), the current

record is a mere 2.59 seconds to achieve 94% training accuracy.

We make use of this repository to ablate on some hyperparameter choices for the Muon optimizer while enjoying minimal experiment runtimes.

### Comparing Newton Schulz approximation to full Gradient Orthogonalization

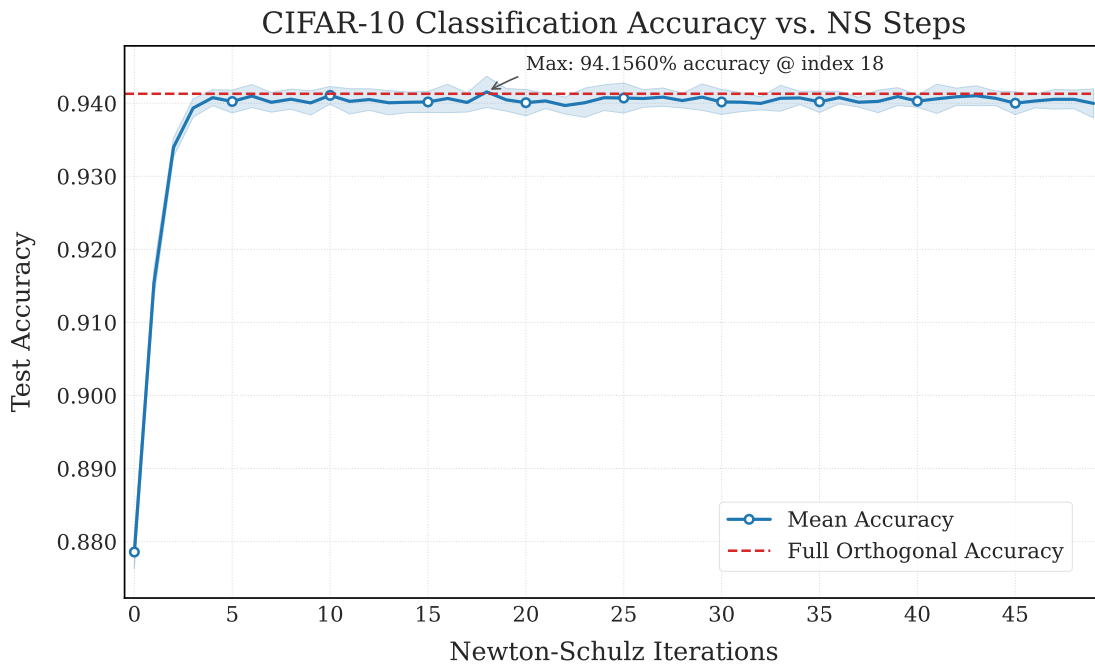


Figure 4.1: Ablation varying number of Newton Schulz steps performed and comparing with an exact orthogonalization computed using the SVD of the matrix. Each data point on the x-axis corresponds to an averaged final test accuracy over 20 random seeds with error bars representing a 95% confidence interval.

## 4.5 Muon on nanoGPT and plainLM

## 4.6 Experimental



## **Chapter 5**

### **Discussion and Future Work**



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## **Appendix A**

# **How Exponential Moving Averages Smooth the Update Signal**

It is generally well accepted that